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Cover: Satellite image captured by JMA (left) and forecasted 10-m wind speed (right) of cyclone 'Burevi' at 0600 UTC on 02 December 2020 See *J.Natn.Sci. Foundation Sri Lanka* 2022 50(3): 675 – 684

EDITORIAL

Seeing is believing

The James Webb Space Telescope, the largest and the most powerful space telescope that has ever been constructed by NASA, has begun its science operations orbiting the Sun in line with Earth on a trajectory located at an antisunward point approximately 1.5 million kilometers away from Earth. In Webb, revolutionary technologies have been used to study the phases of cosmic history of the universe. The continual expansion of the universe has caused the very first stars and galaxies formed over billions of years ago that emitted ultraviolet and visible light to undergo redshifts to observe that radiation in the infrared range. Unlike the Hubble space telescope which views the universe in visible and ultraviolet light, Webb is equipped with an infrared telescope to focus on infrared radiation capable of peering through gas and dust clouds to see distant objects with an unprecedented resolution and sensitivity. Webb is also capable of studying the nearby universe making it a powerful tool to explore the planets and other bodies in our solar system. Already, to the excitement of scientists and readers, headlines appear in news reports on exoplanets, old star clusters and distant galaxies with their spectacular images taken by Webb with much improved clarity. Scientists, with time will be able to come up with new information about the universe with interpretations, models and even new theories.

The newest technologies NASA has used in the Webb include a foldable sunshield which prevents the heat and light from the Sun, Earth, and the Moon reaching the telescope's heat-sensitive optics, a foldable mirror system in the appropriate dimensions to receive the infrared radiation emitted from celestial objects and maintenance of lasting cryogenic operating temperatures to prevent emission of infrared radiation by this massive observatory itself. Webb is considered an engineering marvel that has become a reality due to meticulous planning over the years and its implementation by an international collaboration between NASA and its partners in Europe and Canada involving thousands of engineers and hundreds of scientists representing a number of universities, organizations, and companies from U.S. and several other countries.

Thus, Webb serves as a classic example that unifies the scientist, the engineer and experts representing other areas in the quest to finding the origin and evolution of the universe. The team on its way to reach this goal has made groundbreaking inventions, innovations leading to new materials and several new technologies which can be used for the benefit of humanity. Applications are already seen in the fields of medicine and engineering. This project shows some parallels to projects launched several decades ago in exploring the subatomic particles which led to the introduction of global communication pathways like World Wide Web that revolutionized the entire communication industry. Webb can be seen as yet another example for how fundamental science can help shape and impact society.

In the context of Sri Lanka, lessons can be learnt from the approach made by NASA to successfully create this massive, amazingly powerful and precisely controllable space observatory. Webb is an example of teamwork which has been guided by proper leadership knowledgeable in correct scientific approach and methodologies which were put into practice. In Sri Lanka, in a culture that focusses on individual glory at personal and institutional level, at times at any cost, Webb shows that strategies must be in place with proper leadership to encourage the collective efforts of scientists, engineers and relevant experts in the country to address the national scientific and technological challenges it is facing at the moment. When considering the size of Sri Lanka, such an approach seems timely and viable. Webb has provided the impetus to believe in ourselves to use human ingenuity to address issues related to energy, environment, agriculture and earth resources etc. recognizing them as high priorities.

Sumedha Jayanetti

REVIEW

Crop Science

Rice production in nutrient-limited soils: Strategies for improving crop productivity and land sustainability

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Summary: Productivity improvement of rice cultivation in most parts of Africa and Asia has slowed down, which is largely due to declining soil fertility. Widely reported soil fertility constraints are deficiencies of nitrogen (N), available phosphorus (P), sulphur (S) and zinc (Zn). The amount of N, P and potassium (K) removed with the harvest of 5 t grain ha⁻¹ is estimated as 27-97 kg N ha⁻¹, 4-20 kg P ha⁻¹, and 8-32 kg K ha⁻¹. Straw produced from one ha of land also contains 25-40 kg N, 3.5-10.5 kg P, 70-85 kg K, 2.5-5 kg S, 15-20 kg Ca, 5-15 kg Mg and 200-350 kg Si. Soils do not have an inherent capacity to replenish these amounts of nutrients over the very short period. Most rice farmers also do not have the ability to apply nutrients due to financial limitations and/or lack of awareness. In order to sustain rice cultivation in those fields, soil fertility constraints must be addressed without prioritising unrealistic synthetic fertiliser applications. Improving soil organic carbon and nutrient pools by incorporating organic manure is needed. Crop rotation, including fallow, minimal or zero tillage, sitespecific nutrient management, and introduction and promotion of biological nutrient enrichment are other alternative and promising soil-carbon and nutrient-improvement strategies. The combination of these measures is known as an integrated plant nutrient management system (IPNS), and it must be implemented to maintain and/or improve soil fertility. Apart from agronomic interventions, cultivar selection and crop improvement programmes need to particularly focus on lowinput soils, as farmers are resource-poor. Due to the complex soil, socio-economic, and institutional structures in these regions, an integrated and participatory strategy incorporating all stakeholders is essential for implementing agronomic interventions and productivity improvement programmes.

CURRENT CONTEXT

Most rice is grown in Asia (143.4 million ha), Africa (10.5 million ha) and the Americas (7.2 million ha). Moreover, 57% of the global rice cultivation is under irrigated lowland conditions. The remaining rice is grown under rainfed lowland (31%) or upland (9%) conditions, and less than 3% as deep-water or mangrove-associated rice culture (Haefele et al., 2014). The rate of increase in global rice yields has slowed down, limiting the availability of milled rice for consumption (Dobermann et al., 2002; Ladha et al., 2003) (Figure 1). It is also estimated that world rice production must increase by approximately 1% annually to meet the growing demand for food resulting from population growth and economic development (Peng et al., 2004; Mishra et al., 2018; Saito et al., 2018). But increases in rice production have remained below the level required in many countries (Cooper et al., 1999; Ray et al., 2012). Despite this growing demand, the extent used for rice cultivation remains unchanged in the recent past (Figure 1). Therefore, it is essential to revisit the necessary efforts need to be made to ensure sustainability and improve the productivity of rice-cultivating systems.

Input used in rice cultivating systems

Declining soil fertility is one of the main factors for stagnating rice yields in Africa and Asia (Timsina & Connor, 2001; Dobermann et al., 2002; Saito &

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Figure 1: Changes in global rice production, consumption and cultivation extent in the past decade (Note: data were extracted from https://www.statista.com/statistics/271969/world-rice-acreage-since-2008/ on 10.04.2022)



Figure 2: Yield gaps of rice and the factors associated. Note: the gaps between actual and attainable, and attainable and potential are not drawn proportionately.

Futakuchi, 2009; Weerakoon *et al.*, 2011; Rodenburg *et al.*, 2014; Katsura *et al.*, 2016; Mayamulla *et al.*, 2017; Stewart *et al.*, 2020). This has been caused to a considerable extent by the continuous crop and residue removal, negative nutrient balances and soil degradation (Cassman *et al.*, 1998; Dobermann *et al.*, 2002; Nhamo *et al.*, 2014; Kekulandara *et al.*, 2019). Therefore, many research studies have been conducted around the globe to improve the productivity of rice-based cropping systems. One of the critical attempts in most of this research is to enhance soil fertility along with other inputs and

management, such as irrigation water supply, pest and disease control, and mechanisation. Such systems can be classified as intensive, high-input rice-cultivating systems (Figure 2). Due to the application of all required inputs and intensive crop management, farmers in high-input rice-cultivating systems could cover a higher percentage of the yield gap between attainable and actual yields by increasing productivity. Even though improving soil fertility to achieve high productivity is desirable, one of the main problems faced by many rice farmers in Asia and Africa is the difficulty in applying recommended rates of synthetic fertilisers. This is due to their poor economic status, high fertiliser costs, unavailability of inputs, and/ or production risks (Weerakoon et al., 2011; Haefele et al., 2003, 2014; Nhamo et al., 2014; Mayamulla et al., 2017; Stewart et al., 2020).

The low-input rice-cultivating systems defined in this study have the nutrient-supplying capacity (either by inherent soil nutrient pools or through the application of fertilisers) less than the amount and rate required by the rice plant to reach attainable grain yields (t ha⁻¹) in a particular environment, when other inputs or characteristics (e.g., drought, pest and diseases, Fetoxicity, low-CEC, low-pH) are not yield limiting (Figure 2). One or more nutrients may act as rate-limiting variables in this system. In low-input rice-cultivating systems, farmers have a variety of options for enhancing soil fertility, despite recommendations from extension services to rely primarily on synthetic fertilisers, which are typically supplied at a subsidised rate (Weerahewa, 2004; Asai et al., 2009; Druilhe & Barreiro-Hurlé, 2012; Nhamo et al., 2014; Stewart et al., 2020). Due to the continuous increase in fertiliser prices, more and more rice fields are likely to become nutrient deficient in the future (Shepherd et al., 2016). Despite this, the importance of rice production in Africa and Asia has significantly increased over the last decades (Nhamo et al., 2014), as rice plays a pivotal role in improving household food and nutritional security as well as national economies in these regions (Diallo et al., 2016; Katsura et al., 2016; Stewart et al., 2020). Therefore, to sustain rice cultivation in lowinput agricultural systems, soil fertility constraints must be addressed before other technologies and policies can become effective in improving productivity. When doing so, soil management options besides applying synthetic fertilisers, such as the addition of organic matter, crop rotation, and tillage practices, should receive high priority as those are vital to overcoming nutrient limitations.

Existing nutrient limitations

Soil fertility differences exist from continental scale to paddock and between and within farms (Tsujimoto et al., 2013; Nhamo et al., 2014; Katsura et al., 2016). Among the three major nutrients, nitrogen (N) is the most widely limiting, while phosphorus (P) ranks second, and potassium (K) deficiency is rarely reported (Dobermann et al., 2003; Ladha et al., 2003; Shen et al., 2004; Rodenburg et al., 2014; Diallo et al., 2016; Somaweera et al., 2016, 2017; Stewart et al., 2020). Moreover, deficiencies in sulphur (S) and zinc (Zn) are common in many regions to variable degrees (Buri et al., 2000; Duxbury et al., 2000; Ladha et al., 2003; Haefele et al., 2014; Rodenburg et al., 2014). Apart from nutrient limitations, constraints related to pH and cation exchange capacity (CEC) also occur in isolated areas or problem soils (Ladha et al., 2003; Haefele et al., 2004, 2014; Shen et al., 2004; Gami et al., 2009; Saito & Futakuchi, 2009; Rodenburg et al., 2014; Diallo et al., 2016; Dossou-Yovo et al., 2016;). In general, the occurrence of nutrient deficiency decreases in the sequence of uplands > rainfed lowlands > irrigated lowlands > deepwater/ mangrove areas (Haefele & Konboon, 2009; Haefele et al., 2014; Nhamo et al., 2014). This is caused by the transport of nutrients and soil particles from higher to lower parts of the landscape and due to the differences in soil moisture availability. In contrast, soils with problems such as Fe-toxicity are more frequent in the lower parts of the landscape and in locations where seepage water accumulates (Becker & Asch, 2005; Haefele et al., 2014; Rodenburg et al., 2014; Suriyagoda et al., 2017; van Oort, 2018). Based on the type of nutrient limitation, rice-growing soils can broadly be categorised as;

- problem/very poor soils (show multiple constraints such as iron and aluminium toxicity, low CEC and pH, high P-fixation, shallow soils, limited waterholding capacity, and deficiency in macro- and micro-nutrients),
- (ii) poor soils (no constraints as above, but low in many macro and micro nutrients including N and P), or
- (iii) good soils (no constraints, mostly limited by N and/ or P only) (Haefele *et al.*, 2014).

All these soils are found both in high and low-input rice cultivating systems (Figure 2). However, in this study, we mainly focus on the second and third categories of rice-growing soils, which are limited by macro- and micronutrients.

Average recovery efficiency (*i.e.* percentage taken up of the amount applied) of fertiliser N by rice plants is

only about 30 to 47% (Dobermann et al., 2002; Ladha et al., 2005). The low recovery efficiency of N is mainly due to high losses, making it unavailable for current or future crops. The recovery efficiency of P is less than 10% in most instances (Sanchez & Salinas, 1981; Dobermann et al., 1998), but values up to 20% have also been reported (Holford, 1997). Unlike N, most excess P is retained in the soil after forming complexes, making it available for future crops after solubilisation at the optimal range of pH (Lambers et al., 2008; Sirisena & Suriyagoda, 2018). In most rice-growing systems in Asia and Africa, K balances are negative (Dobermann et al., 1998; Ladha et al., 2003; Haefele et al., 2004) unless crop residues are incorporated (Wihardjaka et al., 1999; Sahrawat, 2000). However, mass balances of K inputs, K uptake, and exchangeable K in rice soil show that a large part of K uptake is from non-exchangeable pools, and the mobilisation of non-exchangeable K is apparently plant-induced (Wihardjaka et al., 1999). Moreover, high soil K reserves can buffer negative K balances for decades (Haefele et al., 2004; Shen et al., 2004; Diallo et al., 2016; Somaweera et al., 2017). Irrigation water also contains more K than N and P, allowing K to move across rice fields in cascade irrigation networks (Suriyagoda et al., 2017). Therefore, K deficiency in rice is less prevalent than N and P and most common in highyielding systems.

Rice grain N, P, and K content (expressed on mg g⁻¹ basis) reported from different countries are mainly in the range of 9-15 mg g^{-1} for N, 1.0-4.3 mg g^{-1} for P and 0.6-4 mg g⁻¹ for K (Dobermann et al., 1996, 2002; Timsina & Connor, 2001; Xu et al., 2015; Somaweera et al., 2016, 2017). Moreover, internal N-use efficiency (i.e. kg grain DW kg⁻¹ N in aboveground biomass) ranges between 31-112 kg grain DW kg⁻¹ N uptake, and for P and K, the values are 192-900 kg grain DW kg⁻¹ P uptake and 32-115 kg grain DW kg⁻¹ K uptake, respectively (Witt et al., 1999; Dobermann et al., 1996, 2002; Dwivedi et al., 2004; Buresh et al., 2010; Das et al., 2009, 2014; Haefele et al., 2003; Singh et al., 2005; Xu et al., 2015). Therefore, the amount of N, P and K stored in aboveground biomass when producing 5 t grain ha⁻¹ are 45-161 kg N ha⁻¹, 5-26 kg P ha⁻¹, and 43-156 kg K ha⁻¹. It is also reported that about 55-65% of N, 65–70% of P, and 15-20% of K absorbed by rice plants are stored in the grains (Dobermann & Fairhurst, 2000; Sahrawat, 2000). Therefore, the amount of N, P and K removed with the harvest of 5 t grain ha^{-1} can be estimated as 27-97 kg N ha⁻¹, 4-20 kg P ha⁻¹, and 8-32 kg K ha⁻¹. The extensive range of N, P, and K removal is due to the differences in grain N, P, and K contents, mainly affected by the variation in soil fertility.

Most farmers cannot apply sufficiently high rates of N, P, and K to support the uptake of elements to reach expected yield targets. Consequently, as soil nutrient limitations develop, an increase in nutrient-use efficiency (or a decrease in tissue nutrient content) can be anticipated. As a benchmark, straw N, P, and K content less than 10 mg N g^-1, 0.6 mg P g^-1 and 1.8 mg K g^-1 at maturity are generally considered deficiency thresholds for N, P, and K when identifying soils, which have a low capacity to supply N, P and K to rice plants (Dobermann et al., 1998; Dobermann & Fairhurst, 2000; Vandamme et al., 2016). Tillering is the most sensitive stage of rice crop development when P deficiency symptoms arise, although N deficiency symptoms appear throughout the crop life cycle (Fageria, 2003; Somaweera et al., 2017). Therefore, the reported N-, P- and K-use efficiency values, straw N, P, and K contents, and sensitive crop developmental stages can be used when identifying the existence and severity of N, P, and K deficiencies in rice culture. Moreover, due to the discrepancy between the growth stage at which critical P level is identified (*i.e.*, at maturity) and the most sensitive growth stage of rice to P deficiency (*i.e.*, at tillering), the relationship between those two needs to be explored.

Zinc deficiency is one of the most widespread micronutrient deficiencies in rice production in Africa and Asia (Buri *et al.*, 2000; Dhanapala, 2000; Duxbury *et al.*, 2000; Timsina & Connor, 2001; Quijano-Guerta *et al.*, 2002; Ladha *et al.*, 2003; Wissuwa *et al.*, 2006). Availability of Zn is generally decreased after flooding rice soils and is associated with an increase in soil pH after flooding (Fageria *et al.*, 2011). Significant positive correlations between total soil carbon and available P and Zn were also found in West Africa (Buri *et al.*, 2000). Thus, knowledge of rice tolerance to many stresses is useful when looking for Zn-deficiency resistant genotypes because much of the information on other nutrient deficiencies, such as P, is available in most rice germplasms.

Apart from Zn, S deficiency occurs in certain parts of Asia, Africa and America (Blair *et al.*, 1979; Dobermann *et al.*, 1998; Yamaguchi, 1999; Buri *et al.*, 2000; Dhanapala, 2000; Duxbury *et al.*, 2000; Tsujimoto *et al.*, 2013). As a benchmark, straw S content less than 0.5 mg S g^{-1} at maturity is generally considered the deficiency threshold for S (Dobermann *et al.*, 1998). When N, P, and K were applied without S in Ghana, biomass production was not increased while the tissue N content and N:S ratios increased, indicating S deficiency (Tsujimoto *et al.*, 2013). In another study, Yamaguchi (1999) found that the maximum quantity of S needed to alleviate the deficiency in rice was about 30% of the N requirement. Moreover, the topsoil sulphate-S content was positively correlated with total soil carbon, available P, and CEC (Buri *et al.*, 2000), indicating the occurrence of multiple nutrient deficiencies and the importance of soil properties in determining the availability of mineral elements. Consequently, S deficiency diagnosis in rice plants with a single variable often leads to an error of judgement. Therefore, the evaluation is suggested to take several variables into account, *e.g.*, total-S, total-N/total-S ratio, sulphate-S, and sulphate-S/total-S ratio (Yamaguchi, 1999), requiring detailed laboratory procedures. This has also become a limitation when detecting S deficiency in most of rice fields at its initial stages.

Silicon (Si) is increasingly considered as a beneficial element for plant growth. Rice is efficient in accumulating Si from the soil, and shoot Si content is about 10% of shoot dry weight (Yamamoto et al., 2012). Silicon in soil may be lost if the Si-cycle is interrupted (Meharg & Meharg, 2015). Most of the Si taken up can be returned to the soil with the incorporation of straw and paddy husk. However, rice cultivation over centuries with considerable straw removal can deplete the soil Si pool, risking the sustainability of rice cultivation in certain regions (Savant et al., 1997). Apart from the positive effects of Si on plant growth, there is evidence that Si fertilisation ameliorates Fe, manganese (Mn), and arsenic (As) toxicities (Fu et al., 2012; Meharg & Meharg, 2015; Suriyagoda et al., 2018), and improves resistance to pest and disease incidences (He et al., 2015). Moreover, Si fertilisation decreases Ca, Fe, Mg, N, and P in rice straw (Ma & Takahashi, 2002), demonstrating a negative relationship between Si and other essential element levels in rice. However, if rice plants can produce the same or higher grain yields with reduced nutrient content (i.e., enhanced nutrient-use efficiency) due to improved Si nutrition, applying Si to Si-deficient fields could be advantageous.

Unlike the occurrence of deficiencies of most of the nutrients in rice fields as discussed above, the presence of a high content of iron (Fe²⁺) in soil has caused toxicity to rice plants, mostly in lowlands in Africa and Asia, causing even complete crop failure (Becker & Ash, 2005; Fairhurst *et al.*, 2007). It is reported that up to 12% of the rice lands in Africa are potentially affected by iron toxicity (van Oort, 2018). The Fe²⁺ concentration (expressed as mg L⁻¹) in soil solution that affects rice yield can vary in the range of 10 - 2000 mg L⁻¹ (Becker & Ash, 2005). Toxicity caused by Fe²⁺ is a complex disorder, as it is associated with soil acidity, low-CEC, and nutrient deficiency (Haefele *et al.*, 2014). Moreover,

due to the presence of a high concentration of Fe^{2+} in lowland soils, availability of Mn, Zn, P, and K in the soil can be low, causing deficiencies of those nutrients to rice plants (*i.e.*, multiple nutrient disorder) (Fageria & Zimmermann, 1988; Suriyagoda *et al.*, 2017).

AGRONOMIC INTERVENTIONS TO IMPROVE FERTILITY IN RICE FIELDS

Improving soil organic carbon (SOC) and nutrient pools

Addition of organic matter

Continuous cultivation on the same land, removal of harvested grains, lack of incorporation of organic matter, including straw produced in preceding seasons, constant and fast decomposition of retained organic matter, and losses due to runoff have led to a decline in organic carbon content in some of the rice-growing soils (Singh et al., 2005; Das et al., 2014). Most of the rice-growing soils in the globe have SOC values in the range of 1-3%, and values less than 0.5% have also been reported (Mukhopadhyay et al., 2013; Xu et al., 2015; Diallo et al., 2016). It is also known that SOC content is lesser in aerobic soils than in anaerobic soil, and in tropical soils than in temperate region. (Sahrawat, 2004). Therefore, tropical soils experiencing aerobic or frequent wetting and drying conditions decompose SOC rapidly and release N (Sahrawat, 2006). Even though the application of external sources of N such as urea is preferred to organic manure low in N and soils deficient in N to enhance the decomposition, such applications are not required in most tropical soils due to the inherently fast rate of organic matter decomposition.

Soil organic carbon can improve nutrient-retention capacity, water-stable aggregates, and porosity, and reduce bulk density, thereby facilitating crop establishment, root growth and nutrient uptake (Lal et al., 1989; Timsina & Connor, 2001; Ladha et al., 2003; Dwivedi et al., 2004; Bi et al., 2009; Suriyagoda et al., 2014) (Table 1). As the C/N ratio of soil organic matter is relatively constant, incorporation of organic matter also improves soil N pools (Cassman et al., 2002). However, the application of synthetic N, P, and K fertilisers does not improve long-term SOC or nutrient pools (Gami et al., 2009). Management practices can significantly affect soil carbon storage through carbon inputs and losses. Still, detection of the changes in SOC is often difficult due to the small magnitude of changes relative to the total stock, except in long-term studies (Conen et al., 2003). Despite the importance of maintaining adequate SOC pools to ensure a favourable soil environment for rice plants and increasing awareness among farmers, there has been insufficient attention to this aspect (Dossou-Yovo *et al.*, 2016; Stewart *et al.*, 2020). Therefore, an understanding of the benefits of increasing SOC should be raised among rice farmers. Moreover, long-term investigations of the changes in SOC and nutrient balances, particularly in nutrient-limited soils, are required.

Improving SOC pools increases grain yield of rice in China (Pan et al., 2009; Huang et al., 2013; Hu et al., 2016; Zhou et al., 2020), Laos (Roder et al., 1995; Asai et al., 2009), Senegal (Diallo et al., 2016), East and South Africa (Nhamo *et al.*, 2014 and references therein), Nepal (Gami et al., 2009), Bangladesh (Hossain et al., 2016), Thailand (Wonprasaid et al., 1996; Homma et al., 2003), Cambodia (Ly et al., 2016) and Sri Lanka (Sirisena et al., 2016). In China, grain productivity increased by 21% and 24%, and C accumulation by 72% and 103%, respectively, when pig manure (8400 kg ha⁻¹) or straw (4500 kg ha⁻¹) was applied (Pan et al., 2009). Hu et al. (2016) reported that the application of a lower amount of organic manure, *i.e.*, 2100 kg ha⁻¹, can be recommended for the implementation of organic rice farming, and higher rates of organic manure application may increase the risks of pests and diseases. Huang et al. (2013) also observed a yield increase by 5.2% when crop residues were retained. In Thailand, grain yield increased, on average, by 49% with the application of 6.25 t farmyard manure ha^{-1} , and by 50% with the application of 60 kg N, 13 kg P and 25 kg K ha⁻¹ in comparison to the fields which did not receive synthetic fertilisers (Khunthasuvon et al., 1998). Similar results are reported by Wihardjaka et al. (1999) in Indonesia. Becker & Johnson (2001) observed an increase in grain yield of upland rice by 26% when SOC content increased by 19% across several sites in West Africa. Therefore, sole application of organic matter or combined application of organic matter with synthetic fertiliser have increased the grain yield of rice and improved soils. Thus the application of organic materials with or without synthetic fertilizer should be promoted in fields low in SOC with site-specific approaches.

Widely available materials to improve the soil organic carbon pool are rice straw, animal manure, farmyard manure, and other crop residues. Returning one t ha⁻¹ of straw (rice, wheat or maize) to the soil each year can sequester about 130 kg C ha⁻¹ yr⁻¹ (Lu *et al.*, 2009). Moreover, about 40% of N, 30–35% of P, and 80–85% of K absorbed by rice remains in the straw at maturity (Dobermann & Fairhurst, 2000;

Das *et al.*, 2014). Incorporation of one t of straw per hectare accounts for the addition of 5–8 kg N, 0.7–2.1 kg P, 12–17 kg K, 0.5–1 kg S, 3–4 kg Ca, 1–3 kg Mg and 40–70 kg Si on a dry weight basis (Dobermann & Fairhurst, 2000, 2002; Dobermann & Witt, 2000; Das *et al.*, 2014). When rice straw is not added to fields, P needs to be re-applied at about 17 kg ha⁻¹ every two crops on acid sandy lowland soils to maintain the grain yield at about 2.5–3.0 t ha⁻¹ in Cambodia (Pheav *et al.*, 2003). Similar results were obtained for N in China (Pan *et al.*, 2009). Therefore, incorporation of rice straw improves soil quality including nutrient content and retention (Kumar & Goh, 1999; Wihardjaka *et al.*, 1999; Sahrawat, 2000; Suriyagoda *et al.*, 2014), and the effects are more pronounced in clayey than in sandy soils (*i.e.*,

due to greater retention), and in flooded than in aerobic soil (*i.e.*, due to higher solubility) conditions (Suriyagoda *et al.*, 2014; Haefele *et al.*, 2016). Rice straw retention also influences availability of micronutrients such as Zn and Fe (Dobermann & Fairhurst, 2000, 2002). Straw incorporation facilitates greater root system growth plasticity with the production of deeper and finer roots allowing the rice plants to explore a large volume of soil in search of nutrients (Suriyagoda *et al.*, 2014). However, straw produced from nutrient-limited soils contains low contents of nutrients, and thus the addition of straw will at best reduce the rate of decline in soil fertility (Table 1). Therefore, nutrient-dense alternative sources should be incorporated into the soil to improve its nutritional condition.

Table 1: Methods widely used to improve soil organic carbon (SOC) and nutrient pools, and the advantages and disadvantages of those methods

Method	Advantages		Disadvantages
Addition of organic matter •	Improve nutrient-retention capacity, water- stable aggregates and porosity	•	Variable in types and quality
		٠	Require large amounts (content of nutrients is low)
•	Reduce bulk density	•	Labour cost for collection and application is high
•	Facilitate crop establishment, root growth and nutrient uptake	•	May fix plant available N early in the season
Fallow or crop rotation •	Improve- productivity, soil fertility, SOC pool	•	Improvement of soil quality depends on the types of crops used during fallow periods or in rotation
•	Reduce bulk density		
•	Avoid risk in cultivation under environmental stresses	•]	Depends on area specific socio-economic, climatological and ecological conditions
Minimum tillage •	Minimize the loss of SOC, soil structure and nutrients	•	Effectiveness is variable and depends on the region, climatic condition, cropping system, rate and time of
•	Increase bulk density		and duration of minimum till adoption
•	Saves labour, energy and water needed for puddling	•	May increase percolation losses during the season
Application of biochar •	Improve SOC and fertility	•	Large quantities are recommended
•	Improves soil physical characteristics	•	Largely variable in quality
		•	Bind mobile nutrients

Soil analyses after six years of experimentation in Nepal showed that treatments receiving organic sources of nutrients (4-12 t ha⁻¹ yr⁻¹) increased total soil C and N by 18% to 62% and 15% to 48%, respectively, compared with the NPK treatment (Gami *et al.*, 2009). The authors also observed a build-up of total and plant-available P in plots receiving farmyard manure. However, total and

available soil K were similar in all treatments (Gami *et al.*, 2001). Farmyard manure also contained a bacterial population of 5.2×10^6 cfu g⁻¹ (Mukhopadhyay *et al.*, 2013). After 11 years of experimentation in Sri Lanka the lowest yield was observed in the plots which did not receive any form of nutrients (2.85 t ha⁻¹). However, the yield in the plots receiving organic manure only (straw

from the previous season, 1 t green manure and 4 t cattle manure ha⁻¹) was 3.75 t ha⁻¹ and that in plots receiving both organic manure and synthetic fertilisers was 5.5 t ha⁻¹ (Sirisena et al., 2016). In China, there has been a prominent topsoil C accumulation of 0.1-0.4 t ha⁻¹ yr⁻¹ when pig manure was applied compared with adding synthetic fertiliser only (Pan et al., 2009). Therefore, incorporation of farmyard manure, in the absence of straw or together with straw, can be promoted to replenish soil C and nutrient pools in nutrient-limited soils. However, the general trend in rice cultivation is a decreasing use of local nutrient sources, mainly because of increasing opportunity costs of the labour needed for collection and application, and an increasing use of synthetic fertilisers when available, incurring high cost (Table 1). Between rice environments, the relative contribution from local nutrient sources increases from irrigated lowlands to rainfed lowlands to uplands, whereas synthetic fertiliser use has an opposite trend (Haefele et al., 2014, 2016). This is, of course, due to the comparatively higher risk and poorer farmers in most rainfed rice environments. Another reason might be the often poor soil fertility in rainfed environments, requiring more organic fertiliser to improve/maintain soil fertility.

Although the beneficial effects of organic manure incorporation, to improve the physical, chemical, and biological properties of soil and the grain yield of rice, have been shown numerous times, they depend on the types and quality of the organic manure available, and its use. Therefore, the amount of organic residues requires to reach productivity targets in different rice-based cropping systems may vary widely and is largely unknown. The amount of organic manure needed, its form, method of application and management, and nutrient releasing patterns are material, site, or region specific. Therefore, localised quantifications are required to maximise the efficiency of an integrated nutrient management system. Based on such calculations, large amounts of organic manure were sometimes recommended, e.g., 13-35 t ha⁻¹ of compost in Africa (Nhamo et al., 2014 and references therein). Such application rates are often far greater than what an average livestock farmer produces annually and what a rice farmer can afford and apply. A potential approacs in such situations is to spread out the application of organic manure across several seasons for gradual improvement of soil fertility, as a component in an integrated plant nutrient management system (IPNS). These practices will ensure long-term development of soil nutrient pools while minimising their losses.

Uptake (*i.e.*, recovery) efficiency of N from a single fertiliser application typically decreases in proportion

to the amount of N fertiliser applied (Reddy & Reddy, 1993). The same principle applies to available N derived from organic N sources such as legume green manures, cover crops, and animal manures. Potential nitrate leaching from manures can be equal or even greater than the losses from synthetic N fertiliser when the available N supply from either source exceeds crop demand by similar amounts for comparable times (Cassman et al., 2002). Similar results were reported for other nutrients across different regions (Zhang & He, 2004; Pratiwi & Shinogi, 2016; Sirisena & Suriyagoda, 2018). Moreover, substantial incorporation of organic fertiliser at once accelerates sudden release of greenhouse gases such as CH₄ and N₂O, and may contribute greatly to global warming (Yuan et al., 2017). In China, the amount of CH₄ emission is greater with organic manure (77.3 kg CH_4 ha⁻¹) than with urea (30.2 kg CH_4 ha⁻¹) and the blank control treatment (17.4 kg CH_4 ha⁻¹) (Yuan *et al.*, 2017). Similar results were observed by Zhao et al. (2015). Therefore, it is important to gradually enhance the soil's nutrient pool in order to minimise nutrient losses.

Incorporation of fallow or crop rotation

One of the most prominent approaches to improving soil fertility is the incorporation of fallow periods in the cropping calendar and improvement of the quality of the fallow vegetation, e.g., introduction of legume cover crops (Balasubramanian & Sekayange, 1992). Improvements in productivity, soil fertility, SOC pool, and reduction in bulk density with the incorporation of legumes in rotation, such as mungbean, pigeon pea, and forage cowpea, were observed in rice-based cropping systems in Bangladesh, India, and Sri Lanka (Timsina & Connor, 2001; Dwivedi et al., 2004; Singh et al., 2005; Malaviarachchi et al., 2016; Hossain et al., 2016) (Table 1). Das et al. (2014) observed that 70-106 kg N ha⁻¹, 10-14 kg P ha⁻¹ and 63-94 kg K ha⁻¹ could be incorporated to the soil with the recycling of 4.8-7.2 t ha⁻¹ of weed biomass under subtropical conditions in Meghalaya, India. In African uplands, where rice is grown with the incorporation of Mucuna spp., Canavalia spp., and Stylosanthes guianensis during the dry season, consistently high N accumulation was found. Grain yields of rice, which had been preceded by a legume fallow, were on average 0.2 t ha^{-1} or about 30% greater than that preceded by a natural weedy fallow control treatment (Becker & Johnson, 1998). However, some contradictory results have also been reported about the use of legume crops in rotations. For example, although including grain legumes in rotation with cereals can reduce N-fertiliser requirements compared with continuous cereal cropping, the degree of soil-N stock replenishment depends on the

amount of N removed with the harvested seeds (Cassman *et al.*, 2002). Overall, the incorporation of legume-based fallow to rice rotation has merits, particularly when noncrop legume species are used, and the net benefit of such systems need to be assessed.

Other studies showed that the SOC pool is affected by the types of crops incorporated into the rice cropping systems; for example, rice-rice (1470 kg ha⁻¹) and ricesoybean (677 kg ha⁻¹) rotations added more dry matter to soil than rice-tobacco (29 kg ha⁻¹) or rice-onion (not detectable) as most of the biomass of onion and tobacco is removed at harvest (Ratnayake *et al.*, 2017). Moreover, intensive land preparation used in onion and tobacco crops resulted in faster decomposition of SOC, depleting the soil carbon reserves. Therefore, the incorporation of high-value cash crops into the rotation, in which either relatively little biomass is produced or most of the biomass is removed from the field, do not fit well in rice rotations in terms of the enrichment of soil quality.

Besides putting good crops in rotation, letting rice fields go fallow is a better way to improve soil fertility than growing high-risk rice crops in dry seasons or areas, even though the soil nutrition benefits are less than when legume species are added (Becker & Johnson, 1998). This can effectively lower production costs and significantly improve soil fertility and biological properties. However, many studies tended to focus on the benefits of crop rotation or fallow in terms of soil fertility, ignoring the diversity of systems, socio-economic characteristics, weeds, and farmers' perceptions and production objectives (Becker & Johnson, 1998). For these reasons, the adoption of fallow as a viable technique by rice farmers in nutrient-limited soils is less successful unless drought becomes a barrier for crop cultivation (Table 1). Thus, future endeavours should consider socio-economic, climatological and ecological concerns when introducing different crops or fallow to rice rotations.

Tillage practices

Minimum tillage can improve SOC and nutritional status compared to puddled rice culture (Table 1). The primary benefits of puddling are the creation of a soft seedbed, reduction of water and nutrient losses and weed control (So *et al.*, 2001). Moreover, deep ploughing could be advantageous in certain systems as a short-term strategy to bring the leached nutrients to top soil layers while enhancing the benefits mentioned above. However, leaching would be negligible when the soil is inherently low in nutrients. Moreover, excessive tillage in continuous rice culture destroys soil structure,

disrupts the continuity of soil pores, reduces the number of residues on the soil surface and degrades soil quality (Osunbitan et al., 2005; Jiang & Xie, 2009). Therefore, puddling disrupts most of these soil qualities. Though the short-term agronomic response to tillage may be either negative (Agboola, 1981; Lal et al., 1989) or positive (Awadhwal & Smith, 1989; Kawakye & Bobo, 1995), most of the long-term crop responses are either neutral or negative (Kirchhof et al., 2000; So et al., 2001). Longterm field experiments conducted in different countries indicate improvements in soil aggregation, higher microporosity and SOC content, and lower bulk density under no-till, compared with conventional tillage systems (Andrade et al., 2010; Das et al., 2014). Moreover, combining minimum tillage and organic residues with reduced synthetic fertiliser application has increased soil nutritional status (Singh & Singh, 1995). For example, instead of conventional land preparation and application of high doses of N fertiliser (120 kg ha⁻¹), a reducedtill system with the incorporation of straw from the previous season and reduced N-application (60 kg ha⁻¹) was shown to improve the SOC pool in Benin (Dossou-Yovo et al., 2016). Therefore, practising reduced-tillage or no-tillage systems has been suggested as a potential approach to reducing the rates of decline of SOC and nutrient stocks (Timsina & Connor, 2001; Gami et al., 2009; Jiang & Xie, 2009; Das et al., 2014; Du et al., 2015). However, the effectiveness of no-till rice culture in improving SOC and nutritional status is site-specific. It depends on region, climatic condition (season), cropping system, rate and time of fertiliser application, rice-crop establishment method (i.e., transplanting, seedling throwing or direct seeding), and duration of notill adoption (Gupta & Seth, 2007; Ghimire et al., 2012; Suriyagoda et al., 2014; Huang et al., 2015) (Table 1). Despite the influence of many factors, minimum tillage shows promise in improving long-term SOC in nutrientlimited soils. Thus site-specific packages, including minimum tillage, should be developed considering the major factors mentioned above.

Use of biochar

Biochar is the product of heating biomass in the absence of or in limited air to above 250 °C, a process called charring or pyrolysis (Lehmann and Joseph 2015). In some instances, the material properties of biochar may overlap with those of charcoal as an energy carrier. Still, many types of biochar do not easily burn, and are typically made to address soil issues. A certain level of organic carbon forms, called fused aromatic ring structures, similar to charcoal, is an important defining feature of biochar. These structures are formed during pyrolysis and are crucial to biochar properties with respect to mineralization or adsorption. Therefore, biochar is typically enriched in carbon and other mineral elements. The chemical properties of the organic carbon structure of biochar are fundamentally different from those of the material that the biochar was produced from, and it is depleted in oxygen and hydrogen. In contrast, the macro-morphological characteristics of biochar typically resemble those of the starting material, which means that it normally looks the same, apart from its black color (Lehmann and Joseph 2015).

Applying biochar to soil can improve soil carbon and fertility (Table 1). Biochar improved the saturated hydraulic conductivity of soil and the xylem sap flow of upland rice in Laos (Asai et al., 2009). Biochar application results in higher grain yield at sites with low P availability and improves the response to N and P chemical fertilisers (Munda et al., 2018; Bi et al., 2019). Rice straw or rice husk biochar application results in increasing grain yield by 8.5-11% or 1-24%, respectively, due to increased nutrient availability and SOC content (Liu et al., 2016; Munda et al., 2018). Similar results were observed by Bi et al. (2019). However, biochar application reduces leaf SPAD values, possibly due to adsorption of NO₂-N on to biochar and reduced N availability, indicating that biochar application without additional N fertiliser application could reduce grain yields in soils with a low N supply (Lehmann et al., 2003; Asai et al., 2009). Recently Wu et al. (2019) revealed that the combined application of biochar and vermicompost increased rice yield by 26.5-35.3% compared with biochar amendment alone. These limited results suggest that biochar application alone or in combination can improve productivity, but the effect of biochar application highly depends on soil fertility and fertiliser management (Haefele et al., 2011; Liu et al., 2019). Moreover, the effects are more evident in highly weathered and infertile tropical soils under upland conditions (Asai et al., 2009; Peng et al., 2011; Liu et al., 2016; Bi et al., 2019).

Biochar application has its own disadvantages. When considering the amount of biochar required, incorporation of large amounts and ranges are often recommended, *e.g.*, 2 - 50 t ha⁻¹ is recommended in Madagascar (Asai *et al.*, 2009; Raboin *et al.*, 2016; Wang *et al.*, 2018; Bi *et al.*, 2019) and farmers cannot afford to apply this amount. Moreover, biochar may bind certain mobile nutrients, making them less available for rice plants, thereby aggravating the nutrient limitations (Asai *et al.*, 2009; Noguera *et al.*, 2010; Pratiwi & Shinogi, 2016; Haefele *et al.*, 2011) (Table 1). Due to the presence of conflicting results with adding biochar, further testing is warranted before making recommendations.

Site-specific nutrient management

Uniform and/or high fertiliser applications across large regions likely result in low fertiliser-use efficiency of rice due to the heterogeneity of soil nutrient availability and crop uptake. One possibility to improve productivity and nutrient use efficiency in such situations is to use the site- or even field-specific nutrient and crop management (Dobermann et al., 1998, 2003; Gupta & Seth, 2007; Haefele & Konboon, 2009; Nhamo et al., 2014; Haefele et al., 2016). The site-specific nutrient management system has gained popularity in rice production in Africa and Asia (Dobermann et al., 2002; Buresh et al., 2010; Saito et al., 2015; Niang et al., 2017). Site-specific nutrient management may increase the grain yield of rice by 300-900 kg ha⁻¹, and N, P and K accumulation by 8-21 % compared with farmers' practice or grain obtained before practising the site-specific nutrient management package in Asia (Peng et al., 1996, 2006; Wang et al., 2001; Dobermann et al., 2002; Khurana et al., 2007; Sui et al., 2013). The amounts and types of nutrients required through site-specific nutrient management may vary across sites, with variable spatial and temporal scales depending on the degree and type of nutrient limitations (i.e., both macro- and micro-nutrients). Moreover, this approach explicitly recognises the need to efficiently utilise organic and inorganic nutrient sources (*i.e.*, IPNS) and minimise losses.

When synchronising nutrient demand and supply to rice plants, the potential use of slow-release fertilisers should receive much attention, particularly for nutrients such as N, as the losses are high, leading to low-use efficiency, and frequent applications are not practicable (Tang et al., 2007; Ye et al., 2013). Site-specific and slow-release synthetic fertiliser management strategies are possible only when farmers have access to them which is unlikely for most rice farmers in low-input agricultural systems. In such situations, the application of organic manure_with a site-specific focus should be considered. However, this requires constant testing of available organic materials for nutrients, nutrient releasing patterns, and deciding on application rates to specific fields, thus making it less attractive to extension officers and farmers.

In-season nutrient management

The nutrient-use efficiency can be increased by improved in-season nutrient management, *e.g.*, N-use efficiency can be increased by 30-40% by increasing the number of splits with lower rates (Cassman *et al.*, 2002; Dobermann *et al.*, 2002). This is particularly important for N, as the losses are relatively high but can only be reduced with higher labour requirements. Leaf-colour charts or chlorophyll meters are helpful in this, as it can determine the time and rate at which N fertiliser application is required to maximise the use efficiency, based on the depletion of leaf greenness (Hussain *et al.*, 2000; Singh *et al.*, 2002; Alam *et al.*, 2005; Gupta & Seth, 2007; Nhamo *et al.*, 2014). Experiments conducted in the Philippines revealed that the N fertiliser rate could be reduced by 30-45% without affecting the grain yield of rice when N is applied based on the leaf colour chart compared with the conventional practice (Hussain *et al.*, 2000). In Bangladesh, N application based on the leaf colour chart increased the grain yield by 0.1- 0.7 t ha⁻¹ compared with farmer's management (Alam *et al.*, 2005).

When leaf colour chart-based N application was combined with site-specific nutrient management for other nutrients (*i.e.*, K, P, S and Zn), further improvement in grain yield by 0.4 t ha⁻¹ was observed (Alam *et al.*, 2005). Despite the benefits of using leaf colour charts and chlorophyll meters in enhancing the N-use efficiency, they are not available for nutrients other than N, and assume that other nutrients are not limiting. Moreover, those charts are not explanatory enough to less educated farmers and thus have limited applicability. The reasons hindering simple and promising techniques should be understood when managing in-season N nutrition efficiently.

Use of microorganisms

Integrated nutrient management with biological and chemical fertilisers can improve rice crop productivity, bio-fortification, soil health, and fertility (Hoseinzade et al., 2016). Contribution from biological forms of fertilisation would be either through the enrichment of nutrient pools (e.g., N) or by increasing the bioavailability of fixed forms of nutrients present in the soil (e.g., P). Arbuscular mycorrhizal fungi (AMF) isolated from rice and non-rice soils increased plant growth and P and Zn nutrition of wetland rice (Khan & Belik, 1995; Roy & Srivastava, 2013). However, the contribution from AMF is usually more significant in aerobic/upland soil conditions. Hoseinzade et al. (2016) evaluated the combined effects of biological fertilisers, including AMF (Glomus mosseae Nicol. & Gerd.), free-living N-fixing bacteria (Herbaspirillum seropedicae), and chemical fertilisers on yield, nutrient concentrations, and contents of wetland rice. Chemical and biological fertilisation interactions resulted in significant improvement in grain Zn, Fe, P, and N, and soil Fe, K, and N nutrition. According to Hoseinzade et al. (2016), the most excellent rates of grain and straw nutrient accumulation and soil nutrient enrichment result from combined use of biological and chemical fertilisers. Moreover, Jeong *et al.* (2015) found that arbuscular mycorrhizal symbiosis is established after four weeks of growth under aerobic soil conditions, and P nutrition of rice was improved. Therefore, introducing beneficial AMF to the soil, and/ or adopting crop management practices that enhance the activity and contribution of AMF, such as aerobic rice culture, may enhance soil fertility and plant nutrition.

Large amounts of N derived from biological fixation have been reported in rice fields, and this contribution can be as much as 50 kg N ha⁻¹ (Cassman et al., 1998). According to laboratory and greenhouse experiments, a significant improvement in rice N has resulted from Herbaspirillum seropedicae (Baldani et al., 2000; Roy & Srivastava, 2013). In a different experiment, 40% N and 25% P chemical fertiliser could be supplemented by the low-cost, natural resource-based bio-fertiliser (Azotobacter sp. and phosphate-solubilising bacteria Bacillus polymyxa) at 12 kg ha⁻¹ and organic manure at 10 t ha⁻¹, possibly enabling long-term sustainability in rice cultivation (Mukhopadhyay et al., 2013). In a separate experiment, Prasanna et al. (2015) tested the effect of cyanobacterial inoculation on the growth and yield of rice while reducing the application of synthetic fertiliser N. The treatments Anabaena-Nostoc consortium, and Anabaena-Trichoderma biofilm Trichoderma, recorded the highest values of grain and straw yield, which are on par with those of plots with recommended doses of synthetic fertilisers. Therefore, the benefits of cyanobacteria in improving nutrient availability, soil fertility, and crop productivity, with savings of 60 kg N ha⁻¹ season⁻¹ from synthetic fertilizer application, seem promising. Cyanobacterial inoculation increases grain yields by 10-24% compared to the non-inoculated control in different rice-cropping systems (De Caire et al., 2000; Prasanna et al., 2012). Moreover, the presence, activity, and associations between microorganisms, and thereby the benefits of soil microorganisms to rice plants, is affected by the duration and intensity of inorganic and organic fertiliser management (Li et al., 2019; Zhou et al., 2020). Biological fertilisers can save at least a fraction of recommended chemical fertilisers and improve the growth and yield of rice, especially if combined with the appropriate rate and type of fertilisers, by enhancing nutrient uptake.

In summary, to date, the general trend in rice cultivation is a decreasing use of organic and biological nutrient sources, mainly because of increasing opportunity costs of labour needed for collection, preparation, and application, and increasing use of synthetic fertilisers when available and affordable. As opportunities are limited for breeding varieties that acquire more nutrients from the soil or have higher internal nutrientuse efficiencies, long-term management strategies must focus on improving soil fertility by applying alternative sources of nutrients, including better management of residues. In this context, improving SOC and advocating for site-specific and integrated nutrient management are promising options, as the application and management of these practices are simple and cost-effective. Moreover, bio-fertilisers act synergistically to improve soil fertility and grain yield of rice. The combined use of nutrient management strategies under the context of cultural and socio-economic conditions is also recognised as the IPNS.

INSTITUTIONAL SUPPORT AND EDUCATION OF FARMERS

Agronomic interventions alone are insufficient to ensure improved productivity and sustainability in rice fields. Improving productivity in nutrient-limited rice fields require collective actions at both field and regional scales. At the field level, there is a need for regular evaluation of soil fertility to make decisions on nutrient management, aiming to lower the cost of inputs and improve the soil fertility status. Challenges remain, as the transfer of technological and intuitive knowledge is restricted by the lack of routine farm-level soil and crop monitoring procedures. Thus, there is a requirement to focus on agronomic advice and support at the local farm and regional scales (Rahman & Parkinson, 2007; Stewart et al., 2020). As a long-term strategy, nutrient limitations/availability heterogeneity can be mapped, along with other soil limitations, at field scale. Once mapped, breeding priorities can be identified in specific areas.

Moreover, simple decision-support tools can be developed using farmers' knowledge of their fields and planned crops to determine the best nutrient management package. When considering such site-specific nutrient management approaches, the major challenge is to reduce the complexity of new technologies disseminated to farmers. Site-specific nutrient management may vary in temporal and spatial scales, and farmers are not adopting new technologies at a fast rate (Dobermann *et al.*, 2002, 2003; Haefele & Konboon, 2009). Therefore, continuous guidance and directions should be provided to farmers until they become independent in using these technologies and improve soil fertility.

Apart from the field-level factors mentioned above, other major constraints to rice production at the regional scale are weed infestation, variable rainfall patterns, low and under-developed irrigation infrastructure, and socioeconomic, institutional, and political considerations (i.e., lack of financial resources, labour shortages, low levels of education, weak infrastructure, lack of conducive policies) (Stewart et al., 2020). Solving these constraints along with the improvement of productivity of nutrient-limited rice fields to bridge the large existing gap between current and the potential productivity is necessary (Nhamo et al., 2014; Stewart et al., 2020). Farmers need access to credits and markets to follow up on recommendations (Nhamo et al., 2014). Subsidies, improved agro-dealer networks, and re-packaging fertilisers and/or manures into smaller portions are measures used to improve fertiliser and/or manure use by resource-poor farmers (Poulton et al., 2006). However, the success of such interventions has been low except for some models such as the OneAcre Fund in Africa and the track (Yaya) approach in Sri Lanka. The successful collective decision-making strategy adopted in the Yaya approach scheme, when making in-farm and off-farm decisions, also contributed to the Yaya approach success.

Agronomic planning (e.g., recycling/importing crop residues or other forms of organic manure and selection of crops in rotation among the farming community) should be carried out with the participation of stakeholders (Gupta & Seth, 2007; Rodenburg et al., 2014; Stewart et al., 2020). This is perhaps the most challenging part of the approach, as getting all stakeholders together has proven difficult in many situations (Warner, 2006). Reaching consensus among a wide range of stakeholders with different interests might be another considerable hurdle. The outcome of such multi-stakeholder platform processes should aim for a compromise between economic, social, and environmental gains (Rodenburg et al., 2014). Therefore, improving the productivity in nutrient-limited rice fields cannot be achieved only through agronomic means, and success depends on the input and commitment made by all stakeholders.

CONCLUDING REMARKS

The increase in productivity of rice lands has slowed down or even declined in the long term in most rice cultivating systems in Africa and Asia. Declining soil fertility is identified as one of the major causes for this. Currently, over two-thirds of the total rice area on the globe is classified as soil with nutrient deficiencies or toxicities (only in localised areas), and yield gaps over 5 t ha⁻¹ have been reported from many of these systems. Widely reported soil fertility constraints are deficiencies of N, available P, S, and Zn, and Fe²⁺ toxicity. In general, the occurrence of nutrient deficiencies decreases in the sequence of uplands > rainfed lowlands > irrigated lowlands > deepwater areas.

Most of the global rice breeding programs are focused on improving productivity in high potential areas under optimal management. Additionally, numerous efforts have been made to breed rice varieties for pest and disease tolerance, flood tolerance, salinity tolerance, iron toxicity, and drought tolerance. Despite these breeding efforts, most rice farmers in Africa and Asia continue to grow rice in nutrient-deficient field conditions, mainly due to their poor economic status, lack of awareness, and/ or lack of access to synthetic fertilisers. This problem is likely to be aggravated further. Thus, optimal management practices recommended for nutrient unlimited soils are not applicable, and the varieties developed for those conditions are not adoptable to nutrient-limited grounds. In order to sustain rice cultivation in nutrient-limited soils, soil fertility constrains must be identified and addressed without prioritising synthetic fertiliser application as a potential option. In addition, parallel implementation of other technologies and policies must be considered. The introduction of credit facilities to farmers, knowledge and technology in weed, pest, and disease management, the selection of suitable varieties, water management, the use of quality seeds, exposure to competitive markets, and IPNS are examples.

Improving SOC and nutrient pools through incorporating organic manure or farm yard manure should be practised without allowing the amount of nutrients released by organic sources to exceed crop nutrient demand to minimise nutrient losses. Apart from that, the practice of fallow or crop rotations, minimal or zero tillage practices, use of site-specific and intensive in-season nutrient management, and introducing and promotion of the activity of microorganisms fixing nutrients, such as bacteria, cyanobacteria and fungi, are promising soil carbon and nutrient improvement strategies. This integrated approach of the above components is recognised as IPNS. Depending on the availability of the above materials, institutional support and socio-economic status, farmers need to be educated to improve the soil fertility status gradually or at the least avoid further decline in soil fertility. The time required to replenish the soil fertility will depend on the type and severity of nutrient limitation, and the balance between nutrient input and output processes. Due to the

complex nature of the soil, and the socio-economic and institutional structures in these regions, an integrated and participatory approach, including all the stakeholders, is required to implement and sustain the agronomic interventions suggested above.

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REVIEW

Food Safety Heavy metals and food safety in Sri Lanka: A review

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Summary: Entry of heavy metals to the food chain leads to food safety hazards. The origins of possible food safety hazards in Sri Lanka due to metalloid arsenic and the heavy metals cadmium, lead, and mercury are reviewed. Of them, arsenic and cadmium draw attention as contaminants in rice. Of the four heavy metals, cadmium in agricultural soils is of anthropogenic origin. Arsenic is of lithogenic origin. In some locations lead appears to be of anthropogenic origin, especially in commercial leafy vegetable cultivating soils. Marine fish, particularly swordfish and yellow fin tuna, occasionally carry cadmium and mercury concentrations above the tolerance limits established by the Codex and European Food Safety Authority. Heavy metals in well water are far below tolerance limits and are safe. Patterns of annual cancer incidences in Sri Lanka do not provide evidence to consider arsenic as a food safety hazard. Food safety hazards may occur with arsenic in the long term if attention is not paid to the quality of fertilizers or the current daily rice consumption level is not reduced. Arsenic being of lithogenic origin, unhealthy exposures cannot be prevented without affecting the food security of the country. High consumption of cadmium containing rice exposes Sri Lankans to health problems. Signs of hotspots of lead are visible. Food safety hazards are predicted by assessing exposure of humans based on their body weight and daily intake of hazardous constituents. Provincial Tolerable Weekly Intakes (PTWI) are calculated for Sri Lanka using information on heavy metals in foods from research publications. International food regulatory limits on heavy metals in foods are summarized. Horizontal standards for heavy metals in foods are developed to minimize food safety hazards in Sri Lanka.

Keywords: Arsenic, cadmium, food chain, lead, mercury.

INTRODUCTION

Entry of harmful agents through the food chain into the human body results in food safety hazards. Heavy metals have drawn scientific attention as a potential food safety hazard entering the human food chain. Of the heavy metals in the soils and the environment, cadmium (Cd), lead (Pb), mercury (Hg), and the metalloid arsenic (As) are of concern globally. Heavy metals are also described as toxic trace metals in the literature. The exposure of humans to heavy metals is linked to their presence in foods, as free metal particles, inorganic compounds, or organic compounds. Exposure of humans to heavy metals in agri-foods depends on the food processing, preparation, and consumption patterns associated with food cultures. Heavy metals of lithogenic origin may be naturally present in the soil and water, or get added to agricultural soils through anthropogenic activities, especially through fertilizers and pesticides. Heavy metals also may enter the agri-foods from organic waste, compost, animal dung, sewage sludges, irrigation water etc. used as manure. There are no mining activities, volcanic emissions, or glacial activities in Sri Lanka that could bring heavy metals from the core of the earth to surface soils. Sri Lankan agriculture depends totally on rain fed and irrigated water. Water from deep aquifers which may carry high concentrations of heavy metals are not used in agriculture. Deep well water in Sri Lanka is free of heavy metals. Heavy metals in foods is suspected

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as one of the probable causes of chronic diseases of unknown aetiology (CKDU) among Sri Lankans. Similar diseases are reported in a few other countries. Understanding the presence of heavy metals in the agronomic soils, their appearance in foods, and causeeffect relationship with chronic diseases in the country need in-depth multidisciplinary scientific studies. This publication examines the presence of heavy metals in Sri Lankan agricultural soils and food production environment, their probable origins, opportunities for them to enter the food chain, the Sri Lankan food consumption patterns and incidences of related effects on health. This review proposes regulatory standards to minimize food safety hazards. The focus is on arsenic, cadmium, and lead entering the food chain during crop production and mercury entering through marine fish.

HEAVY METALS IN AGRICULTURE AND FOOD SAFETY

Heavy metals present naturally in varying are concentrations in soils. They are distributed heterogeneously in the Earth's crust, resulting in differing views of their presence, and mapping of heavy metal hotspots. The heavy metal concentrations the Earth's crust is different from that of the core. The heavy metal concentrations on the crust change continuously due to environmental, agronomic, and natural factors. The suitability of soils for cultivation are decided based on the "threshold values" of heavy metals, above which there is a risk of heavy metals entering food crops. The aim is to ensure public health through a safe food supply. The average concentrations of heavy metals in the Earth's crust, their threshold values based on information from European and Indian studies, and those established by Finland, are presented in Table 1 (Ministry of Environment, Finland, 2007).

Most heavy metals of concern in Sri Lanka have been established to arise mainly from weathering of rocks and soils and not from agrochemical contamination of soil surfaces or from deep well water (Jayawardene *et al.*, 2012). The homogeneity of arsenic in Sri Lankan paddy soils is evident from the presence of similar concentrations in 70 locations of 14 villages (Chandrajith *et al.*, 2005). In a study comparing the trace metals in Sri Lankan soils from agricultural and non-agricultural highlands and lowlands of the Anuradhapura district by principal component analysis, calculating the geo-accumulation index and pollution loading index, it is established that copper, nickel, zinc, and lead in agricultural soils are of lithogenic origin and cadmium is of anthropogenic origin (Sanjeevani *et al.*, 2017). Cadmium is one of the problematic heavy metals in human food chains in many countries.

Entry of heavy metals into the food chain

Every atom of a heavy metal or every molecule containing a heavy metal in soil does not end up in the food plate. The physical properties of the soil, the ability of soil particles to absorb, desorb or bind heavy metals, the redox potential of soils governed by oxygen availability, the soil pH, and biological properties of the plants govern the entry of heavy metals into plants. Redox potentials of paddy soils have opposing effects on the absorption of arsenic and cadmium by plants, whereas soil pH affects the bioavailability of arsenic and cadmium (Zhao & Wang, 2019). It is thus challenging to work towards food safety through adjustment of soil conditions. The heavy metals should be available in the soil at depths down to 30 cm on average, for them to be absorbed by food crops. The depth of extension of roots in the soils limit the opportunity for the heavy metals to enter plants. The plants are selective in up-taking heavy metals and in translocating them to the edible components of crops. These complexities make it difficult to scientifically predict the extent of movement of heavy metals from soil to food of plant origin. The entry of each toxic heavy metal into the human food chain needs to be understood in working towards food safety.

Table 1: Abundance of As, Cd, Pb and Hg in the Earth's crust and their threshold values (mg/kg)

Distribution	As	Cd	Pb	Hg	References
Average on Earth's crust*	1.8**	0.15	10	0.05	https://en.wikipedia.Earth%crust
Natural abundance	8.93	0.48	29.9	0.13	Arunakumara et al., 2013
Thresholds for agricultural soils	5	1	60	0.5	Govt. Decree 214.2007.doc (finlex.fi) of Finland
After sewage sludge applications in agriculture	50	3	300	1	EC Directive 86/278/EEC

*The values vary in different publications; **Range 0.1 - 40 mg/kg

Arsenic in Sri Lankan food chain

The average concentration of arsenic in the earth crust is 1.8 mg/kg with a possible range of 0.1 - 40 mg/kg in non-contaminated soils. The threshold value for arsenic in agricultural soils is 5 mg/kg. Arsenic could be present in 200 different forms in soil of which 60% are arsenates, 20% are arsenites, oxides, arsenides, silicates etc., and 20% as sulphosalts and sulphides (Lim *et al.*, 2014). Arsenites are more toxic than arsenates. Out of these, only a few forms of arsenic enter the rice grain. Though total arsenic in foods was used in assessing food safety hazards in the past, current toxicological studies focus on inorganic arsenic in foods. Of the total arsenic content in rice, 60-80% is inorganic arsenic which may increase up to 90% (Jose *et al.*, 2009).

Organic arsenic, mainly present as arsenobetaine, forming more than 90% of total arsenic in fish, pose no recognizable threat to human health. Organic arsenic is not absorbed in the human gut. No regulatory limits are established for inorganic arsenic in fish in the food regulatory system in developed countries, as its presence is low and negligible (Codex, 2015; EC- 2006; FDA, 2021).

The food safety hazards through arsenic in rice occurs due to following reasons.

- 1. High volumes of rice consumed by Sri Lankans make rice a major entry pathway.
- 2. Arsenic exists in rice as the toxic inorganic form constituting 60 90% of total arsenic.

- Removal of arsenic in rice is limited to polishing and washing prior to cooking. Each washing takes away 15% of arsenic in rice.
- 4. There is no mechanism to avoid food safety hazards from rice cultivated in areas where arsenic content in soil or irrigation water is naturally high.
- 5. Reduction of arsenic regulatory limit of 0.2 mg/kg for rice to 0.1 mg/kg would eliminate 70% of rice in the market leading to food security problems culminating in undernutrition of the populations (FDA, 2016).

Absorption of arsenic by plants from soil and its accumulation finally in the edible tissues occur along a concentration gradient (Jose et al., 2009). With 7.19 -18.63 mg/kg of arsenic in rice plant roots in Bangladesh, the observed arsenic content in rice grains was 0.25 - 0.73 mg/kg (Bhattacharya et al., 2009). In rice plants, the arsenic content in roots is reported to be 28-fold higher than in the shoot, and 75-fold higher than raw rice grains (Raman et al., 2002). Approximately 1% of arsenic in the soil ends up in the rice grains. Raman et al., (2002) postulate that rice in soils containing less than 14 mg/kg of arsenic could be considered safe for human consumption. West Bengal in India and Bangladesh continue to face food safety hazards due to higher concentrations of arsenic in rice and water. The main source of arsenic in Bangladesh is the deep well water used for irrigation, resulting in contamination of the rice soils over years. A comparison of total arsenic concentrations in soil, water, and rice from West Bengal, India, where severe exposure of populations to arsenic occurs, with that of Sri Lanka are given in Table 2.

Location	Soil arsenic (mg/kg)	Irrigated water arsenic ^a (µg/L)	Rice grain arsenic ^b (mg/kg)	References
West Bengal	1.38 - 12.27	110 - 760	0.25 - 0.73	Battacharya et al., 2009
Dry zone Sri Lanka	0.45 - 1.04 Dry zone; 0.5 - 24 (tank sediments)	0.015 - 0.361 Nikawewa, Girandurukotte	0.09 - 0.26 Girandurukotte, Nikawewa	Chandrajith <i>et al.</i> , 2011 Chandrajith <i>et al.</i> , 2005 Chandrajith <i>et al.</i> , 2008
Dry zone paddy soils	1.18 ± 0.59			Balasooriya et al, 2021
Wet zone paddy soils	1.32 ± 0.85			Balasooriya et al, 2021
CKDU hotspot soils	1.33 ± 0.60			Balasooriya et.al, 2021

a = FAO permissible limit for irrigation water is 0.10 mg/L (= 100 μ g /L); Value for drinking water is 10 μ g /L; b = Codex limit for arsenic in rice is 0.2 mg/kg

Arsenic in Sri Lankan soils are reported to originate from sulphide minerals of the basement rocks and the concentrations are insignificant to affect plant growth (Jayawardena, 2012). The concentrations of arsenic in well water in Sri Lanka are far below the internationally accepted maximum tolerance limit of 10 μ g/L (Table 22). The concentrations are reported by Herath *et al.*, 2018 (Table 3) and in the WHO report on CKDU, 2016 (Figure 1).

Table 3: Average As, Cd and Pb concentrations in well waters from the 25 districts in Sri Lanka (µg/L) (Herath *et al.*, 2018) Values are computed from district averages

Metal	Country average	SD	Range	MTL (µg/L)	Below MTL	High district averages \pm SD & (maximum) $\mu g/L$
Total arsenic	1.38	2.68	0 - 66	10	21/25 districts	Batticaloa = 3 ± 3.2 (14) Mannar = 7 ± 11.7 (66) Mullativu = 3 ± 3.7 (13) Puttalam = 4 ± 4.1 (15)
Cadmium	0.008	0.007	0 - 0.05	3	All	
Lead	0.133	0.155	0 - 0.5	50	All	Galle: one sample had 228

Concentrations of 0.5 - 24 mg/kg of arsenic and 10 - 33 mg/kg of lead have been reported from the sediments at different levels in Malagane Tank, Deduruoya. The values probably reflect accumulation of arsenic from weathering rocks (Chandrajith, 2008). These concentrations do not reflect alarming anthropogenic contributions from paddy cultivations in the surroundings.

In pot experiments with arsenic concentrations of 60 mg/kg in soil, the corresponding arsenic concentrations in cultivated rice plants were 21 mg/kg in panicles at initiation stage and 23 mg/ kg in the panicles at maturity stage (Rahman *et al.*, 2008). At these artificially high exposure levels to soil arsenic, less than 50% appears to enter the rice grains. The information in Table 2 indicates that the exposure of rice plants to arsenic through soils

Table 4: Average As, Cd, and Pb in rice from CKDU endemic and non-endemic areas (mg/kg) (Herath et al., 2018)

Heavy metal [Sample number]	Average \pm SD	Maximum	Average \pm SD	Maximum	Area related to CKDU	Codex mg/kg
	Polishe	ed rice		Unpolished rice		
Total arsenic [67]	0.03 ± 0.04	0.20	0.03 ± 0.04	0.20	Endemic	0.2
Cadmium [67]	0.12 ± 0.19	0.87	0.16 ± 0.17	0.52	Endemic	0.4
Lead [67]	0.01 ± 0.02	0.08	0.00 ± 0.02	0.08	Endemic	0.2
Total arsenic [24]	0.03 ± 0.03	0.00	0.04 ± 0.03	0.09	Non endemic	0.2
Cadmium [24]	0.21 ± 0.24	0.65	0.18 ± 0.36	1.43	Non endemic	0.4
Lead [24]	0.00 ± 0.00	0.00	0.00 ± 0.00	0.02	Non endemic	0.2

Table 5: As, Cd and Hg in Sri Lankan rice (mg/kg) (Jayasekara & Fretas, 2005)

Heavy metals	Raw polis	hed grains	Parboile	ed grains	Rice flour	
	Producer 1	Producer 2	Producer 1	Producer 2	Producer 1	Producer 2
Total arsenic	0.034 ± 0.006	0.034 ± 0.001	0.065 ± 0.012	0.092 ± 0.001	0.035 ± 0.001	0.061 ± 0.006
Cadmium	0.192	< 0.5	< 0.5	< 0.5	< 0.5	< 0.5
Mercury	< 0.01	< 0.01	0.033	< 0.01	< 0.01	< 0.01

[Codex tolerance limits for rice As = 0.2, Cd = 0.4 mg/kg]

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in Sri Lanka is far below that of the pot experiments and in West Bengal. The soils and agronomic practices in Sri Lanka do not seem to result in entry of hazardous arsenic concentrations into the rice production-processing chain. Independent studies reported in Tables 4 and 5 by different groups of scientists support the view that rice arsenic concentrations cannot cause food safety hazards in Sri Lanka as the concentrations are well below Codex maximum tolerance limits.

The presence of arsenic in rice is a global phenomenon. Rice arsenic concentrations in USA and in several countries are given in Tables 6 and 7. A study of 901 polished white rice samples from 10 countries has shown 7-fold variation from the median. The lowest values were 0.04 mg/kg in Egypt and 0.07 mg/kg in India. The highest values were 0.25 mg/kg in USA and 0.28 mg/kg in France (Meharg *et al.*, 2009).

The arsenic concentrations in rice in Sri Lanka (Tables 2 and 4) are notably less than what is in Tables 6 and 7 indicating a low exposure. The arsenic concentrations reported in Tables 2 and 4 are less than the Codex tolerance limit of 0.2 mg/kg.

Correlation of the information on arsenic in the WHO study (2016) in Sri Lanka is given in Figure 1. The figure presents the food chain in blue boxes and arrows. The values given above the food chain (in red) indicate potential food safety hazards and outcomes associated with arsenic. The values given below the food chain (in green) represent values that do not point towards food safety hazards.

The evidence does not suggest a food safety hazard or health effects on humans arising from arsenic in agricultural soils or water. A long-term risk may arise from pesticides and excessive use of phosphate fertilizers containing arsenic on continuous application. The long-term effects are generally predicted by examining the exposure of agricultural lands to contaminated fertilizers over a period of 45 years. Arsenic available in added phosphate fertilizers may get diluted in the soil allowing only a fraction to reach rice as discussed by Raman *et al.*, (2002). The pesticides along with arsenic in them are absorbed through leaves and panicles leading to increased risk. However, the current evidence does not indicate food chain (rice and water) as a pathway for exposure of humans to arsenic in Sri Lanka.

Table 6: Concentrations of As in rice products in USA (Meharg et al., 2009)

No	Rice type	No of samples	Mean (mg/kg)	SD (mg/kg)	Range (mg/kg)
1	All samples tested	193	0.194	0.144	0.006 - 0.723
2	Rice (non-organic)	88	0.205	0.122	0.047 - 0.559
3	Rice (organic)	13	0.174	0.142	0.086 - 0.526
4	Rice products (non- organic)	67	0.214	0.171	0.010 - 0.723
5	Rice products (organic)	25	0.125	0.142	0.006 - 0.620

Table 7: Concentrations of As in rice from several countries (Duxbury & Zavala, 2005)

Country	No. of samples	Arsenic (mg/ kg)	Country	No. of samples	Arsenic (mg/kg)
Argentina	1	0.136	Korea	2	0.045
Bangladesh	3	0.046	Lebanon	1	0.169
Bhutan	1	0.032	Pakistan	3	0.033
China	2	0.146	Spain	2	0.186
Egypt	2	0.032	Thailand	9	0.093
Greece	1	0.114	USA	22	0.181
India	16	0.037	Venezuela	12	0.084
Italy	7	0.158	TOTAL	84	Mean 0.107



Figure 1: Concentrations of As linked to agronomic practices, appearing in food chain and in tissues of persons affected by CKDU.

While precaution is important, haste does not seem to be scientifically justifiable in assessing food safety hazards.

The major concern that may arise in Sri Lanka is the increased exposure to arsenic due to higher consumption of rice compared to developed nations, rather than high concentrations of arsenic in rice. The comparable per capita consumption values for rice in kg are Bangladesh 268; Vietnam 220; Thailand 178; Sri Lanka 160; China 125; India 98; Australia 13; USA 10; UK 8 and World average of 80 (Helgi library, 2017). The effects of volumes of rice consumed is addressed in Table 17.

Among other food sources, concerns about arsenic in canned fish has been expressed at times. Assessment of weekly, monthly or daily exposure to arsenic in foods is used in recognizing risks leading to food safety hazards. The average consumption of rice by a Sri Lankan is 438 g per day. The corresponding value for fish in Gampaha district, which get plenty of fish from Negombo is 43 g per day (Jayasinghe et al., 2018). With a 10-fold lower consumption of fish than rice, there is extremely low possibility of food safety risks through arsenic in fish. The arsenic in fish is 90% arsenobetaine and other organic forms which are non-toxic. Organic arsenic is not absorbed by the human gut. The European Food Safety Authority (EFSA), Joint Expert Committee of FAO/WHO (JECFA) guidelines used by Codex Alimentarius Commission (Codex), and the Food and Drugs Administration of USA (FDA) have not established regulatory limits for arsenic in fish, considering the insignificant contribution of toxic

inorganic arsenic in fish to human diet. Arsenic in fish, fruits and vegetables are 90% organic (Mandel & Suzuki, 2005).

Cadmium in Sri Lankan food chain

Cadmium could enter the human body through food and water. The average concentration of cadmium in the Earth's crust is 0.15 mg/kg. The threshold value for agricultural soils is 1 mg/kg (Table 1). Cadmium compounds contaminate agricultural soils from sewage, manure, and phosphate fertilizers. Soils naturally contain low concentrations of cadmium. Gunadasa *et al.*, (2021) reported that the cadmium and arsenic concentrations in paddy soils and in rice are less than 0.1 mg/kg. The soil cadmium concentrations observed in uncultivated soils in Sri Lanka is 25 - 50% below the threshold value (Sanjeevani *et al.*, 2017) and vegetable soils reach threshold values (Sanjeevani *et al.*, 2017) and vegetable soils reach threshold values (Table 8).

Rice, potatoes, grains, and vegetables tend to absorb cadmium from soil more readily than other plants. Green vegetables appear to possess a tendency to accumulate cadmium. Unacceptable cadmium concentrations have been reported in sword fish from most oceans. Entry of cadmium to plants is reported to be 2-fold higher than arsenic. Cadmium accumulate in kidneys over long periods, and its half-life in the human body is around 30 - 38 years, and is 10 years in the kidney. Cadmium is a Group 1 carcinogen affecting mainly the kidneys, followed by bones (IARC, 2018).

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Area	Sub area	No. fields	Cadmium (mg/kg)	Lead (mg/kg)
Low country	Sedawatta	4	0.61-3.28	39-113
	Welewatta	3	0.46-1.37	34-66
	Kotuwilla	3	0.98-1.31	20-56
	Kahathuduwa	10	0.49-1.55	17-33
	Bandaragama	2	0.53-0.89	15-15
	$Mean \pm SD$		1.18 ± 0.82	54 ± 29
	Control soil		0.26	49
Up country	Sithaeliya	4	0.51-0.88	56-311
	Kandapola	6	0.39-1.96	27-97
	Haputale	3	0.51-3.86	26-242
	Bogahakumbura	3	1.30-1.42	45-75
	Rahangala	2	1.22-1.29	97-116
	$Mean \pm SD$		1.21 ± 0.80	88 ± 75
	Control soil		0.51	40

Table 8: Cd and Pb in low-country and up-country vegetable soils (Premarathna et al., 2011)

Concentrations of cadmium observed in water and rice in Sri Lanka are below the Codex tolerance limit of 3 μ g/L and 0.4 mg/kg respectively (Tables 3, 4, and 5). Meharg *et al.* (2013) reported the highest quantities of rice consumption in Sri Lanka and Bangladesh, among 12 countries, and the concentration of cadmium is the highest in Sri Lankan rice. The mean cadmium concentration

reported by them is 0.081 ± 0.024 mg/kg, with the highest value of 0.80 mg/kg in the range for Sri Lanka. Though the concentration of cadmium in Sri Lankan rice is below the Codex tolerance limits (Table 9), the exposure of Sri Lankans appears high. The study has estimated the exposure of Sri Lankans and Bangladeshis to be 100 µg/kg of body weight per week.

 Table 9:
 Concentrations of Cd in market rice and projected maximum weekly exposures in several countries including Sri Lanka. (Meharg *et al.*, 2013)

Country	Number of samples	Mean (mg/kg)	Range (mg/kg)	Max weekly Cd intake in µg/kg body weight
Bangladesh	260	0.099	< 0.0005 - 1.31	100
Cambodia	14	0.006	0.0010 - 0.03	5
France	37	0.010	0.0030 - 0.10	1
Ghana	428	0.020	< 0.005 - 0.27	10
India	58	0.078	0.0020 - 1.00	12
Italy	114	0.038	0.0030 - 0.16	10
Japan	18	0.059	0.0101 - 0.14	10
Nepal	12	0.050	0.0139 - 0.08	10
Spain	92	0.024	0.0008 - 0.14	10
Sri Lanka	75	0.081	< 0.0005 - 0.80	100
Thailand	18	0.027	0.0057 - 0.07	8
USA	21	0.018	0.0095 - 0.04	7

Leafy vegetables possess a higher tendency to accumulate cadmium than other plants (Westfall *et al.*, 2005; Gupta *et al.*, 2019). Vegetables from commercial plots with long term cultivations and vegetables available along the roadside on the Kesbewa – Kalutara road are reported (Premarathna, *et al.*, 2011; Kananke *et al.*, 2014) to carry cadmium concentrations above the Codex tolerance limit and EU tolerance limit of 0.2 mg/kg for leafy vegetables and 0.1 & 0.05 mg/kg for other vegetables (Table 10 and Figure 2).

Premarathna *et al.*, (2011) also reported high concentrations of cadmium in upcountry and low country vegetable soils, indicating the probable origin of cadmium in leafy vegetables (Table 8). The reported concentrations are 8-fold higher than the average cadmium concentrations in earth crust and 10-20% higher than the threshold values for agricultural soils indicated in table 1.

Area	Vegetable	Cd ^a (mg/kg)	Cd ^b (mg/kg)	Pb ^a (mg/kg)	Pb ^b (mg/kg)
Low country	Kankun	0.28 - 0.53	0.09 - 0.19	7.4 - 11.3	0.27 - 0.45
	Mukunuwenna	0.17 - 1.10	0.08 - 0.90	5.6 - 10.36	0.18 - 1.32
	Sarana	0.48 - 0.65		8.3 - 12.7	
	Niwithi	0.30 - 0.51	0.18 - 0.72	6.6 - 12.3	0.44 - 0.97
	Thampala	0.55 - 0.62	0.11 - 0.54	8.15 - 11.72	0.54 - 1.04
	Gotukola	0.54		8.75	
	Kohila		0.24 - 0.97		1.37 - 1.59
Up country	Potato	0.22 - 0.86		3.66 - 9.95	
	Lettuce	0.3		10.12	
	Cabbage	0.37 - 2.02		5.68 - 10.10	
	Leeks	0.48 - 0.54		3.84 - 10.45	
	Carrot	0.41 - 2.05		3.81 - 10.10	
	Knol-khol	1.28		7.65	

Table 10: Cd and Pb in low-country and up-country vegetables (Premarathna, et al., 2011^a; Kananke et al., 2014^b)



Figure 2: Concentrations of Cd linked to agronomic practices appearing in the food chain and in tissues of persons affected by CKDU.

Heavy metals and food safety

The concentrations of cadmium in the market triple super phosphate (TSP) from locations in Anuradhapura, Medawachchiya, Medirigiriya, Girandurukotte, and Kandy are reported to be above the acceptable limits and vary widely compared to other fertilizer components urea and Nitrogen – Phosphorus – Potassium mixtures (NPK) (Chandrajith *et al.*, 2010). The results are summarized in Table 11.

Reports on cadmium concentrations in other possible sources of fertilizers and manures clearly indicate the probable origin of cadmium toxicity in Sri Lankan foods leading to food safety hazards (Table 12).

The extremely heavy concentrations of cadmium in imported triple superphosphate, against local rock phosphate and other sources of phosphate fertilizers of

Table 11: Summary of Cd and Pb concentrations in market fertilizers in 6 locations of Sri Lank	a.
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Fertilizer	Cadmium (mg/kg)		Lead (mg/kg)		
	$Mean \pm SD$	Range	$Mean \pm SD$	Range	
Urea	0.3 ± 0.2	n.d - 0.4	4.2 ± 0.9	3.7 - 6.0	
NPK	0.5 ± 0.1	0.4 - 0.5	3.6 ± 0.5	2.6 - 3.9	
TSP	49.9 ± 29.2	39.8 - 80.2	79.2 ± 53.5	58.2 - 166.0	
Tolerance limits, Texas, USA	39		300		

Table 12: Cd and Pb in phosphate fertilizers and manures in Sri Lanka (Premarathna et al., 2011)

Fertilizer / manure	Number of samples	Cd (mg/kg)	Pb (mg/kg)
Triple super phosphate	5	23.50	5.15
Eppawala rock phosphate	4	1.92	13.00
Rock phosphate (USA)	2	13.54	12.00
Rock phosphate (India)	2	12.18	13.50
Rock phosphate (Bi Ru – China)	2	14.1	14.20
Rock phosphate (Lucille)	2	15.8	12.00
Dolomite	2	9.06	16.9
Poultry manure	12	0.97	3.20
Cattle manure	10	0.43	1.10

 Table 13:
 Pb, Cd, and As in household foods in CKDU endemic area (Ananda Jayalal *et al.*, 2019) and Codex tolerance limits

Food item	Number	As (mg/kg)	Cd (mg/kg)	Pb (mg/kg)
Rice	65	0.046	0.022	0.236
Flower vegetables	88	0.045	0.018	0.232
Leafy vegetables	43	0.066	0.019	0.212
Legume vegetables	19	0.020	0.019	0.185
Fruits	24	0.020	0.015	0.188
Fat and oil	6	0.020	0.027	0.154
Inland fish	2	0.020	0.015	0.057
Root & tuber crops	27	0.020	0.015	0.186
Codex MTL		0.20	0.40	0.20

Note: These foods represent farmer cultivations for their use

global origin, indicate the possible entry of cadmium at unsafe concentrations into the food chain in Sri Lanka. Contrary to the above observations on cadmium in foods, a study on heavy metals in household foods in farmer families indicated low arsenic and cadmium concentrations, and high lead concentrations (Table 13). The samples tested perhaps represent foods produced under controlled agronomic practices for family consumption by farmers.

Correlation of the information in the WHO study (2016) points towards unacceptable cadmium concentrations in soils, in vegetables, and the tissues from persons exposed in CKDU endemic and non-endemic areas (Figure 2).

Cadmium tends to accumulate in marine fish. A study on heavy metals in fish in Sri Lanka has revealed high cadmium concentrations in sword fish compared to other fish (Table 14). Sword fish accumulates more cadmium than others. The values also represent fish ready to be exported.

Exposure of Sri Lankans to cadmium appears to occur through several routes with a common anthropogenic origin. Sanjeevani *et al.* (2017) identified cadmium as the only toxic heavy metal of anthropogenic origin in agricultural soils from Anuradhapura district. The distribution pattern of cadmium in agricultural soils of CKDU endemic and non-endemic regions, in commercial vegetable plots and the market vegetables (in non-endemic areas) indicates high exposure of crops to cadmium leading to food safety hazards. Evidence on fertilizer analysis further confirms this origin. Cadmium, probably arising from imported TSP threatens food safety by burdening the soil and food with cadmium. Fish does not seem to carry cadmium concentrations above the tolerance limits.

Table 14: Summary of total Hg, Cd and Pb in sea fish in Sri Lanka

Fish (number of samples)	Hg (mg/kg)	Cd (mg/kg)	Pb (mg/kg)	Reference
Cooked sword fish (11)	nd – 1.47	nd - 0.029	-	Jayasinghe et al.,
Cooked yellowfin tuna (11)	nd-0.95	0.007 - 0.049	-	2018
Cooked sardinella (11)	nd-0.09	0.007 - 0.021	-	
Sword fish (35)	$\begin{array}{c} 1.24 \pm 0.72 \\ (0.20 - 2.58) \end{array}$	0.13 ± 0.08 (0.03 - 0.36)	$\begin{array}{c} 0.03 \pm 0.04 \\ (nd-0.15) \end{array}$	Jinadasa et al., 2010
Yellow fin tuna (25)	$\begin{array}{c} 0.39 \pm 0.19 \\ (0.14 - 0.88) \end{array}$	0.02 ± 0.02 (nd - 0.09)	$\begin{array}{c} 0.06 \pm 0.06 \\ (nd-0.24) \end{array}$	
Red snapper (12)	0.17 ± 0.06 (0.09 - 0.28)	0.02 ± 0.01 (nd - 0.04)	$\begin{array}{c} 0.04 \pm 0.05 \\ (nd-0.15) \end{array}$	
Export sword fish (176)	0.90 ± 0.51	0.09 ± 0.13	0.08	Jinadasa et al., 2014
Export yellow fin tuna (140)	0.30 ± 0.18	-	0.11 ± 0.16	
Export red snapper (28)	0.16 ± 0.11	0.01 ± 0.01	0.04 ± 0.04	
Export marlin (24)	0.49 ± 0.37	0.02 ± 0.02	0.05 ± 0.05	
European Commission specifications (mg/kg)	1.0 = sword fish, tuna; $0.5 =$ other	0.3 = sword fish; 0.1 = tuna; 0.05 = other	0.3 = tuna, sword fish	EC 1881 (2006)
Codex	Predatory fish = 1; fish = 0.5	Bivalve mollusks = 2	Fish = 0.3	Codex 193-1995

Lead in Sri Lankan food chain

The average concentration of lead in the Earth's crust is 10 mg/kg. The threshold value for lead in agricultural soils is 60 mg/kg. Sewage sludges used in agriculture elsewhere carry up to 300 mg/kg of lead. It is not possible

to prevent the entry of lead naturally present in soil into the food chain through plants.

Lead in agricultural soils in Sri Lanka is mainly of lithogenic origin. Acceptable lead concentrations of 15 -32 mg/kg have been reported from soils in Medirigiriya,
Talawa and Padawiya (Jayawardene *et al.*, 2012) and 4 mg/kg in rice soils recently (Balasooriya *et al.*, 2021). The average lead concentrations reported in soils of vegetable plots in the low country are within, yet close to the threshold value for agricultural soils, while the ranges of lead in upcountry vegetable soils reach 2 to 5-fold higher at the maxima of the ranges (Table 8).

The lead concentrations observed in urea and NPK mixtures of market fertilizers are around 4 mg/kg indicating a low contribution (Table 11). By contrast, the market TSP contributes 15 to 40-fold higher concentrations of lead compared to urea and NPK mixtures.

Herath *et al.* (2018) reported lead concentrations below 0.03 mg/kg in rice from CKDU endemic and nonendemic areas (Table 4). The lead concentrations observed in leafy vegetables from up country and low country are 20-fold higher compared to Codex and EU regulatory limits of 0.3 mg/kg (Table 10). These observations point more towards the probable anthropogenic origin of lead in some vegetable soils. The concentrations of lead observed in agricultural inputs, vegetables, and human test materials are correlated in Figure 3.

Ananda Jayalal *et al.*, (2019) observed lead concentrations of 0.20 mg/kg in rice and homegrown vegetables in CKDU affected families in the Anuradhapura district against the Codex tolerance limit of 0.3 mg/kg for leafy vegetables and 0.2 mg/kg for legume vegetables and pulses (Table 13). Table 14 shows lead concentrations below the tolerance limits for sea fish (Jinadasa *et al.*, 2010; 2014a) and in Table 13 for inland fish (Ananda Jayalal *et al.*, 2020).

Though the lead concentrations in rice and vegetable were notably high in home-grown vegetables, the cadmium and arsenic concentrations were low (Table 13). This may be an indicative sign of food safety hazards in the local foods due to lead hotspots.

The average lead concentrations in water from wells in 25 districts in Sri Lanka is $0.133 \pm 0.155 \ \mu g/L$ against the Codex regulatory limit of 50 $\mu g/L$ (Table 3). Though the lead concentrations in soils reported in the WHO study (2016) are within the threshold value, contribution from agricultural inputs appears high (Figure 3).

Lead concentrations in blood indicate human exposure. Madhavan *et al.* (1989) suggests that a concentration of 600 mg/kg of lead in soil contributes not more than 50 μ g/L to blood in children under 12-years of age. In an indicative communication of blood lead levels, 100 - 220 μ g/L has been reported in persons exposed to traffic polluted air in Kelaniya against 40 μ g/L in control groups (Gunasekera *et al.*, 2015). If the level in the control group is considered as normal exposure through foods, it stands below the WHO accepted limit of lead in blood of 100 μ g/L for adults and 50 μ g/L for children. Concentrations of lead in bones of CKDU affected persons were about 60% more compared to control groups (Ananda Jayalal *et al.*, 2020) suggesting a risk through lead in diet at least to some population groups.



Figure 3: Concentrations of Pb linked to agronomic practices, appearing in the food chain and in tissues of persons affected by CKDU.

The high concentrations of lead observed in soils of vegetable plots (Table 8), in up-country and low-country vegetables (Table 10), in household vegetables of CKDU patients (Table 13) and the WHO study (Figure 3) together indicates probable exposure of Sri Lankans to unacceptable lead concentrations leading to food safety hazards needing attention at least in certain locations.

Mercury in Sri Lankan food chain

The average concentration of mercury in the Earth's crust is 0.05 mg/kg. The threshold value for agricultural soils is 0.5 mg/kg. Contamination of agricultural soils by mercury is not possible as there are no ores containing mercury in Sri Lanka. Mercury enters the food chain in Sri Lanka mostly through sea fish. Studies on mercury content in the Pacific and Indian Oceans have shown a mean total mercury concentration of 5.3 ng/L with a range of 3-6 ng/L in sea water with no significant variations at different depths of the oceans. Mercury occurs mostly as methyl mercury in fish and shellfish. Of the total mercury in fish, organic mercury could be 80% (Kannan *et al.*, 1998). Methyl mercury is of high toxicity (Nishimura *et al.*, 1983).

Of the different types of fish, swordfish and tuna are known to commonly accumulate mercury. Among four types of fish tested for mercury in Sri Lanka, swordfish is observed to carry high mercury concentrations, sometimes beyond the regulatory limits established by the Codex and EFSA (Table 14). Mercury concentrations in tuna fish also reach regulatory limits at times.

Mercury is not considered a serious food safety problem associated with rice, though the concentrations in rice are generally described to be high among grains. Mercury concentrations reported in rice in Sri Lanka are very low (Table 5). There is no tolerance limit established for mercury in rice in the Codex or EFSA. In China, mercury of anthropogenic origin has been reported in rice from districts where mercury is extracted from the ores.

Mercury could become a food safety hazard only with high consumption of swordfish and tuna.

FOOD SAFETY AND HUMAN HEALTH LINKED TO HEAVY METALS

Chronic exposure of humans to heavy metals could occur through air, water, and food, or even through skin absorption. The major health problems associated with heavy metals of interest to Sri Lanka, identified from World Health Organization information, are as follows.

Arsenic (Arsenic key facts: WHO 2018)

Skin lesions leading to cancer, bladder cancer, and lung cancer. Increased risk of diabetes, pulmonary diseases, and cardiovascular diseases.

Cadmium: (Exposure to Cadmium, WHO 2019) Primarily toxicity to kidney, leading to tubulointerstitial damage, affecting lung function inducing cancer, and causing bone demineralization due to interference with calcium metabolism.

Lead: (Lead poisoning – key facts, WHO 2021) Cardiovascular effects, nervous disorders, decreased kidney function, and fertility problems

Mercury: (Mercury and Health, WHO 2017) Toxic effects on nervous, digestive, and immune systems.

Health problems arising from exposure to heavy metals

Effects of heavy metals on public health are established by understanding the exposure dose and resulting toxicity or carcinogenicity.

Arsenic related health problems are associated with inorganic arsenic. The patterns of cancer reported for 2019 by the National Cancer Control Program of Sri Lanka (Cancer Incidence data book, 2019), published in November 2021, provide the following information on the crude rate of new cancer cases per one hundred thousand of population in 2019 (Table 15).

Skin, bladder or lung cancer, which are the predominant health problems arising from exposure to arsenic do not appear to be high in the classified information in the Table 15. The value for skin cancer is approximately 2% of total cancers. The incidence of lung cancer is higher with males associated with smoking. The ratio of male to female CKDU patients reported in Sri Lanka is approximately 1:1 (Jayatilake *et al.*, 2013). Bladder cancer and lung cancers are the other major type of cancers caused by arsenic in food. Comparison of bladder cancer rates in Colombo with the three agricultural areas is given in the Table 16.

The reported cancer patterns are not congruent with the districts where agrochemicals containing arsenic are applied. Review of the variations of the different cancers in Sri Lanka over the period 2005 to 2019 do not show relative differences in the variations of cancer incidences by types. The patterns suggest that the factors affecting cancers in Sri Lanka are common to all types of cancers. Heavy metals in the food cannot be identified as a major contributor to cancer in Sri Lanka.

The incidence of CKDU in the North Central region of the country is well studied. Its symptoms, mainly the effects on the tubular cells of the kidney, are documented (Jayatilake *et al.*, 2013). It appears to be a toxicological problem different from cancer. There are no published evidence from the health sector in the country to suspect direct food safety hazards related to lead and mercury. There are mechanisms based on food safety studies to predict the possibilities of food safety hazards associated with exposure to heavy metals under discussion.

 Table 15:
 Crude cancer rate (CR*) in Sri Lanka by major types of cancers in 2019.

Males		Females	
Туре	CR	Туре	CR
Lip, tongue and mouth	20.6	Breast	39.5
Trachea, bronchi and lung	12.0	Thyroid	19.6
Colon and rectum	11.9	Colon and rectum	11.1
Oesophageal	10.4	Cervix, uteri	9.9
Prostate	9.5	Uterus	8.8
Pharynx	6.3	Oesophagus	7.3
Larynx	5.9	Lip, tongue, and mouth	8.8
Lymphoma	5.8	Ovary	8.5
Bladder	5.5	Trachea, bronchi and lung	4.1
Stomach	4.3	Lymphoma	3.9

CR* = number of new cancer cases diagnosed divided by at risk population multiplied by 100,000.

Table 16:Comparison of crude rates (CR) of bladder, tracheal, and
bronchial & lung cancers in 4 districts of Sri Lanka in
2019.

CR bladder cancer among males		CR Tracheal, bronchial and lung cancer among females	
District	CR	District	CR
Colombo	5.3	Colombo	5.8
Anuradhapura	0.4	Anuradhapura	0.6
Polonnaruwa	3.7	Polonnaruwa	3.6
Batticaloa	8.4	Batticaloa	1.7

Assessment of exposure of Sri Lankans to heavy metals through food chain

Assessment of safety levels of hazardous substances are done by examining the quantity of hazardous agent that would be ingested by subjects taking into consideration the quantities of contaminated food, the body weights and durations of exposure. Durations may be a month, a week, or a day. This information is examined against the adverse effects on public health to establish Provisional Tolerable Weekly Intakes (PTWI). The PTWI values are revised based on new evidence from the disease patterns associated with the hazardous component in the food. The units describing PTWI values are weight (mg / μ g / ng) of the hazardous substance per kg of body weight per week. Si Lankan data on heavy metals in foods are examined using this concept to recognize the degree of food safety. The values are compared with international norms.

Risks through exposure to arsenic

A study predicted the global burden due to arsenic by applying models designed by the US Environment Protection Agency to WHO data on cancer. According to the predictions, increased incidence of bladder cancer from 9129 to 119,176, lung cancer from 11, 844 to 121,442, and skin cancer from 10,729 to 110,015 annually attributable to inorganic arsenic in food is expected (Oberoi *et al.*, 2014). The prediction represents an approximately 11-fold increase in the three types arsenic induced cancers. In contrast to the above global prediction, the information from the cancer records in Sri Lanka (Cancer Incidents data book – 2019, 2021)

does not indicate an increase in lung, skin or bladder cancer patterns different from other cancers in Sri Lanka. Patterns of all cancer incidences have remained the same with only a 3-fold increase of total cancer, and each type of cancer incidences over 10 years.

A risk assessment study on exposure to inorganic arsenic in rice was done by the FDA (2016) in USA. The study assessed the risks quantitatively by examining the population affected by bladder cancer and lung cancer, as influenced by inorganic arsenic in rice consumed by pregnant mothers and infants. A parallel qualitative examination looked at exposure-effect relationships linked to other health concerns.

Important information from the above study of relevance to Sri Lanka is as follows.

- 1. Heavy metal toxicity associated with rice is mainly due to inorganic arsenic, which is a large fraction of total arsenic in rice.
- 2. The availability of inorganic arsenic in rice is 0.092 mg/kg for white rice and 0.154 mg/kg for brown rice in the USA, which were used in the study.
- 3. It is noted that 28-60% of inorganic arsenic is lost during the washing of rice prior to cooking.
- 4. A single rinsing of rice with water removes up to 15% of inorganic arsenic in rice.
- 5. A selected population group in USA consuming 2-3 meals of rice per day are exposed to 435 ng inorganic arsenic per kilogram of body weight per day.
- The regulatory limit of 0.2 mg/kg for arsenic in rice could reduce the cancer risk by 11%, and a limit of 0.075 mg/kg could reduce the risk by 79%.
- Reducing regulatory limit arsenic from 0.2 to 0.1 mg/kg would decrease the market availability of rice by 4 - 93% depending on other factors.

8. A healthy balance between availability of rice (food security) and risk (food safety) needs to be established scientifically for consumer benefit.

To examine the exposure levels and predict the risks associated with arsenic in the Sri Lankan food chain, PTWI values were calculated (Table 17). The following assumptions were made in the calculations.

- a) The average body weight of a Sri Lankan is 55 kg.
- b) A Sri Lankan consumes 438 g of rice per day.
- c) Inorganic arsenic (I-As) in rice is 80% of total arsenic (T-As).

The Table 17 compares the exposure of Sri Lankans against recommendations of JECFA. An attempt to compare it with the USA situation for those consuming 2-3 meals per day, as reported by the FDA, was also made. The basis used here is the calculation of PTWI in μ g of inorganic arsenic per kg of body weight over one week. Calculations and predictions here involve varying amounts of uncertainty due to approximations and assumptions.

Based on above calculations, it appears that the concentrations of arsenic naturally present in rice in Sri Lanka would not pose a significant risk to human health through weekly exposure. If the PTWI value is reduced to 10 or 12 in the future by JECFA based on new exposure-effect evidence, the calculation would show a tendency towards an increased risk through rice at the maximum reported values from Girandurukotte, but not with values in the other two studies. Scientifically it is not possible to reduce the amount of natural arsenic in soil. The concentrations of arsenic naturally present in Sri Lankan soils are about 1/5 threshold value for agricultural soils. It may be possible to minimize

 Table 17:
 Comparison of exposure of Sri Lankans to As through rice on different published data converted to PTWI.

I – As in rice (mg/kg)	- As in rice (mg/kg) Notes	
	JECFA exposure value withdrawn in 2015	15*
0.20 Codex limit	Applied to 438g rice / day in SL	11.1
	Based on 435 ng/kg bw/day (FDA, 2016)	3.0
0.07 = 80% T-As	Maximum reported (Jayasekara & Fretas, 2005)	3.9
0.21 = 80% T-As	Max. Girandurukotte (Chandrajith et al., 2011)	11.7
0.03 = 80% T-As	Mean reported (Herath et al., 2018)	1.7

* Protection provided by Provisional Tolerable Weekly Intake (PTWI) of 15 μ g I-As/kg bw was found to be inadequate and withdrawn in 2015

risk by preventing entry of inorganic arsenic to rice soils through rigorous testing of agricultural inputs or reduction of rice consumptions perhaps from 3 meals to 2 meals a day. The fertilizer import policy needs to address this food safety issue. The arsenic, cadmium, and lead concentrations in fertilizer are concerns in most countries. The State of Texas has established a limit for arsenic as 41 mg/kg in fertilizer (Westfall *et al.*, 2005). Other countries address the heavy metal concentrations in phosphate fertilizers or by examining heavy metals added to soil over 45-year period (Agriculture and agrifood Canada, 1997).

Risk through exposure to cadmium

Food safety concerns about cadmium in the Sri Lanka arise mainly due to its nephrotoxic effects. The high quantities of rice consumed in Sri Lanka and Bangladesh are known to expose the populations to unhealthy concentrations of cadmium compared to other rice consuming countries. PTWI values are used to recognize the exposure to cadmium from different food sources based on the information from published research data. The PTWI values for Sri Lanka arising from rice, vegetables and fish are presented in Table 18.

The following assumptions were made in calculating PTWI for exposure to cadmium.

- a) The average body weight of humans is 55 kg.
- b) The average daily consumption of rice is 438 g.
- c) The average daily consumption of vegetables is 1.73 portions (approximately 130 g)
- d) The average daily leafy vegetable consumption is 50 g.
- e) Weekly fish consumption is 300 g.

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Thresholds derived for "safe" exposure to cadmium by international regulatory agencies are given as the basis to understand the risks.

The PTWI values indicates high exposure of Sri Lankans to cadmium from rice and green leafy vegetables. Uraguchi and Fujiwara (2012) reported the intake of cadmium in Japanese population is $3.0 \ \mu g/kg$ bodyweight per week, which is above the PTWI limit of EFSA. In this respect the PTWI values from Sri Lankan foods in the Table 18 indicates a very high degree of food safety risk.

The high concentrations of cadmium observed in urine, hair and postmortem kidneys of CKDU affected persons further indicate the exposure of Sri Lankans to cadmium through the food chain (Figure 2). Accumulation of cadmium in bones, which is a feature of cadmium toxicity is however not detected in the WHO study (2016) or by Ananda Jayalal *et al.* (2020). Bone demineralization arising from exposure to cadmium is described more as an indirect influence via induction of renal dysfunction (Godt *et al.*, 2006). The observations on relationship of cadmium with bone demineralization differ, though there is evidence supporting bone demineralization with Itai-Itai disease linked to cadmium in rice.

Patterns of daily intake of cadmium as a function of rice consumption is presented in Figure 4. The exposure due to rice consumption at 438 g per day in Sri Lanka is represented by the brown upward arrow. The red dotted line indicates the exposure of Sri Lankans to cadmium through rice at different concentrations of cadmium in rice and quantities of rice consumed.

 Table 18:
 Comparison of exposure of Sri Lankans to Cd through foods based on published research data converted to PTWI.

PTWI	Notes
0.7	US Agency for toxic substances and diseases registry
2.5	European Food Safety Authority (EFSA, 2012)
5.8	Equivalent for JEFCA value of 25 μ g/kg PTMI (monthly intake)
9.5	Rice based on Herath et al., 2018 (Table 4) for 0.17 (average) mg/ kg
5.3	Vegetables based on Cd observed in endemic area (Figure 2)
1.0	Vegetables based on Cd observed in non-endemic area (Figure 2)
3.2	Leafy vegetables based on (Table 9); Approximated as 0.5 Cd mg/kg
2.5 - 7.0	Rice, vegetables and water combined based on Jayalal <i>et al.</i> (2015) 60 kg body weight (consumption: 152-419 μ g Cd / week)
0.23	Fish at reported maximum value of 0.049 mg/kg (Jinadasa <i>et al.</i> , 2010); for maximum exposure with consumption of 300 g per week

The curves indicate the need to either reduce the cadmium in rice or reduce consumption of rice to levels around 2/3 of the current consumption to be in line with Codex standards or to a level of 1/3 to be in line with EFSA standards.

Evidence leading to food safety hazards through cadmium in the food chain is apparent through several studies. Sanjeevani *et al.* (2017) indicated only cadmium, among the heavy metals under discussion here, to be of anthropogenic origin. High cadmium concentrations in TSP used as a fertilizer in Sri Lanka is documented in Table 11 (Chandrajith *et al.*, 2010) and the heavy cadmium concentrations associated with vegetable production in Tables 8 and 10. The observations of Meharg *et al.*, (2013) in Table 9 provide independent evidence on high exposure of Sri Lankans to cadmium. Cadmium is a known cause of tubulointerstitial damage to the kidney. Tubulointerstitial damage is the main lesion among CKDU patients.

When the contribution of cadmium from leafy vegetables (Figure 3) is added to the contribution from rice, the need to prevent the use of TSP as a fertilizer and application of better quality phosphate fertilizer becomes evident. It must be mentioned that the cadmium concentrations in Eppawala rock phosphate is much less compared to TSP detected in Sri Lankan markets (Table 11 and 12).



Figure 4: Daily intake of cadmium as a function of rice consumption modified from Zhao and Wang (2020).

Regulating the entry of heavy metals present in fertilizer into the human food chain is difficult. However, there are regulatory limits for heavy metals in fertilizers in other countries. California has regulations for arsenic, cadmium, and lead in fertilizers. The regulatory limits are linked with the main components in fertilizers such as 1% P_2O_5 or 1% Zn arising from minerals. The regulations in Canada specifie application limiting to accumulation over a 45-year period expressed as kg per hectare (Agriculture and Agri-foods Canada, 1997). The regulations in Texas specify the limits of arsenic, cadmium, lead, and mercury as 41, 39, 300 and 17 mg/kg of fertilizer respectively (Westfall et al., 2005). A mechanism to limit the entry of cadmium into agricultural soils is a need, in order to prevent food safety hazards arising through commercial agriculture. Limiting the entry of cadmium to agricultural soils is highly relevant to Sri Lanka, in the light of evidence regarding high cadmium concentrations in rice and leafy vegetables arising from use of TSP.

Risk through exposure to lead

Risks in the food chain may arise from lead present naturally in soil or added through contaminated agricultural inputs. The lead content reported in rice is less than 0.03 mg/kg on the average (Table 4), which is much below the Codex tolerance limit of 0.2 mg/kg. Norton *et al.* (2014) have reported lead concentrations of 0.020 mg/kg and 0.048 mg/kg in white and brown rice from Sri Lankan markets respectively. Based on findings of Norton *et al.* (2014) and Herath *et al.* (2018), 0.03 mg/kg was taken as a basis for estimating the contribution to the PTWI of lead from rice.

With the lead content in water reported as $0.133 \ \mu g/L$ in well water (Herath *et al.*, 2018) against the Codex tolerance limit of 10 $\mu g/L$ (WHO 2016), the risk arising from drinking water would be insignificant. Maximum lead concentrations of 0.486 mg/kg reported in vegetables (WHO, 2016) and in leafy vegetables (Premarathna *et al.*, 2011) are of concern, and used in calculating PTWI.

The following assumptions were made in calculating the PTWI and PTTI values for exposure to lead through food. Each contribution may be used to recognize cumulative contribution.

- a) The average body weight of humans is 55 kg.
- b) Rice consumption is 438 g per day containing 0.03 mg/kg of Pb (Herath *et al.*, 2018)
- c) A mean contribution of 8.5 mg/kg from leafy vegetables (Table 10, Premarathna *et al.*, 2011)
- A weekly consumption of 300 g of fish containing a mean of 0.11 mg/kg lead.
- e) The total diet of farmers from lead hot spots consists of 330 g of food with 0.200 mg/kg lead.

The analysis indicates high exposure to lead through leafy vegetables.

In estimating the provisional tolerable intake of lead through food, a factor of 10 was applied to obtain an exposure level to achieve some margin of safety based on lead concentrations in blood (Carrington & Bolger, 1992). Estimation of lead in blood is the only reasonably acceptable level of understanding exposure to lead from a variety of sources, *i.e.*, food, water and air. The FDA classifies a population into 4 groups by age. When the reported lead concentrations in Sri Lankan foods is viewed from the daily exposure angle, the resulting Provisional Tolerable Total Intake (PTTI) values could be compared (Table 20). The values reflect daily exposure in micrograms.

The risk due to lead through exposure to vegetables appear to reach the PTTI value of the FDA for adults.

Table 19:	Comparison of exposure of Sr	Lankans to Pb through the	foods based on published	research data converted to PTWI
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PTWI	Notes
25	This Codex value was declared inadequate in 2010
1.67	At the rate of 0.03 mg/kg in rice (Herath et al., 2018; Norton et al., 2014); Consumption of 438 g rice per day
8.04	Based on WHO, 2016 (0.486 mg/kg); consumption of 130 g vegetables per day
52.2	Based on Table 9 (approx. maximum of 8.2 mg/kg); consumption of 50 g leafy vegetables per day.
7.7	Household rice and vegetables 330 g/day from CKDU patients with approx. 0.200 mg/kg of lead (Ananda Jayalal <i>et al.</i> , 2019); 60 kg body weight.
4.2	Fish (observed mean) 0.11 mg/kg in tuna fish (Jinadasa et al., 2014)

Table 20: Comparison of exposure of Sri Lankans to Pb through foods based on published research data converted to PTTI

PTTI (µg/day)	Notes	
6	Limit for children below 7 years	
15	Limit for children above 7 years	
25	Limit for pregnant women	
75	Limit for adults	
13.4 (Rice)	At the rate of 0.03 mg/kg in rice (Herath et al., 2018; Norton et al., 2014); Consumption 438 g rice per day	
63.2 (Vegetables)	Based on WHO (2016) (0.486 mg/kg); consumption 130 g per day	
410 (Leafy vegetables)	Based on Table 9 (approximate maximum lead of 8.2 mg/kg) and leafy vegetable consumption 50 g per day	
66	Household rice and vegetables 330 g/day from CKDU patients approximated to 0.20 mg/kg of lead (Ananda Jayalal <i>et al.</i> , 2019)	
4.4	Fish (mean) 0.11 mg/kg in tuna (Jinadasa <i>et al.</i> , 2014); consumption 43 g per day [mean exposure may be 1/10 of this]	

Risk through exposure to mercury

CKDU affected subjects.

Risk associated with mercury in the food chain occurs mainly on exposure to methyl mercury in fish, which is of higher toxicity than its inorganic forms. Methyl mercury accounts for 83% of total mercury in fish muscle from south Florida estuaries (Kannan *et al.*, 1998). In the same location, methyl mercury in estuary sediments were only 0.77%, and that of water is less than 52% of total mercury. Separate studies by the same research group in Sri Lanka have shown mean total mercury concentrations

concentrations of lead in bones of autopsy samples of

U Samarajeewa

of 0.43 in cooked, 0.90 in export, and 1.24 mg/kg in fresh swordfish, with a maximum value of 1.47 mg/kg (Table 14). These values were used in estimating the PTWI values for comparison with international norms.

The following assumptions were made in calculating the PTWI values for exposure to total mercury through fish.

- a) The average body weight of humans is 55 kg.
- b) The concentrations of mercury in fish is 1.24 mg/kg (mean value for sword fish) and 2.58 mg/kg (maximum reported for sword fish) (Jinadasa *et al.*, 2010).
- c) The average consumption of fish is 300 g per week, as observed in a study in Gampaha district (Jayasinghe *et al.*, 2018).
- d) The average consumption of pelagic fish, mainly swordfish, is 83 g per week (Jinadasa *et al.*, 2014b).

On the above basis, PTWI levels for mercury are shown in Table 21.

 Table 21:
 Comparison of exposure of Sri Lankans to mercury through sea fish based on published research data converted to PTWI.

PTWI	Notes
1.6	JECFA PTWI for methyl mercury (2003)
5.0	JECFA PTWI for total mercury (1978)
2.3	Assuming consumption of 300 g per week of cooked fish containing 0.43 mg/kg total mercur (Jayasinghe <i>et al.</i> , 2018)
6.7	For a mean value of 1.24 mg/kg of total mercury in swordfish (Jinadasa et al., 2010)
3.9	Maximum of 2.58 mg/kg of total mercury in swordfish (Jinadasa <i>et al.</i> , 2014b) with a weekl consumption of 83 g.

Table 22: Summary of food standards for heavy metals in developed countries (mg/kg)

Regulator	Arsenic	Cadmium	Lead	Mercury
Codex - food	0.1 [rice 0.2]	0.05-0.2	0.001 - 0.3	0.5 -1.0
Codex - water	10 µg/L	3 µg/L		0.1 µg/L
European Commission (ESFA)	0.1-0.3 [rice products]	0.05 -0.5 [meats] 0.05 [root vegetables] 0.2 [leafy vegetables]	0.02 - 0.5 0.1 [vegetables]	0.5 - 1.0 [fish]
USA - FDA	10 achievable in apple juice	50 ppb suitable for juices		
USA - EPA - water	10 µg/L	7 μ g/L for tap water	0 µg/L for tap water	1 μ g/L for tap water
FAO/WHO	10 μg/L for water	0.2 for food	0.3 for food 100 $\mu g/L$ water for adults 10 $\mu g/L$ water Canada	1 μg/kg body weight per week other than seafoods

If the data in the Table 21 are examined assuming that methyl mercury is 75% of total mercury (Florida estuaries showed 80%), the calculated PTWI values would be almost the same as JEFCA value for cooked fish, and would exceed that for fresh swordfish. Jinadasa *et al.* (2014a), in a study of 140 and 176 samples of yellow fin tuna and swordfish, concluded that they contribute only 9% and 27% total mercury respectively to the PTWI of JCEFA, based on consumption of 83 g large pelagic fish per week. They also reported that 32% of swordfish samples exceeded the EU tolerance limit of 1 mg/kg for mercury, but not any of the tuna fish samples.

The results suggest that there is a food safety risk to persons consuming tuna or swordfish as the only type of fish in Sri Lanka in high volumes.

Regulations on heavy metals in foods

Each country decides on the level of protection needed for its population from food safety hazards. This results in variations in food standards. A summary of the food standards by major food safety authorities in the World is given in Table 22.

Fable 23 : Proposed horizonta	food standards for	heavy metals	in Sri Lanka.
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Heavy metal	Commodities	MTL mg/kg	Notes
Arsenic (Total)	Rice and rice- based products for adults	0.2	Recommendations take into consideration what is agronomically possible based on reported levels of arsenic in soil and water, leaving out what may be contributed by agricultural inputs, and accepting what is recognized internationally as safe.
Arsenic (Total)	Rice and rice -based products for infants	0.1	Risk reductions need to be addressed for more vulnerable groups, the infants, children, and pregnant mothers consuming special foods through rice-based cereals <i>etc</i> .
Arsenic (Total)	Drinking water	10 µg/L	Codex limit would be satisfactory under Sri Lanka conditions as total arsenic levels are 7 to10-fold less than the limit.
Arsenic (Total)	Vegetables and fruits	0.2	General Codex limit for arsenic in foods as there is no evidence of arsenic in vegetables and fruits in Sri Lanka
Arsenic (Inorganic)	Fish and fish products	0.2	The bulk of arsenic in fish is in a non-toxic form as arsenobetaine.
Cadmium	Rice and rice products	0.1	This appears achievable under current cultivation conditions and would prevent development of cadmium hot spots. The need is linked with observed high consumption of rice in Sri Lanka
Cadmium	Water	3 µg/L	Codex standard
Cadmium	Vegetables and fruits	0.2	Stringent conditions are needed considering current field observations. Controls through fertilizer and pesticides is possible. The suggested value is the Codex limit.
Cadmium	Fish & fish products	0.1	In line with the limits in many countries. It is achievable.
Lead	Rice & rice products	0.1	A more reasonable tolerance limit to discourage use of contaminated agricultural inputs while accommodating natural soil concentrations.
Lead	Water	10 µg/L	This low limit is suggested as it is already reflected in reports and would be beneficial to be used as a mechanism to reduce the overall exposure and the risk through lead
Lead	Vegetables and fruits	0.3	A low tolerance limit for vegetables to discourage use of agricultural inputs contaminated with lead. Same as Codex limit
Lead	Fish & fish products	0.3	Achievable. It is in line with EFSA limits.
Mercury	Rice & rice products	0.02	Mercury is not a food safety hazard associated with rice. It is best to use the Codex standard
Mercury	Water	$1 \ \mu g/L$	Same as Codex standard
Mercury	Vegetables and fruits	0.02	Mercury is not a food safety problem associated with vegetables. It is best to use the Codex standard.
Mercury	Fish & fish products	1.0 methyl mercury	For predatory fish (shark, swordfish, tuna). 0.5 for other fish. Same as Codex standards

All developed countries and most of the Southeast Asian countries moved to new food safety authorities and horizontal food safety standards during 2012 to 2020. Preparation of country specific standards needs deep understanding of the exposure levels and risks arising from heavy metals in foods and dietary patterns.

Recommendations

Considering this national need, the recommendations given in Table 23 are made to develop Horizontal Food Safety Standards in Sri Lanka.

CONCLUSION

Sri Lanka needs to understand the critical points in the food chain where controls should be introduced to ensure the food safety of the nation without affecting its food security. It requires addressing the problems with deep scientific understanding. This review is meant to provide the analyzed scientific information for Sri Lanka in its efforts to ensure a safe food supply.

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RESEARCH ARTICLE

Machine Learning

Fuzzy based congestion detection and control algorithm for energy efficient wireless sensor network (WSN)

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Abstract: A wireless sensor network (WSN) is built by numerous tiny, resource aware sensor nodes to facilitate and develop various fields such as the military, health care, disaster management etc. One of the major design issues in WSN is congestion detection and control. Data collision, channel contention, and buffer overflow are considered to be a few important factors that may lead to a congestion problem. Congestion causes increase in packet drop rate and thus network efficiency is decreased. Hence, the network performance and quality of service may be degraded. Therefore congestion detection and control are vital and critical challenges to be addressed. In this paper, a novel energy efficient fuzzy based congestion detection and control algorithm (EE-FBCDCA) has been proposed with the objective of addressing the energy aware congestion detection and control mechanism. The proposed work consists of three phases, namely, 1) Fuzzy inference system based congestion detection, 2) Congestion notification phase and 3) Fuzzy inference system based congestion control. In this work, once the congestion is detected and upon receiving the notification message, the intermediate nodes will adjust their respective transmission rate to prevent further drop of packets. Thus, EE- FBCDCA detects and control the congestion more effectively to improve the QoS of the network. The proposed work is implemented and analysed by using MATLAB, and the performance is compared with the existing HTCCFL and FBACC. The simulation results of EE-FBCDCA shows enhanced network lifetime, energy efficiency, and reliability.

Keywords: Congestion detection, control, fuzzy inference system, energy, network lifetime, WSN.

INTRODUCTION

A wireless sensor network (WSN) consists of numerous tiny sensor nodes deployed over a vast geographical area with limited resources such as energy and bandwidth availability (Bhushan & Sahoo, 2010; Banimelhem & Khasawneh, 2012; Antoniou et al., 2013; Chen et al., 2016; Rao & Rama Rao, 2022). In today's era, WSNs have wide applications in various fields, namely healthcare systems, military, multimedia etc. The sensor nodes are capable of collecting, processing and transmitting the data through the intermediate nodes to reach the sink nodes (Farzaneh & Yaghmaee, 2015). In WSN many design issues such as queuing delay, packet drop, packet collision, and congestion need to be addressed. Among all the above stated issues, one of the major problems in WSN is congestion, which needs to be addressed to enhance the network lifetime and performance. Congestion leads to numerous issues especially inefficient energy consumption by nodes, buffer overflow, and high packet loss (Gholipour et al., 2015). Thus, this problem will degrade the network performance and quality of service. The primary working principle of nodes is to sense, gather, process, and transmit the data to the sink nodes for analysis and decision making. In WSN, each node transmits the gathered data to the sink node for processing. Thus, vast amounts of data are forwarded towards the base station at

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any given instant of time, leading to network congestion and packet loss. As WSN play a vital role in various real time application domains, congestion will jeopardize the network lifetime and performance that need to be avoided (Jan et al., 2014). Thus, congestion is considered as a very important design challenge. Hence, it becomes essential to implement an efficient and reliable congestion control mechanism to address the problem (Sonmez et al., 2014). Moreover, sensor nodes are power constrained devices and therefore energy is considered to be one of the major performance metrics during congestion detection (Uthra & Raja, 2014). Much research has been done towards the enhancement of energy conservation to prevent packet collision and congestion (Wang et al., 2012). But one of the major drawbacks in all this work is that energy consumption is not equally distributed among all the nodes (Ahmed et al., 2014; Yi & Yang, 2016; Zou & Qian, 2019). Different mechanisms and schemes were utilized by various researchers towards congestion detection and control process. Moreover, many research projects are focused towards fuzzy based congestion control and detection mechanisms (Munir & Man, 2007; Jutan et al., 2012; Rekha & Gomathy, 2012). Therefore, designing a novel energy efficient congestion detection and control process with a better solution is considered as a major research challenge. In this work, a novel fuzzy based energy aware congestion detection and control algorithm for WSN has been proposed. This algorithm yields a better solution for congestion detection and control when compared to other existing algorithms.

The primary focus of the proposed work is to develop an energy aware, high reliable congestion detection and control algorithm for WSN. The proposed algorithm consists of three phases. The first phase is defined as a novel fuzzy based energy aware congestion detection phase with fuzzy input variables, such as packet delivery rate, residual energy level, and queue occupancy, with the aim of performing the task of congestion detection to enhance the energy efficiency of the network.

The next phase, the congestion notification phase, will be triggered to transmit the congestion signal to the nodes whenever congestion occurs in the network. Finally, in the third phase, *i.e.*, the fuzzy based congestion control phase, the data transmission rate of the nodes is controlled to prevent further congestion in the network. Thus, the proposed work ensures energy efficient, high throughput, minimal end-to-end delay, a smaller packet drop rate, and overall enhancement in the network performance. Much research has been carried out in the field of congestion detection and controls in WSN (Misra *et al.*, 2009; Jenolin *et al.*, 2011; Jan *et al.*, 2014; Mahdizadeh *et al.*, 2014; Ghanavati *et al.*, 2015). Ghaffari (2015) and Rekha Chakravarthi *et al.* (2010) have carried out comparative surveys on the congestion control algorithm. Congestion control (Chen & Zhang, 2006) is defined as a technique of controlling the amount of data flow in the network without any loss of information (Rezaee *et al.*, 2014). It consists of three primary phases namely detection (Chen & Yang, 2006), notification (Wan *et al.*, 2011), and control (Yin, 2009).

Ahmed & Paulus (2017) developed congestion avoidance and mitigation for multipath routing and load balancing in WSN. Cao *et al.* (2012) proposed a novel delay based congestion control for a multipath TCP feedback congestion control protocol. The experimental results of this protocol shows better energy and throughput.

Huang *et al.* (2009) developed an FACC protocol to detect congestion based on the classification of the nodes adjacent to the source and adjacent to the base station. Tao and Yu (2010) presented an enhanced congestion detection and avoidance protocol for detecting congestion by using buffer threshold and buffer weight. This protocol supports a hop-by-hop congestion control scheme. Adjeroh & Yaghmaee (2008) presented a queue based congestion control protocol with priority support to congestion control by considering the queue length as the index of density degree. Thus, in this protocol the transmission rate is controlled depending upon the priority and density degree.

Basaran *et al.* (2010) developed a control of sensor queue (CONSEQ) protocol with the aim of detecting congestion by considering the queue length and data transmission among the one hop neighbouring nodes. Akyildiz *et al.* (2003) proposed an event-to-sink reliable transport protocol. In this protocol, the packet transmission rate to the sink node is observed and queue overflow is identified. The intermediate nodes will detect the presence of congestion in the network by referring to the congestion bit in the packet.

A fuzzy inference system can be utilized to implement the congestion detection and control scheme (Rezaee & Pasandideh 2017; Sangeetha *et al.*, 2019). Many fuzzy inference system based models have been proposed to address various issues in WSN (Logambigai & Kannan, 2016; Rajeswari *et al.*, 2019). A fuzzy temporal congestion predication model was proposed to identify the congestion free path. This model consists of two phases, namely, congestion prediction and localization phases (Mukherjee & Dasgupta, 2013). Hatamian *et al.* (2016) developed a fuzzy rate controller based congestion aware routing by assigning priority to data packets. Ghanavati *et al.* (2013) presented a type 2 fuzzy logic algorithm to measure and control the congestion level, to enhance the QoS in WSN. This algorithm improves the network lifetime and decreases the packet drop rate.

Ghaffari (2014) proposed a star algorithm based routing mechanism to ensure improved link quality and reduced hop count. But in this method end-to-end delay is more and thus the complexity is greater. Sayyad and Choudhari (2014) proposed a Hierarchical Tree based congestion control using fuzzy logic (HTCCFL) using the following metrics, namely, packet service ratio, number of competitors, and buffer occupancy. The above algorithm gives increased energy efficiency by selecting an alternate path with less congestion for data transmission.

Hatamian and Barati (2015) developed a priority based congestion control schema using fuzzy rules in WSN. In this method, buffer occupancy and congestion level are considered as input parameter for a fuzzy rate controller to monitor the packet transmission rate. Damaso *et al.* (2014) proposed a reliable framework with energy as a primary constraint. Blocks and models were used to express the sensor reliability elements using the energy factor. Mostafaei (2019) proposed distributed learning automation (DLA) to identify the small group of nodes to enhance end to end reliability.

Ullah *et al.* (2017) proposed an energy and congestion aware routing for an AMI network in a smart city. In this method, residual energy and queue occupancy are considered as major factors monitoring and controlling the congestion level. Song *et al.* (2019) developed a unified modelling framework known as GLS for space terrestrial networks. Jaiswal and Yadav (2013) developed a fuzzy based adaptive congestion control (FBACC) that enable congestion estimation by considering the factors buffer occupancy and traffic rate. FBACC enhances the energy efficiency and reduces the packet drop rate.

Gharajeh and Khanmohammadi (2016) developed DFRTP based upon the distance and numbers of neighbouring nodes, which were considered as the input parameters. DFRTP yields better network lifetime and packet drop rate. Chen *et al.* (2015) developed a RLAN to detect the node position under varying magnitude across various directions. Khan *et al.* (2020) proposed and implemented the ant colony optimization Technique and repetitive route configuration with reactive routing protocol for obstruction of Black Hole Attack in mobile *ad hoc* networks. Perumal and Subramaniyan (2020) developed the statistical Markov model based natural inspired glow worm multi-objective optimization (SMM-NIGMO) technique for Energy efficient data delivery in MANET.

In spite of the availability of much related work in the literature survey, an energy efficient congestion detection and control in WSN still remains as a major design goal, following the metrics buffer size, service time, and channel load. These studies do not focus much on energy efficiency based congestion detection and control process. Hence there is a need to design an energy efficient congestion detection and control algorithm for WSN for enhancing the throughput, reducing delay, and increasing energy efficiency and reliability. This novel challenge is addressed in this work. Moreover, most of the related work focuses on Energy Efficient Fuzzy Based Congestion Detection and Control Algorithm, and This work is proposed with the objective of improving the network performance and lifetime.

The rest of this paper is structured as follows. Section 2 describes in detail the Proposed Methodology. Section 3 illustrates and discusses the simulation based results. Section 4 finally concludes the paper.

MATERIALS AND METHODS

Energy efficient fuzzy based congestion detection and control algorithm (EE- FBCDCA)– energy model

The energy model considered in this proposed work, EE-FBCDCA, is similar to the work developed by Heinzelman and Balakrishnan (2002). According to the first order radio model, the amount of energy consumed during the transmission of a packet with 1 bit over d distance is given by the Equation (1). The E_{elec} indicates the amount of energy consumed per bit to run the transmitter or receiver circuitry. E_{fs} and E_{mp} are the amplifier energies in free space and multipath, respectively.

$$E_{Tran}(L) = \begin{cases} LE_{elec} + L\varepsilon_{fs}d^2 ford < d_0 \\ LE_{elec} + L\varepsilon_{mp}d^4 ford > d_0 \end{cases} \qquad \dots (1)$$

where
$$d_0 = \sqrt{\frac{\varepsilon_{fs}}{\varepsilon_{mp}}}$$

The energy $E_{RX}(1)$, which indicates the amount of energy consumed in receiving a packet with 1 bit, is given by Equation (2).

$$E_{RX}(\mathbf{l}) = E_{elec} \qquad \dots (2)$$

EE-FBCDCA

An energy aware congestion control mechanism is considered to be a major design challenge for WSN.

Hence in this work, a novel EE- FBCDCA based on a fuzzy inference system has been proposed with the objective of detecting and controlling the congestion. The residual energy level and reliability are considered as major aspects for measuring the network performance. In the proposed algorithm the network congestion is detected by considering the data flow from sources to base station. Moreover, buffer capacity is considered to be an important metric for the process of congestion monitoring and detection. The proposed work consists of the following three phases, namely i) Congestion Detection Phase, ii) Congestion Notification Phase, and iii) Congestion Control Phase. Figure 1 Shows the architecture of the proposed EE-FBCDCA.



Figure 1: Architecture of the proposed EE- FBCDCA

Design of energy efficient fuzzy inference system for congestion control and detection Phases.

In this proposed work, the FIS is defined under the following two scenarios: firstly, for congestion detection phase and secondly, during the process of congestion control.

The fuzzy inference system (FIS) employed in this work consists of the following 4 main components:

- (i) Fuzzifier
- (ii) Fuzzy inference engine (FIE)
- (iii) Fuzzy rule base, and
- (iv) Defuzzifier.

Congestion detection and control are the most important design challenges in the WSN environment. Thus, to address these challenges an EE- FBCDCA is developed in this work. The major goal of the proposed work is address detection as well as controlling congestion in the network, in order to to improve the network reliability and lifetime. Thus, this section elaborates in detail the proposed work, EE- FBCDCA. The main flow of the proposed EE- FBCDCA is shown below in Algorithm 1.

The EE- FBCDCA has been proposed with the following two major objectives.

1. A novel energy aware fuzzy based congestion control scheme is proposed to prevent the congestion.

2. Energy conservation is improved by congestion control using the fuzzy rules. Hence, the network lifetime is enhanced.

The proposed EE- FBCDCA consists of the following three phases:

- 1. Fuzzy inference system based congestion detection phase
- 2. Congestion notification phase
- 3. Fuzzy inference system based congestion control phase

Algorithm 1 - EE- FBCDCA

Input : Senor Nodes **Output : Congestion Free Network Environment** For each Node n do Random number = rand(0,1)If $(q_{len} < TH)$ then Congestion Level Fuzzy logic1 (REL, PDR, Buffer Capacity) If (*Congestion_Level < Random_Number*) Add Packet 'P' to the node's buffer Else Send Congestion Warning Signal (Node ID,REL) to its Neighbour Nodes End If For each Neighbour (M) on receiving the Congestion Warning Signal Transmission Rate Control Fuzzy logic 2 (Congestion Level, REL) End For End if End For

The following subsection describes each of the phases in detail.

Fuzzy inference system based congestion detection phase

In the proposed work, the fuzzy inference system is employed to detect the congestion level of the network. The main objective of this proposed work is to improve the network lifetime and performance by controlling and managing the transmission rate of the nodes, so that the congestion level will be reduced. The primary role of this phase is to detect the congestion level of every node in the network. Thus, in this phase, queue length is measured periodically to detect the traffic level in the network. Moreover, in this proposed work the fuzzy logic inference system is used for measuring the congestion level. The three fuzzy input variables used are Buffer Capacity, Residual Energy, and Packet Delivery Rate.

Description of the fuzzy variables

Buffer capacity

The buffer capacity of any node, say (n), is defined as the maximum amount of data packets that a node can store before processing. Thus, the buffer capacity parameter plays a vital role during the congestion detection scheme. The buffer capacity is measured by using Equation (3), given below:

$$Buffer_Capacity = \frac{Buffer_{Size} - Buffer_{Packet}}{Buffer_{Size}} \qquad ...(3)$$

where $Buffer_{Size}$ indicates the buffer size of a node, and $Buffer_{Packet}$ indicates the number of packets available in the buffer at given instant 't'.

Hence, if the $Buffer_{Packet}$ is zero, then it indicates that the node's buffer is empty, whereas if $Buffer_{Size}$ is equal *to* $Buffer_{Packet}$ it indicates that the buffer is full and thus all the arriving packets are dropped, which may lead to congestion in the network.

The threshold Buffer length (B_{TH}) is calculated by using Equation (4), given below:

$$B_{TH} = \frac{\text{Packet}_{\text{T}}^{\text{N}}}{Buffer_{Size}} \qquad \dots (4)$$

where $Packet_T^N$ is the amount of data packets arrived at node 'N' at given time 't'.

Packet delivery rate

The packet delivery rate at any node 'n' is estimated with respect to the packet service time at node 'n'. P_{ST} is defined as the time taken by the node for successfully transmission of a single packet over the communication path.

Thus, the packet average service time can be measured by considering an exponential weighted sum formula as shown below, Equation (5):

$$P_{ST} = (1 - \alpha)P_{ST} + \alpha P_{ST} \qquad \dots (5)$$

where α is constant such that $0 \le \alpha \le 1$

Thus, by using the measured P_{ST} , the node rate can be estimated by using Equation (6), given below:

$$PDR = \frac{1}{P_{ST}} \qquad \dots (6)$$

Residual energy level (REL)

The REL of a node is calculated by using Equation (7), given below:

$$\operatorname{REL}(N) < E_{\operatorname{thres}} \qquad \dots (7)$$

where REL (N) is the Residual Energy Level of the node N, and E_{thres} is the threshold energy. The threshold energy (E_{thres}) is calculated by using Equation (8).

$$E_{thres} = \frac{\sum_{i=1}^{N} REL_i}{N} \qquad \dots (8)$$

where REL is the Residual Energy Level, and N is the Number of nodes in the network.

Congestion in a network leads to packet drop or collision that may increase the retransmission of same set of data packets by node. Hence, the node's energy level drastically decreases. Thus, congestion acts as one of the primary source of decrease in network lifetime as well as energy efficiency. Table 1 shows the fuzzy rules for estimating the congestion level in the system.

The following parameters such as Buffer Length, Residual Energy Level, and Packet Delivery Rate are considered as fuzzy inputs whereas the Congestion Level act as fuzzy output. Thus, the fuzzy inputs variables and its linguistic variables used for Congestion Level estimation are shown below.

Buffer Length has the following linguistic variables, namely Low, Medium, and High. Residual Energy Level holds linguistics variables Low, Medium, and High, whereas the Packet Delivery Rate has the following linguistic variables, namely High_Flow, Medium_Flow, and Low_Flow. The output variable Congestion_Level has the following five linguistics variables: Low, Very Low, High, Very High and Medium.

Table 1: Fuzzy Rules for measuring the Congestion_Level

Set of "IF- THEN" Rules

Rule_I: IF Buffer_Length is High and Residual_Energy_Level is High and Packet_Delivery_Rate is High-Flow THEN Congestion_Level is High. Rule_II : IF Buffer_Length is Low and Residual_Energy_Level is Low and Packet_Delivery_Rate is High_Flow THEN Congestion_Level is Low. Rule_III: IF Buffer_Length is Low and Residual_Energy_Level is Low and Packet_Delivery_Rate is High_Flow THEN Congestion_Level is Low. Rule_IV: IF Buffer_Length is High and Residual_Energy_Level is Low and Packet_Delivery_Rate is Low_Flow THEN Congestion_Level is Very High.

Rule_V: IF Buffer_Length is Low and Residual_Energy_Level is High and Packet_Delivery_Rate is Low_Flow THEN Congestion_Level is Medium.

Congestion notification phase

Once the congestion is detected, in order to prevent the worst case of more packets begin transmitted over the network all other intermediate nodes should be notified about the congestion issue. Thus in this proposed work, through the congestion notification phase, the nodes are given a warning signal about the congestion in the network. In this phase, the congestion notification mechanism based on the ICN scheme of Yi *et al.*(2008) is employed. Thus, the nodes will transmit the piggyback packet holding the congestion warning signal. The main advantage of this phase is to prevent a network form getting more congested with additional control packets

and in turn to enhance the network lifetime and energy efficiency.

Fuzzy inference system based congestion control phase

In this phase, we have developed a fuzzy based rate control scheme that will perform congestion control and prevent its subsequent issues in network such as delay in transmission, inefficient energy consumption, reduced network lifetime, etc. Moreover, the primary objective of this phase is to control the congestion by initiating the intermediate nodes to reduce its transmission rate once after receiving the congestion notification signal. Hence, this may leads to prevent further more congestion in network and enhance the network performance. Once again Fuzzy Inference System is utilized in this phases. Congestion Control phases is carried out by considering the following two metrics namely Congestion_Level and Residual Energy. Hence each node evaluates the Transmission_Rate_Control by using the fuzzy logic to determine the nodes probability to control the data transmission rate. Table 2 shows the fuzzy rules for evaluating the parameter Transmission_Rate_Control. The two fuzzy input variables and its corresponding linguistics variables for Transmission_Rate_Control are shown below.

Congestion_Level has the following linguistic variables, namely Low, Medium and High. The Residual_Energy_Level holds linguistic variables Low, Medium, and High. The output variable Transmission_Rate_Control has the linguistics variables Increase_Low, Increase_High and Decrease.

 Table 2:
 Fuzzy Rules to evaluate the Transmission_Rate_Control

Set of "IF- THEN" Rules
Rule_I: IF Congestion_Level is Low and Residual_Energy_Level is Low and THEN Transmission_Rate_Control is Decrease.
Rule_II: IF Congestion_Level is Low and Residual_Energy_Level is Medium and THEN Transmission_Rate_Control is Increase.
Rule_III: IF Congestion_Level is Medium and Residual_Energy_Level is High and THEN Transmission_Rate_Control is Increase.

In our fuzzy inference system, the Congestion_Level and Transmission_Rate_Control are determined using both the triangular and trapezoidal membership functions. The triangular membership functions are used to represent the intermediate variables whereas for boundary variables the trapezoidal membership functions is used.

The triangular membership functions and trapezoidal membership functions (Rezaee *et al.*, 2014) are calculated using Equation (9) and Equation (10), respectively.

$$A = \begin{cases} 0, x \le a1 \\ \frac{x - a1}{b1 - a1}, a1 \le x \le b1 \\ \frac{c1 - x}{c1 - b1}, & b1 \le x \le c1 \\ 0, & x \le a2 \end{cases}$$
...(9)

$$A = \begin{cases} 0, x \le a^2 \\ \frac{x-a^2}{b^2-a^2}, a^2 \le x \le b^2 \\ \frac{d^{2-x}}{d^2-c^2}, c^2 \le x \le d^2 \\ 0, d^2 \le x \end{cases}$$
...(10)

The finale step in FIS is the defuzzification. The primary objective of this step is to obtain the crisp output equivalent to the fuzzy output. Moreover, in this work we have used the Center of Area COA method for defuzzification. This is measured by using the Equation (11).

$$COA = \frac{\int \mu_A(x) x dx}{\int \mu_A(x) dx} \qquad \dots (11)$$

where $\mu_A(x)$ denotes the fuzzy values for the membership functions.

Illustration

In this section we have illustrated the fuzzy based congestion_level estimation mechanism in detail, with a suitable example.

For example, assume that the Packet_Delivery_Rate is 50, Buffer Length is 0.35 and Residual Energy Level (REL) is 0.9. Table 3 shows the fuzzy variables and their corresponding values. Table 4 shows the corresponding minimum and maximum values for the fuzzy variables.

The input value Z = 50 indicates the packet delivery rate of the node. Thus by setting the trapezoidal member function to low, we get

$$f(z; k, l, m, n) = max \left(min \left(\frac{z - k}{l - k}, 1, \frac{n - z}{n - m} \right), 0 \right)$$

$$f(162; 70, 135, 162, 165) = max \left(min \left(\frac{50 - 0}{1 - 0}, 1, \frac{65 - 50}{65 - 40} \right), 0 \right)$$

$$= max(min(50, 1, 0.6), 0)$$

$$= max(0.6, 0)$$

$$= 0.6$$

Similarly, by setting the triangular and trapezoidal member functions for buffer length to low and medium, respectively, we get,

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$$Z = 0.35 \text{ (Low)}$$

$$f(z; k, l, m, n) = max \left(min \left(\frac{z - k}{l - k}, 1, \frac{n - z}{n - m} \right), 0 \right)$$

$$f(162; 70, 135, 162, 165) = max \left(min \left(\frac{0.35 - 0}{1 - 0}, 1, \frac{0.40 - 0.35}{0.40 - 0.30} \right), 0 \right)$$

$$= max (min (0.35, 1, 0.5), 0)$$

$$= max (0.35, 0)$$

Z = 0.35 (Medium)

$$f(z; k, l, m) = max \left(min \left(\frac{z - k}{l - k}, \frac{m - z}{m - l} \right), 0 \right)$$

$$f(z; k, l, m) = max \left(min \left(\frac{0.35 - 0.25}{0.45 - 0.25}, \frac{0.70 - 0.35}{0.70 - 0.45} \right), 0 \right)$$

$$= max (min (0.5, 1.4), 0)$$

$$= max (0.5, 0)$$

$$= 0.5$$

= 0.35

Similarly, by setting the triangular and trapezoidal member functions for residual energy level to medium and high, respectively, we get,

$$Z = 0.9 \text{ (Medium)}$$

$$f(z; k, l, m) = max \left(min \left(\frac{z - k}{l - k}, \frac{m - z}{m - l} \right), 0 \right)$$

$$f(z; k, l, m) = max \left(min \left(\frac{0.9 - 0.1}{0.4 - 0.1}, \frac{0.9 - 0.8}{0.9 - 0.4} \right), 0 \right)$$

$$= max(min(2.66, 0.2), 0)$$

$$= max(0.2, 0)$$

$$= 0.2$$

Z = 0.9 (High)

$$f(z;k,l,m,n) = max\left(min\left(\frac{z-k}{l-k},1,\frac{n-z}{n-m}\right),0\right)$$

$$f(z; k, l, m, n) = max \left(min \left(\frac{0.9 - 0.4}{0.8 - 0.4}, 1, \frac{1.2 - 0.9}{1.2 - 1} \right), 0 \right)$$
$$= max(min(1.25, 1, 1.5), 0)$$
$$= max(1, 0)$$
$$= 1$$

Applying the obtained values in fuzzy rules 1 to 4,

 $Rule 1: \min(0.6, 0.35, 0.2) = 0.2$ $Rule 2: \min(0.6, 0.35, 1) = 0.35$ $Rule 3: \min(0.6, 0.5, 0.2) = 0.2$ $Rule 4: \min(0.6, 0.5, 1) = 0.5$

The maximum value obtained from Rule 1 to Rule 4 is 0.5. Rule 4, corresponding to the maximum value, represents the medium probability of congestion_level in the node. The crisp value ranges from 0.25 to 0.7. During the defuzzification process, the corresponding values are substituted to obtain the value of the congestion_level as 0.61.

Table 3: Fuzzy variables and their input range

Fuzzy Input	Fuzzy variable	Input range
Packet_Delivery_Rate	Low	0-60
	Medium	55-170
	High	150-200
Buffer Length	Low	0-0.40
	Medium	0.25-0.70
	High	0.60-1
Energy	Low	0.0 - 0.5
	Medium	0.1 - 0.8
	High	0.5 - 1
Fuzzy Outputs	Fuzzy variable	Input Range
Congestion_Level	Low	0.01-0.05
	Medium	0.25-0.7
	High	0.5-1
Transmission_Rate_	Increase_Low	0.01-0.06
Control	Increase_High	0.35-0.7
	Decrease	065-1

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Table 4:	Minimum	and	maximum	value	for	fuzzy	variabl	es

Fuzzy variables	Minimum crisp value	Maximum crisp value
Packet_Delivery_Rate	0	200
Buffer Length	0	1
Energy	0	1
Congestion_Level	0.01	1
Transmission_Rate_Control	0	1

RESULTS AND DISCUSSION

The proposed work has been tested using MATLAB software because the MATLAB fuzzy tool box consists of all the fuzzy membership functions. Hence, MATLAB is the most efficient and convenient software to carry out the experiments of the proposed work EE- FBCDCA. The experiment were carried out with different nodes, starting from 100 nodes up to 200, deployed over network area of (500 x 500) m². The simulation parameters are listed in Table 5.

Table 5: Simulation parameters

Parameters	Value
Simulation Area	500 m x 500 m
Number of Nodes	200 Nodes
Initial Node Energy	0.5 J
Packet Size	512 Bytes

Packet delivery rate

Figure 2 shows the packet delivery rate of proposed algorithm, and it is compared with other existing algorithms. From Figure 2, it is observed that the packet delivery rate of EE- FBCDCA is greater when compared to other existing work. This is because, in the proposed work, the congestion control phase is initiated to control the packet transmission rate of the intermediate nodes, to control the congestion level in the network. Thus, in EE- FBCDCA congestion control phases play a vital role to control the packet loss ratio and hence the network efficiency is enhanced.



Figure 2: Packet delivery rate

Network lifetime

Figure 3 illustrates the network lifetime comparative analysis of the EE- FBCDCA and other existing algorithms. From Figure 3, it is observed that the proposed work shows better network life than other algorithms. This is because, in the proposed work, once congestion is detected, the packet forwarding rate of the nodes in the network is reduced. This may prevent furthermore congestion and loss of data in the network. Thus, EE- FBCDCA enhances the network lifetime and improves the performance of the network.



Figure 3: Network lifetime

Reliability

Figure 4 depicts the reliability comparison between the EE-FBCDCA and other existing algorithms. Reliability is the network parameter that estimates the network lifetime for a given period of time, say 't'. From

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Figure 4 it is observed that the proposed algorithm gives a high reliability rate when compared with other existing algorithms. This is because in EE- FBCDCA, further congestion will be avoided by activating the congestion notification phase and hence the packet drop in the network will be prevented. Therefore the reliability rate is improved in the proposed work and network performance is enhanced.



Figure 4: Reliability



Figure 5: Packet drop rate

Packet drop rate

Figure 5 illustrate the comparative analysis of the packet drop rate of the EE-FBCDCA and other existing algorithms. The packet drop rate is measured by considering the formula given below, Equation (12).

$$PacketDropRate = \frac{P_{\alpha}}{P_{\alpha} + P_{\beta}} \qquad \dots (12)$$

Where P_{α} represents the number of packets dropped by a node P_{β} represents the number of packets received by a node

From Figure 5 it is observed that the packet drop probability of the proposed work is less than the that of other existing algorithms. This is because in the proposed algorithm, both the fuzzy based congestion detection and control phases ensure reduced packet loss rate. Thus, the proposed work yields improved network lifetime with decreased packet drop rate.

Delay

Figure 6 shows the comparative analysis of delay of the EE-FBCDCA with the existing algorithms. End to end delay is defined as the average time taken by the packet to reach the destination node successfully. Thus, minimal delay indicates a better performance of the network. From Figure 6 it is observed that the delay of the proposed work is less than that of the other existing algorithms. This is because the EE-FBCDCA algorithm contains a very efficient congestion control process that may lead to reduced delay in the network.



Figure 6: Delay

Energy consumption

Figure 7 depicts the average energy consumption per bit comparative analysis of the EE- FBCDCA and other existing algorithms. The average energy consumption per bit is defined to be the energy required to transmit a data packet successfully from the source to the destination. From Figure 7, it is clear that the proposed work consumes less energy than existing algorithms. This is because the EE-FBCDCA uses fuzzy based congestion control phases that reduce the amount of data transmitted by the intermediate nodes. Since the congestion is controlled to greater extent that may lead to reduce the data drop rate and therefore energy conservation is improved.



Figure 7: Energy consumption

Throughput

Figure 8 shows the throughput analysis of the proposed work and other existing algorithms. The experiments are conducted for a varying number of nodes, from 200 to 1000. From Figure 8 it is observed that network throughput increases continuously with an increase in the number of nodes. The throughput in the proposed work is improved by applying a fuzzy based congestion detection and control phase that reduces the packet loss rate. Hence, in EE- FBCDCA the number of data packets that are transmitted successfully is more, and in turn network throughput is improved to a greater extent.



Figure 8: Throughput

Residual energy

Figure 9 depicts the residual energy analysis of the EE-FBCDCA algorithm with other existing algorithms. From Figure 9 it is clear that the proposed work gives more residual energy than other algorithms. This is due to the fuzzy based congestion detection process utilized in the proposed work. Hence the packet drop rate is reduced, and so, more energy is conserved. Therefore, the proposed work gives high residual energy.



Figure 9: Residual energy



Figure 10: Network performance analysis

Network performance analysis

Figure 10 depicts the network performance analysis of the EE-FBCDCA algorithm and other two existing algorithms. From Figure 10 it is clear that the EE-FBCDCA algorithm gives better network performance in terms of the metrics Packet Delivery Rate (PDR), Delay, and Network Lifetime.

CONCLUSION

In this paper, a novel energy efficient fuzzy based congestion detection and control algorithm has been

proposed with the objective of detecting and controlling congestion in WSN for enhancing the energy efficiency and lifetime of the network. Moreover, fuzzy rules are used in the process of both detection and control of congestion. Through simulation results using MATLAB, the proposed algorithm is evaluated and it has been proved that EE- FBCDCA shows better results when compared to the existing traditional processes, such as HTCCFL and FBACC, in terms of network lifetime, stability and overall performance. Thus, from the simulation results it is observed that the proposed process, EE-FBCDCA, outperforms the other existing processes in improving the lifetime of the network by 35%. Similarly, EE-FBCDCA shows 30% - 33%, and 35% - 38% better throughput in comparison to HTCCFL and FBACC, respectively. Moreover, from the proposed work, it is observed that above 78% of the residual energy in the network is sustained, which enables the transmission of data with reduced delay. Finally, from the experiments conducted it has been proved the proposed algorithm, EE-FBCDCA, gives better congestion detection and control and reduced end to end delay, as well as increased throughput, energy efficiency, and reliability.

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RESEARCH ARTICLE

Agricultural Statistics

Evaluating the potential of an open sensor network to support reservoir pre-release decision making

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Abstract: Still in most countries, reservoir flood warnings are threshold-based alerts issued when water levels exceed thresholds. This current practice of releasing water from reservoirs causes flash floods in downstream areas and increases the likelihood of dam failures and public outrage. Pre-release of water from reservoirs is therefore an important strategy for downstream flood mitigation. Hydrological models can simulate river flow with sufficient lead time. Thus, the resulting outputs can be effectively applied to pre-release decision making in the reservoirs. Since the beginning of computer-aided applications, many attempts have been made to establish a decision support system for reservoir flood control. However, this is hampered by manual stations, low quality data, high cost of software and data, unknown parameter values, and lack of expertise, especially in developing countries. Therefore, a total opensource solution combined with low-cost open-source hardware, free and open-source software, and open standards was seen as the only way to overcome reservoir-related flood risk. Moreover, research studies on open-source hardware, software, standards, and data are limited to a few case studies reporting real-time data on certain environmental parameters. Therefore, the application of integrated open-source technologies for reservoir flood control remains an unexplored area. In this background, a hydrological model powered by integrated opensource technology is presented in this research for reservoir prerelease decision making. The model was tested for the Deduru Oya watershed using the SWAT (Soil and Water Assessment Tool) toolkit. The calibration results appear to be satisfactory for both daily and hourly time intervals. Thus, this model helps to simulate the inflow of the reservoir and determine the level of reservoir gate opening.

Keywords: 4ONSE, Deduru Oya basin, open-source technologies, reservoir pre-release.

INTRODUCTION

A reservoir is an artificial or natural lake or pond which is used to collect and store water for versatile activities. In Sri Lanka, reservoirs are called tanks or 'wewa', which is the Sinhala word. Reservoir flood control measures can be broadly classified as structural and non-structural. Structural measures involve the mitigation of floods through physical constructions. In ancient Sri Lanka, floods associated with tanks were mainly controlled through the 'Ellanga gammana' system which is also known as the 'Cascaded tank-village' system (Figure 1). This system includes a network of small to large tanks in a micro or mesoscale catchment for storing, conveying, and utilizing water from an ephemeral rivulet (Bandara, 1995). It can be considered as one of the structural measures adopted by ancient Sri Lankans to create a buffer against seasonal flooding and to store water during droughts. However, in certain circumstances, the functioning of this system has been negatively affected by the filling of paddy lands and the availability of abandoned tanks.

Compared to structural methods, non-structural methods of flood control have been accepted in the present day as a more proven method for reducing flood risk and damages. This is mainly due to its long-term sustainability and minimal cost for operation and maintenance. Out of the different non-structural measures, the flood warning is the best measure to undertake for the areas that deserve prompt attention. Flood warnings differ from forecasts

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as they are issued when an event occurs or is about to occur (WMO, 2013). In the Asian context, China and Japan have established reservoir flood control systems, taking advantage of real-time hydro-meteorological data (Takeuchi et al, 1998; Guo et al, 2004). However, like most developing countries, Sri Lanka lacks a reliable weather station network, which could offer continuous near-real-time data for decision-making. Limited weather stations, costly and offline data, unavailability of parameter values, expensive modeling software, and limited resource persons are some of the major reasons which hinder the application of hydro-meteorological data for reservoir pre-release decision making. In many countries, even in Sri Lanka, reservoir flood warnings are issued under conditions where the reservoir is already at full capacity. This creates flash floods in downstream areas due to sudden release of water at a high speed.



Figure 1: Schematic diagram of a tank cascade system (Panabokke et al., 2002)

The project 4ONSE (4 times open and non-conventional technologies for sensing the environment) was launched in this setting to introduce an open-source weather station

network to support disaster warnings. IoT (internet of things) enabled technologies, open-source standards, and software have now made a significant turning point in the future of disaster warning from high-cost sophisticated devices to low-cost, open-source solutions. Evans (2011) defined the term internet of things (IoT) as a moment where more things are connected to the internet than people. Wireless connectivity and smart sensors are the two technologies that shape up the IoT network. Hence, IoT usually does the collecting of data through smart sensors and sharing them through the internet. Hart & Martinez (2015) have stated that most of the IoT-based applications and IoT-oriented research have been applied in cities and indoor environments where the relevant infrastructure facilities such as internet connectivity, accessibility, and electric power supply are available. Therefore, the most essential requirement to form a global environmental monitoring sensor network is to introduce IoT to remote environments where the IoT systems are powered by energy harvesting systems composed of sustainable energy source/s and energy storage units and wireless internet connectivity. Several of the world's most recognized companies such as IBM and HP have already started some initiatives in this respect (IBM,2010; HP, 2013; Liang & Huang, 2013).

The most cost-effective IoT applications have started to become popular after the addition of lowcost sensors, open-source hardware platforms, opensource software and standards in system development (Bitella et al., 2014; Sadler et al., 2014; Chemin et al., 2015; Formisano et al., 2015; Prescott et al., 2016; Rao et al., 2016; Saini et al., 2016). However, only a few cases of integration of these open-source platforms for monitoring various environmental parameters have been reported. Valenzuela et al. (2018) developed a turbidity data acquisition system using Arduino as open hardware and MyOpenLap free software as open-source software. Sabatini (2017) has presented an approach of step-by-step installation of an automatic weather station in remote sites. Daniele et al. (2016) have developed an open hardware device based on Arduino to monitor the soil water potential for irrigation activities. A similar kind of application was developed by Bitella et al. (2014) to monitor the soil water content integrating the soil, vegetation, and atmosphere parameters. Prescott et al. (2016) discussed a hydro-climatic monitoring station that observes six water quality parameters. Mesas-Carrascosa et al. (2015) developed an open-source hardware device to record environmental parameters and a smartphone application to analyze the data. Sadler et al. (2014) developed a low-cost environmental monitoring system that measures air temperature and

relative humidity and automatically sends the collected data to Hydrologic Information System. Samourkasidis & Athanasiadis (2014) demonstrated an automated data archival system integrated with OGC-SOS, low-cost sensors, and Raspberry Pi as open hardware. Therefore, the integration of open-source technologies for reservoir management in river basins, remains an unexplored area, especially with regard to reservoir pre-release decision making. Therefore, the main objective of this research is to evaluate the potential of such a system for reservoir pre-release decision-making with the support of a hydrological model.



Figure 2: Hydrological network, sub-catchments & sub-basins of Deduru Oya basin

MATERIALS AND METHODS

Study site

Figure 2 shows the river basin chosen to establish the 4ONSE network, the Deduru Oya river basin. It's catchment area is about 2687 km² and the length of the main channel is 115 km. There are 8 major reservoirs and 2408 minor reservoirs in the basin. Among them, Deduru Oya reservoir is the largest and main reservoir in the basin. Thus, it has been selected for testing the reservoir management decision support system. Its significant geographical location is a major factor in flood control in the lower basin. The upper catchment area can be divided into four sub-catchments based on the four stream networks originating from the Central Highlands. Among them, the reservoir receives the largest amount of water from the Deduru Oya sub-catchment.

Open-source Framework

Arduino, istSOS (Instituto Scienze della Terra Sensor Observation Service) and OGC-SOS (Open Geospatial Consortium – Sensor Observation Service) are the opensource hardware, software and standards, respectively, used in system development. The istSOS allows access to all sensor observations from a centralized location based on OGC-SOS standard. The data are visualized in istSOS at a rate of 10 min. It has an automatic data validation procedure to identify the quality of near-realtime data. This validation procedure assigns a code for each datum after the validation test.

All the 4ONSE stations were built on Arduino Mega 2560 open hardware platforms. In addition to the weather stations, several river gauges were built using the same Arduino Mega version, to measure the water levels of the streams. Each station is powered by 30W

solar panel and 12V 35Ah rechargeable battery. The sensors of the stations, measured parameters, units, accuracy, and measuring range are shown in Table 1. Cost, WMO standards, and durability are the main factors that have been considered when selecting sensors for the stations (Cannata *et al.*, 2017; Sudantha *et al.*,

2018, 2019). Figures 3 (a) and (b) shows photographs of a weather station and a river gauge developed under the 4ONSE project. Figure 4 shows the locations where the 27 weather stations and 06 river gauges are deployed. The quality of the 4ONSE data was checked using some reference stations' data at daily and 10 minutes intervals.

Sensor	Parameter	Unit	Accuracy	Measuring range
DS18B20	Temperature	Degrees Celsius (°C)	±0.5°C	-10 to 85 °C
BME280	Relative Humidity	Percentage (%)	±3%	0% - 100%
BME280	Barometric pressure	Hectopascals (hPa)	±1 hPa	300 – 1000 hPa
ZHIPU wind speed sensor	Wind speed	Meters per second (ms ⁻¹)	$\pm 1 \text{ m/s}$	0-32.4 m/s
Anemometer 485 wind direction sensor	Wind direction	Degrees	±3°	16 different directions and any angle values can be identified
6465 Davis AeroCone Rain Gauge with Mountable Base	Precipitation	Millimeters (mm)	0.2mm	N/A
BH1750 light sensor module	Light intensity	Lux (lx)	1.44 times, Sensor Out / Actual lx	(1-65535lx)
Soil moisture module	Soil moisture	Percentage (%)	±2%	0 to 22%
River gauge module MB7062 XL-MaxSonar-WR1 Ultrasonic sensor	Water Level	Meters	±0.5cm	0 - 10m

Table 1: Sensors of the 4ONSE stations



Figure 3: 4ONSE (a) weather station and (b) river gauge

In this study, the SWAT (Soil and Water Assessment Tool) open-source tool has been used to develop the hydrological model. The model calibration part has been done using the SWAT-CUP open-source tool. Figure 5 shows the complete open-source framework used to develop the hydrological model to support pre-release decision making in the reservoir.

Development of hydrological model

The SWAT model has shown its capability in regionalscale hydrological modelling for simulation of river discharge (Ghoraba, 2015; Bailey *et al.*, 2016; Zhang *et al.*, 2016; Rafiei *et al.*, 2017; Warusavitharana, 2020). The model was developed using QSWAT plugin version 1.9 (Dile *et al.*, 2016) embedded in QGIS software. The QSWAT plugin successfully works in QGIS 2.6.1 Brighton version. The processing of input data such as DEM (Digital Elevation Model), land use, and soil was performed using QGIS Brighton version. SWAT's algorithms for infiltration, surface runoff, flow routing, impoundments, and lagging of surface runoff have been modified to allow flow simulations with a sub-daily time interval as small as one minute, and evapotranspiration, soil water contents, base flow, and lateral flow are estimated daily and distributed equally for each time step (Jeong *et al.*, 2010). Therefore, using precipitation on a sub-daily basis and the other input data (relative humidity, temperature, wind speed and solar radiation) on a daily basis was sufficient.

Since SWAT is a continuous hydrological model, it requires a warm-up period of several years to stabilize

the model. Since 4ONSE is a newly deployed sensor network, the required data for the model's warm-up period has been estimated through SWAT's weather generator. The required monthly statistical data to operate the weather generator has been calculated using CFSR (Climate Forecasting System Reanalysis) data. Table 2 shows the models and methods used in the SWAT weather generator to estimate the missing weather data.



Figure 4: Locations of the 4ONSE stations in Deduru Oya basin



Figure 5: 40NSE open-source framework for reservoir pre-release decision making

Climatic variable	Model / Method
Daily precipitation	Model developed by Nicks (1974)
Sub-daily precipitation	Double exponential function
Daily maximum and minimum air temperature	Normal distribution
Daily average relative humidity	Triangular distribution
Daily solar radiation	Normal distribution
Daily mean wind speed	Modified exponential equation

 Table 2:
 Models and methods used in SWAT weather generator

Model calibration and uncertainty analysis

The model calibration and uncertainty analysis has been accomplished through the SWAT-CUP (SWAT-Calibration and Uncertainty Procedures) public domain programme. The four upper sub-catchments in the Deduru Oya basin have been separately calibrated in this study by developing four separate SWAT models. However, this research extends the results of the model associated with the Deduru Oya sub-catchment.

Primarily, the effectiveness of the model at daily timestep was checked using the default parameter values in the tool. As there was a substantial difference between the simulated flow and the observed flow, the model was regionalized first. Generally, the terms regionalization, parameterization, and calibration have similar meaning, which is adjusting the parameter values to reduce the difference between a simulated result and observed values. The parameterization / regionalization scheme of SWAT-CUP tool is as follows:

x_ < parname >.< ext > _ < hydrogrp > _ < soltext > _ < landuse > _ < subbasn > _ < slope >

The components in the scheme namely, *parname*, *ext*, *hydrogrp*, *soltext*, *landuse*, *subbasn* and *slope* represents the name of the parameters (as it appears in SWAT), the extension of the parameter, soil hydrologic group, land use type, sub-basin number, and the slope, respectively. As per the scheme, x_{-} represents the type of change to be applied to the parameter. The type of changes represented by x_{-} are as follows:

l) V_{-} - replacing the existing parameter

2) A_- - (existing parameter) + (given value)

3) R_- - (existing parameter) × (1+ given value)

In this study, several rules have been applied when selecting the appropriate x_{type} . The usual principle in the SWAT_CUP tool to use type R_ for spatial parameters (*e.g.*, land use and soil type). Moreover, considering the

ease of examining a wide range of values, type $R_{\rm was}$ applied to parameters with a large range. Type $V_{\rm was}$ applied to all the other parameters.

sensitive/dominant The parameters the of hydrological model were identified first through the OAT (One-at-a-time) sensitive analysis in SWAT-CUP. It shows the sensitivity of each parameter if all other parameters are kept constant. The model calibration and uncertainty analysis were performed according to the SUFI 2 (Sequential Uncertainties Fitting Version 2) algorithm integrated in SWAT. The fitness of the model was assessed statistically using the Nash-Sutcliffe Efficiency (NSE) method (Nash, 1957) (Equation 1), Goodness of fit linear regression model (R²) (Equation 2), P factor (percentage of observed data simulated in the model) and R factor (average thickness of the 95% predication uncertainty (95PPU) divided by the standard deviation) (Equation 3). An NSE value between 0 and 1, $R^2 > 0.5$, P-factor > 70% and R-factor of around 1 are considered acceptable.

$$Maximize : NSE = \frac{\sum_{i=1}^{n} (O_i - \bar{O})^2 - \sum_{i=1}^{n} (P_i - O_i)^2}{\sum_{i=1}^{n} (O_i - \bar{O})^2} \dots (1)$$

$$R^{2} = \frac{\sum_{i=1}^{n} (O_{i} - \bar{O})(P_{i} - \bar{P})}{\sqrt{\sum_{i=1}^{n} (O_{i} - \bar{O})^{2}} \sqrt{\sum_{i=1}^{n} (P_{i} - \bar{P})^{2}}} \dots (2)$$

$$R - factor = \frac{\frac{1}{n_j} \sum_{t_{i=1}}^{n_j} (x_s^{t_i > 5.5\%} - x_s^{t_i > 2.5\%})}{\sigma_{o_j}} \qquad \dots(3)$$

where *n* is the number of observations, O_i is the *i*th observed value, \overline{O} is the mean observed value, P_i is the *i*th model-predicted value, \overline{P} is the mean model-predicted value, $x_s^{t_i, 27.5\%}$ is the upper boundary and $x_s^{t_i, 2.5\%}$ is the lower boundary of 95PPU.

As a summary, Table 3 lists all the open-source tools used in developing the hydrological model to support reservoir pre-release decision making.

Open-source tool	Function	Link
QGIS Brighton	 Analyse vector and raster data 	http://qgis.org/downloads/QGIS-OSGeo4W-
Version	GIS interface to run the SWAT model	2.6.1-1-Setup-x86.exe
QSWAT	 SWAT plugin used to run the model in QGIS software 	https://swat.tamu.edu/software/qswat/
SWAT-Editor	 Reading project databases 	https://swat.tamu.edu/software/swat-editor/
	 Generating missing weather data 	
	Executing SWAT run	
	Calibrating the model	
SWAT-CUP	 Identifying the dominant parameters & their ranges 	https://www.2w2e.com/home/SwatCup
	Calibrating the model	
	 Validating the model 	
istSOS	 To view and download the data of 4ONSE stations 	https://geoservice.ist.supsi.ch/4onse/admin/

RESULTS AND DISCUSSION

Before applying the 4ONSE meteorological data in the model, its quality was checked using data from some reference stations in Sri Lanka. Table 4 shows the coefficient of determination of all key meteorological parameters measured at 10 minutes and daily intervals.

Table 4:	Coefficient	of	determination	(\mathbb{R}^2)	between
	40NSE stat	ions	and reference s	station	s

Interval	R ² value
10 minutes	0.9678
Daily	0.9921
10 minutes	0.7292
Daily	0.7784
10 minutes	0.7448
10 minutes	0.9184
Daily	0.9811
10 minutes	0.9771
Daily	0.9929
10 minutes	0.9889
Daily	0.9916
	Interval 10 minutes Daily 10 minutes Daily 10 minutes 10 minutes Daily 10 minutes Daily 10 minutes Daily

After running the model using daily 40NSE data with SWAT's default parameter values, several key features of

simulated flow and the actual flow were observed. They are:

- 1. Simulated flow peaks are higher than actual flow
- 2. The baseflow of the simulated flow is higher than the actual flow
- 3. The discharge in the simulated flow shifted to the left

To correct the above issues, the model was first regionalized. Table 5 shows the related dominant/ sensitive parameters identified through the OAT analysis and applied modifications to correct the above deviations.

In the SWAT model, CN2 (curve number), CANMX (maximum canopy storage), SOL AWC (soil available water content) and ESCO (soil evaporation compensation factor) are the parameters that contribute to the peak fluctuation of streamflow. However, CANMX was the only parameter that showed sensitivity to high peaks, especially for the dominant land use categories of coconut, rice, low-density residential, and rubber. The CANMX parameter represents the maximum amount of water that trees can be hold. This value is zero by default in the SWAT database. The peaks can be reduced by increasing the CANMX value of dominant land use classes. Since CANMX is a parameter that introduces water into the system it was taken separately and 50 runs performed, to obtain optimal values related to the four dominant land use categories. Table 6 shows the sensitive parameters and their values for daily and hourly intervals obtained during model calibration.

Observation	Reason	Related parameters	Applied modification
High peaks	High surface flow	CN2, SOL_AWC, ESCO, CANMX	Increase CANMX
Model over predicts the flow	High baseflow and/or little evapotranspiration	GWQMN, GW_REVAP, REVAPMN	Increase GWQMN & GW_REVAP
Discharge was shifted to left	Simulated flow leads the actual flow	SLOPE, OV_N, SLSUBBSN, CH_N2	Increase CH_N2

Table 5: Regionalized parameters

 Table 6:
 Sensitive parameters and their value ranges related to daily and hourly intervals.

Sensitive parameter	Description	Type of change	Value range – daily interval	Value range – hourly interval
CN2	Initial SCS runoff curve number for moisture condition II)	R_	0.14 - 0.25	(-0.3) – 0.1
SOL_AWC	Available water capacity of the soil layer	R_	(-0.14) – 0.03	Not sensitive
ESCO	Soil evapotranspiration compensation factor	V_	0.68 - 0.95	Not sensitive
SOL_BD	Moist bulk density	R_	(-0.12) – 0.06	(-0.08) - 1.77
MSK_X	Weighting factor for wedge storage	V_	0 - 0.14	0 - 0.1
MSK_CO2	Muskingum coefficient for low flow	V_	(-0.04) – 0.59	0 - 8.1
MSK_CO1	Muskingum coefficient for normal flow	V_	0.96 - 1.31	1.0 - 5.2
ALPHA_BF	Baseflow alpha factor	V_	(-0.17) – 0.23	0 - 0.2
SOL_K	Saturated hydraulic conductivity	R_	(-1.21) – 0.06	(-0.47) – (-0.04)
CH_K2	Effective hydraulic conductivity in main channel alluvium	V_	4.27 - 6.73	1.1 – 27.3
CH_N1	Manning's "n" value for the tributary channels	V_	(-0.13) - 0.20	(-0.3) – 0.7
CH_N2	Manning's "n" value for main channel	V_	0.04 - 0.06	0 - 0.7
GW_DELAY	Ground water delay time	V_	38.90 - 89.94	Not sensitive
GW_REVAP	Groundwater "revap" coefficient	V_	0.24 - 0.45	0.1 - 0.2
GWQMN	Threshold depth of water in the shallow aquifer required for return flow to occur	R_	3.14 - 4.32	0.8 - 2.0
SURLAG	Surface runoff lag coefficient	V_	Not sensitive	(-0.5) – 1.0

Except the SOL_AWC, ESCO, GW_DELAY, and SURLAG parameters, all the other parameters were received as sensitive for both daily and hourly time intervals for the Deduru Oya sub-catchment. For hourly simulation, the effect of ESCO, SOL_AWC, and GW_DELAY was not significant while SURLAG parameter was significant to simulate hourly flows.

Figure 6 shows the simulated and actual inflow in the Deduru Oya sub-catchment, after model calibration. The statistical results regarding P factor, R factor, R², and NSE presented in Table 7 appear to be satisfactory. The Davis rain gauge used at 4ONSE weather stations typically has a margin of error of $\pm 4\%$ for rainfall rates up to 50 mm/hour and $\pm 5\%$ for rainfall rates in the 50 mm/hr to 100 mm/hr range. This is the main reason why some peaks do not reach the desired level.

Accordingly, the calibrated parameter values of the model can be easily applied to the model to simulate the inflow of the Deduru Oya reservoir. The time interval of the model can be changed according to the needs of the decision-makers who decide on water pre-release. Customization of weather data and optimization of parameters should be done according to the selected time interval. Moreover, the application of near real-time weather data in the model is more valid when the time required for decision-making is greater than the time required to concentrate water from the upper basin to the reservoir. The model developed in this research cannot be directly used for flood modelling. It only helps to simulate the flow into the reservoirs / tanks and to determine the level of opening of the reservoir gates, which helps in reducing downstream flood inundation. However, the inflows simulated by the model can be applied to SWAT's reservoir management tool to estimate reservoir capacity and determine reservoir outflow.



Figure 6: Simulated flow and Observed/Actual flow

 Table 7:
 Statistical results of the hydrological model

Period	Interval	P factor	R factor	R ²	NSE
1 st August to 6 th October 2019	Daily	0.87	0.98	0.69	0.69
1 st to 15 th June 2019	Hourly	0.96	0.76	0.76	0.75
15 th to 30 th June 2019	Hourly	0.96	0.89	0.89	0.88
4th to 9th August 2019	Hourly	1.0	0.53	0.77	0.55
1 st to 15 th August 2019	Hourly	0.74	0.54	0.67	0.63
16 th to 23 rd August 2019	Hourly	0.83	0.00	0.67	0.43

CONCLUSION

Reservoir flood management applications require generating modelling results with the shortest lead time using near real-time meteorological data as input data. Due to the limitations of the existing setup, the incorporation of near real-time and quality hydro-meteorological input data to produce accurate hydrological estimates has never been used in Sri Lanka. Thus, this study intends to introduce a cost-effective decision support system to reservoir pre-release decision making. It is built entirely using several open source technologies for model implementation and data entry. QGIS Brighton version, QSWAT, SWAT Editor and SWAT-CUP are the opensource tools used to implement the hydrological model and Arduino, istSOS and OGS-SOS are the open-source technologies used to build meteorological stations and feed data to the model. Estimating parameter values for different time periods is tedious, time consuming, and labour and capital intensive. Therefore, the modelling approach presented in this research avoids the need for pre-specified parameter values and allows users to determine them at any time. The approach presented in this study is more appropriate for simulating reservoir inflow. If the basin has any series of tanks/cascade systems, the daily and sub-daily discharge flow of the upper basin tanks should be considered while developing the model. Furthermore, this study introduces a new approach for conducting hazard warnings in more remote environments where environmental processes cannot be realistically observed or studied due to the lack of access and facilities.

Conflict of interest statement

The authors declare that they have no conflict of interest

Data availability statement

The data that support the findings of this study are available from the corresponding author, [E.J. Warusavitharana], upon reasonable request.

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RESEARCH ARTICLE

Ophthalmic Genetics

High prevalence of glaucoma-associated CYP1B1 mutation (p.G61E) in primary congenital and open angle glaucoma patients in Pakistan

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Abstract: The purpose of this research study was to investigate the prevalence of the p.G61E variant of the CYP1B1 gene among primary congenital and open angle glaucoma (PCG and POAG) patients in the province of Punjab, Pakistan. A total of 112 POAG and 50 PCG patients were enrolled in this study. Detailed clinical examination was carried out on all patients. Screening of G61E was done by direct Sanger sequencing. Different in silico tools, e.g., Clustal Omega, PSIPRED, and Franklin tools were used to check the conservation, secondary structure, and pathogenicity, respectively, of this variant. Sanger sequencing of the whole CYP1B1 gene revealed a homozygous missense transition, c.182G>A, p.G61E in 25/50 (50%) of PCG and in 42/112 (37.5%) of POAG cases, which co-segregated with the disease phenotype. This study revealed that p.G61E is the relatively major contributor of PCG. However, 42 POAG patients harbouring the G61E mutation showed moderate to severe phenotype, suggesting the genetic heterogeneity of this variant in Pakistani population.

Keywords: Juvenile open angle glaucoma, intra ocular pressure, latent transforming growth factor-beta protein 2, primary congenital glaucoma.

INTRODUCTION

Glaucoma is a heterogeneous group of visual neuropathies which leads to optic nerve damage and permanent loss of vision if left untreated (Allingham *et al.*, 2009). Primary congenital glaucoma (PCG) is a more severe form of the disease categorized by atypical development of trabecular meshwork and elevated intraocular pressure (IOP) at the time of birth or within the first three years of life, which manifest itself with photophobia, excessive tearing, and buphthalmos (Liu & Allingham, 2017). PCG mostly shows autosomal recessive form of inheritance (Firasat et al., 2008). In the Western region, the incidence of PCG is 1 in 10,000-20,000 live births while in the Middle Eastern region the incidence is 1 in 2500 to 8200 (Lewis et al., 2017). Primary open angle glaucoma (POAG) is a common cause of irreparable blindness in the world. It is an optic neuropathy characterized by open anterior chamber, complete cupping of nerve head, loss of visual field and elevated IOP (Liu & Allingham, 2017). On the basis of age of onset, POAG is subcategorized into adultonset open angle glaucoma (AOAG) and juvenile-onset open angle glaucoma (JOAG). JOAG as an idiopathic glaucoma occuring in children above three years of age and it is associated with myopia and elevated intraocular pressure with fluctuation (Kwun et al., 2016). The age of onset of JOAG is 3-40 years and shows autosomal dominant inheritance; however a few cases of autosomal recessive inheritance from Saudi Arabia and India have been reported (Khan et al., 2011). Adult-onset POAG manifest itself after the age of 40 years and shows a complex pattern of inheritance because multiple genes and environmental factors are involved (Rauf et al., 2016). The estimated prevalence of JOAG ranges from 0.38 to 2 in 100,000 in individuals between 4 and 20 years of age (Taqi et al., 2011). JOAG comprises 4% of childhood glaucoma (Kwun et al., 2016). POAG is the

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most common form of glaucoma worldwide accounting for 90% of all cases reported (Qureshi *et al.*, 2006). POAG is also the common types of glaucoma in Pakistan (Taqi *et al.*, 2011).

The CYP1B1 (OMIM 601771) gene is a major contributor in glaucoma pathogenesis (Achary et al., 2006). CYP1B encodes a member of the Cytochrome p450 superfamily, subfamily I. The CYP1B1 gene is localized in chromosome number 2 and consists of three exons out of which two are coding (López-Garrido et al., 2015). It has been reported that the CYP1B1 protein plays role in the development of trabecular meshwork, iridocorneal angle, and in the formation of the ciliary (Rauf et al., 2016). Pathogenic and potentially pathogenic mutations in CYP1B1 have been reported in JOAG, POAG, and PCG patients from various populations around the globe (Micheal et al., 2014). Mutation p.G61E has been reported previously in POAG and PCG patients from different ethnicities (Qashqai et al., 2018). Recent investigations on the functional role of the homozygous p.G61E CYP1B1 mutation in a glaucoma family have shown that this mutation disturbs the microfibrils of the extracellular matrix (ECM), making its presence less abundant, and increases more fragmented protein in affected individuals. This study has suggested that disruption in development of the ECM-trabecular meshwork leads to increased resistance to aqueous outflow, causing elevated IOP, and eventually causes development of glaucoma (Banerjee et al., 2016). The G61E mutation has been reported to be the commonest mutation in Lebanon, Iran, Egypt, and other Middle Eastern countries (Qashqai et al., 2018). This variant is a founder mutation in Middle Eastern populations (Saudi Arabia, Iran, Morocco, Kuwait, Israel, Oman, and Egypt) (Campos-Mollo et al., 2009; Li et al., 2011). In Saudi POAG patients the frequency of this variant accounted for 64.3% (Qashqai et al., 2018). Heterozygosity for the mutation p.G61E has already been reported as a cause of adult-onset POAG in two Pakistani sporadic cases (Micheal et al., 2014). In heterozygous form, p.G61E has been reported with mild phenotype in PCG and POAG patients of different ethnicities (Khan et al., 2011; López-Garrido et al., 2015; Wiggs & Pasquale, 2017). Previously, our study demonstrated that the p.G61E variant was segregated in a consanguineous Pakistani family with co-existence of JOAG and PCG in two successive generations (Bashir et al., 2015). Only one study showed heterozygous p.G61E mutation in sporadic POAG patients from Pakistan (Micheal et al., 2014). Mutation Taster and PolyPhen predicted this

variant as disease causing, and the Franklin tool qualified this variant to be deleterious.

POAG and PCG are the most prevalent types of glaucoma in Pakistan (Iqbal *et al.*, 2011). However, few studies are available on the contribution of the p.G61E mutation in the Pakistani population. In the current study, the aim was to investigate the frequency of the p.G61E variant in PCG and POAG cases.

MATERIALS AND METHODS

Selection of study subjects and clinical examination

The study followed the guidelines of the Declaration of Helsinki. This study was approved by the Ethical Review Committee of the Department of Biotechnology, Lahore College for Women University, Lahore, Pakistan with reference number 2019-07. Informed consent was obtained from all participants or from the parents in case of infants.

Patients identified as having POAG and PCG were recruited from 2017-2020 (a 3-year period) from the Mughal Eye Hospital Lahore and the eye ward of the Children Hospital and Institute of Child Health (CHICH) Punjab. CHICH is the major ophthalmology hospital recommended for congenital cases in Punjab, and patients travel from all parts of Pakistan to this hospital. 112 unrelated Pakistani POAG and 50 PCG patients from consanguineous marriage were enrolled. 196 unrelated ethnically matched controls were also recruited.

For this study, the following inclusion criteria were used for POAG and PCG cases: (1) Age at the time of diagnosis, (2) Intraocular pressure > 21 mm Hg in adults and >12mm Hg in at least one eye in case of PCG, (3) Anterior chamber angle that was open to scleral spur or ciliary body, (4) Evidence of optic nerve damage and loss of visual field consistent with glaucoma in at least one eye, and (5) cup-to-disc (C/D) ratio > 0.3 of optic nerve. Megalocornea with corneal diameter >12 mm and rupture in Descemet's membrane have been considered. Inclusion criteria for POAG: ethically matched controls (50 males, 50 females) and age ranges from 40 to 67 years were selected after ophthalmic screening, and individuals with normal range of IOP (not having antiglaucoma medication), open angles after gonioscopy examination and healthy optic disk with no history of eye disease(s) or any ophthalmic surgeries.

For PCG, ethnically matched controls (56 males, 40 females), age ranges from 2 to 15 years with normal IOP range, normal cup/disc ratio, normal corneal diameter and intact Descemet's membrane were included.

Screening of p.G61E mutation

Primers were designed for coding exon 2 of CYP1B1 with Primer3 including 50–80 bp of intronic region from the upstream and the downstream region of coding exon with Primer3 software (http://www.bioinformatics.nl/cgi-bin/ primer3plus/primer3plus.cgi). Polymerase chain reaction (PCR) amplification was performed from genomic DNA. Each reaction was performed in a final volume of 50 µL under standard conditions with 50 ng DNA in 2.5 µL of Taq buffer, $2 \text{ mM MgCl}_{2}(1 \mu L)$, $0.2 \text{ mM DNTPs}(1.5 \mu L)$, 0.24 µL forward primer, 0.24 µL reverse primer, 25 ng genomic DNA and then PCR water was added to raise its volume to 50 µL. Denaturation of PCR products was performed at 95 °C for 30 s and annealed at 55 °C for 30 s. The initial extension was performed at 75 °C for 1 min and final extension was performed at 75 °C for 5 min. PCR products were then purified by gene cleaning column (Gel Extraction Kit Thermo Scientific GeneJet USA). After purification, samples were sequenced by direct sequencing with Big Dye Terminator[®] (V 3.1) Cycle Sequencing Kit (Applied Biosystems, Foster City, CA).

In silico analysis

ClustalW alignments were performed to determine evolutionary conservation, Figure 1C. To predict secondary structure, elements from amino acid sequence PSIPRED v.3.3 were used, Figure 1D. Franklin tool classified the variant as potentially pathogenic (*https:// franklin.genoox.com/clinical-db/variant/snp/chr2-*38302350-C-T?app=assessment-tools).

RESULTS AND DISCUSSION

Table 1 shows the clinical characteristics of all PCG and POAG patients clinically examined. In 50 PCG and 112 POAG patients, the mean age in POAG was 52.8 ± 11.1 years with range 40-70 years. In PCG cases the mean age was 12.2 \pm 9.7 years and range was 1 month to 9 years. The mean age of IOP in PCG cases was 19.5 ± 6 years whereas in POAG the mean age of IOP was 24.3 ± 7.8 years (Table 1). Clinical features of POAG patients showed watery eyes with enlarged globe and hazy cornea. They felt pain, headache, itching, and blurred vision. Some of them had maintained normal IOP pressure by regular treatment and prescribed medications. Clinical features of PCG patients showed hazy cornea, photophobia, complete opacification of corneas, and excessive tearing with enlarged globe. Most of the PCG patient underwent trabeculectomy in both or one eye to maintain normal IOP pressure along with the use of prescribed medications. IOP of PCG and POAG was positively associated with glaucoma development (Figure 1 and Figure 2).

Sequencing results of CYP1B1 showed a missense mutation c.182G>A (p.G61E) in exon 2 which was homozygous in 25/50 (50%) PCG cases and in 42/112 (37.5%) POAG cases; the mutation co-segregated with the disease phenotype in 50 PCG and 42 POAG patients (Figure 2A, 2B). The variant was absent in 196 ethnically matched control chromosomes, further confirming that this mutation is pathogenic in the Pakistani population. MutationTaster and PolyPhen predicted this variant as disease causing. Franklin tool qualified this variant to be deleterious. PSIPHRED sequence plot showed that residue G61 lies in the Protein Helix region. Genetic variants in *CYP1B1* gene are major contributing factors in the pathogenesis of PCG and POAG in

 Table 1:
 Clinical parameters of PCG and POAG patients and control groups

Parameters	Status	PCG	POAG	Control	Control
				POAG	PCG
Gender	Male	30	66	50	56
	Female	20	46	50	40
Age	$Mean\pm SD$	12.18 ± 9.7	52.8 ± 11.1	65.9 ± 11.6	$13.2\pm\textbf{-}6.4$
Unit: (mth,yrs)	Range	1 mth-14 yrs	$40-70 \ yrs$	$40 \; yrs - 67 \; yrs$	2–15 yrs
Max. IOP	$Mean \pm SD$	19.46 ± 5.96	24.32 ± 7.8	15.0 ± 2.7	16.5 ± 3.7
(Unit: mmHg)				8-21	
CD ratio	Range	0.3–0.9	0.4 - 0.9	0.2–0.3	0.2–0.3

C/D: cup to disk ratio, mth: month yrs: years

various populations of the world (Qashqai *et al.*, 2018; Youngblood *et al.*, 2019; Ling *et al.*, 2020). The variant p.G61E has been found with varying frequency in different populations; however, data on genetic spectrum of this variant in Pakistani population is very limited. To explore the clinical spectrum of this variant, the present study was performed to determine the frequency and to make genotype-phenotype comparison of the p.G61E mutation associated with PCG and POAG in the Pakistani population. This genetic study was done to demonstrate the role of the p.G61E variant of *CYP1B1* gene in Pakistani PCG and POAG patients. This cohort included 162 patients (n = 50 PCG, n = 112 POAG) as well as 196 control.



Figure 1: Electrogram of G61E mutation sequence. (A) normal sequence in control sample; (B) G>A transition in patient's DNA; (C) ClustalW sequence alignment indicates conservation of p. G61E mutation in *CYP1B1, CYP1A1* and *CYP1A2* among different vertebrate species; (D) PSIPhred sequence plot indicating that residue G61 lies in Protein Helix.



Figure 2: Graph showing IOP among both groups, i.e., PCG patients (A) and POAG (B) patients.

The p.G61E mutation has already been reported to be a frequent cause of PCG and POAG in many different populations of the world including Arab, Middle Eastern and Mediterranean countries (Khan et al., 2011; Micheal et al., 2014; López-Garrido et al., 2015), whereas this variant has been reported with incomplete penetrance as well in different populations (Campos-Mollo et al., 2009 Lewis et al., 2017;). This mutation has been shown to be a 'hypomorphic' allele, having reduced enzymatic activity of the CYP1B1 protein (Campos-Mollo et al., 2009). In previous studies, it was reported that the homozygous p.G61E mutation was pathogenic in PCG patients of Saudi Arabia (Abu-Amero et al., 2011). In a recent study from Pakistan, this variant has been identified in homozygous state in a PCG patient belonging to a consanguineous family. However, in some cases the p.G61E mutation may exhibit decreased penetrance which has been reported previously in Spanish and Saudi Arabian patients (Campos-Mollo et al., 2009; Youngblood et al, 2019). In a cohort of Saudi POAG patients, the p.G61E variant accounted for 64.3% (Abu-Amero et al., 2011). However, in another study of the Saudi population, the mutation was reported in heterozygous form and accounted for 4% of POAG cases and 2% of controls (Abu-Amero et al., 2011). Mutation p.G61E has been the most frequent mutated CYP1B1 allele in Iranian PCG patients. It is also the most commonly detected mutation among the POAG patients (Qashqai et al., 2018). In adult-onset POAG patients, the variant p.G61E is mostly reported in heterozygous and compound heterozygous state, with a milder disease phenotype (López-Garrido et al., 2015). In homozygous state the p.G61E variant showed severe phenotype in Indian and Saudi Arabian PCG patients. In Saudi JOAG patients, this mutation is reported to result in a severe phenotype in recessive form. In a report from the Pakistani population, the heterozygous p.G61E variant was reported in sporadic cases of POAG (Micheal et al., 2014). Moreover, in our previous study from Pakistan we reported the homozygous p.G61E variant, which, co-segregating in two generations of JOAG and PCG patients from a consanguineous family, showed a severe phenotype with variable penetrance of this p.G61E variant (Bashir et al., 2015).

The residue p.G61E lies in the functional part of a very highly conserved hinge region of the *CYP1B1* protein, from which region the polypeptide chain shows a sharp turn (Panicker *et al.*, 2004). It is present right after the N-terminus of a region rich in proline. The proline–proline–glycine–proline motif might make the

link between the membrane binding N-terminus and the globular region of the P450 protein. The change of glycine to glutamic acid at position 61 causes a complete loss of haem binding, resulting in minimal catalytic activity of the enzyme and results in disease manifestation (Panicker *et al.*, 2004). Furthermore, MutationTaster, SIFT, Franklin tool, and PolyPhen predicted this variant as disease causing.

In this study, screening for mutation p.G61E in the CYP1B1 gene revealed that 50% of PCG cases and 37.5% of POAG cases carried this mutation in homozygous state. The frequency of the p.G61E variant in PCG cases is higher in the Pakistani population as compare to Chinese, Iranian, and Spanish populations (López-Garrido et al., 2010; Safari et al., 2016; Abu-Amero et al., 2018). These variations in mutational frequency may be due to differences in geographical region, and epigenetic factors such as age, gender, race, and family history also contribute in causing PCG and POAG. IOP and CD ratio of PCG cases in p.G61E positive patients was high, resulting in opacification of cornea and need of surgical interventions. The patients affected with PCG showed severe phenotype indicating complete penetrance of this mutation in congenital cases. However, 19 patients affected with POAG showed mild phenotype whereas 23 cases showed severe phenotype, which showed the genetic heterogeneity of this variant in POAG.

CONCLUSION

We conclude that the *CYP1B1* gene is important in the aetiology of PCG as well as POAG, in the Pakistani population. This study reveals that p.G61E is more common among primary congenital glaucoma in the Pakistani cohort. However, the phenotypic variability associated with p.G61E variant in POAG patients of our population is possibly the effect of genetic modifiers or epigenetic factors that may modify the manifestation and variable penetrance. Further detailed studies applying whole-exome sequencing on the DNA of a large number of severely affected cases and those of less affected patients could reveal further genetic modifiers.

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RESEARCH ARTICLE

Environmental Toxicology Effect of cylindrospermopsin on the hepatotoxicity in wistar rats

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Abstract: A naturally-derived cyanotoxin, cylindrospermopsin (CYN), present in freshwater systems, poses a threat to human health due to potential hepatotoxicity. The present study aimed to determine the hepatotoxic effects of CYN on male Wistar rats. Following ethical clearance, 35 rats were divided into five groups. Test groups were administered with pre-prepared solutions to provide three doses of CYN: 0.175 µg/kg, 0.140 µg/kg, and 0.105 µg/kg. Well-water collected from Padaviya (0.161 µg/kg of CYN) was given to the environmental-exposure group (EN) and distilled water was administered to the control. The total duration of exposure was 90 days. Blood samples were collected at 0, 7, 14, 28, 42, 60, 90 days. Aspartate amino-transferase (AST), aspartate alaninetransferase (ALT), and full blood count (FBC) were analyzed. At 90 days, hepatic tissue samples were taken for histology. The mean body weight of the treated and control groups of rats gradually increased until the 90th day. A statistically significant reduction (p < 0.05) in increment of body weights was observed in CYN-treated rats, compared to the control, at 12th and 13th weeks. Relative weights of the livers of treated groups were significantly lower (p < 0.05) than those of the control group. The highest AST and ALT concentrations were recorded in rats given a CYN dose of 0.175 µg/kg. Haematology revealed a significant (p < 0.05) reduction in monocyte and lymphocyte counts in the EN group given well-water compared to the control. Ballooning degeneration, Kupffer cell hyperplasia, lobular haemorrhage and necrosis, perivenular inflammation, and sinusoidal congestion were observed in the livers of treated groups. The findings show that prolonged exposure to CYN contaminated water leads to hepatotoxicity in Wistar rats.

Keywords: Cylindrospermopsin (CYN), hepatotoxicity, histopathology, Wistar rats.

INTRODUCTION

Cyanobacteria are a group of over 2000 prokaryotic organisms commonly named 'blue-green algae' (Woese, 2002). Several genera of cyanobacteria produce two major toxin groups, the microcystins (MCs) and cylindrospermopsin (CYN). The Microcystis spp. and Cylindrospermopsis spp. of cyanobacteria grow well in a wide range of fresh, brackish, and marine habitats (Sethunga & Manage, 2010; Kulasooriya, 2011), dug wells (Abeysiri et al., 2018a, 2018b; Abeysiri & Manage, 2018), terrestrial ecosystems (Codd et al., 2005), extreme habitats of hypersaline localities (Chatchawan et al., 2011), hot springs (Wijesekara & Manage, 2017; Sadeepa et al., 2019a; 2019b) and arid deserts (Kulasooriya, 2011). Approximately 40 cyanobacteria species have been recorded as potential producers of cyanotoxins (Merel et al., 2013; Paerl et al., 2013). Cyanotoxins are secondary metabolites categorized chemically as cyclic peptides and alkaloids with a stable chemical structure (Zanchett & Oliveira-Filho, 2013). These are recognized as a potential hazard in drinking water worldwide, resulting in organ toxicity in humans and animals (Falconer & Humpage, 2005).

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CYN, a cyanotoxin contaminant of water supplies, is produced by freshwater, non-bloom-forming filamentous cyanobacteria; Cylindrospermosis raciborskii, Anabaena bergii, Anabaena lapponica, Lvngbva wollei, Aphanizomenon flosa-aquae, Aphanizomenon ovalisporum, Umezaka natans, and Raphidiopsis curvata (Falconer & Humpage, 2005; Sethunga & Manage, 2010; Mazmouz et al., 2011). C. raciborskii and Anabaena sp. are abundantly present in lakes, reservoirs, ponds, and rivers in many tropical and subtropical areas globally, with increased colonization recorded in temperate regions of Australia, Europe, Japan, India, Sri Lanka, and South America (Sethunga & Manage, 2010; Piyathilaka et al., 2015; Wijewickrama & Manage, 2019). CYN is a water-soluble, stable, tricyclic guanidine combined with uracil by the hydroxyl bridge (Runnegar et al., 2002) (Figure 1).



Figure 1: Chemical structure of cylindrospermopsin

The toxicity of CYN to the humans was first recorded by the "Palm Island Mystery Disease" of 1979, in Queensland, Australia, which affected 148 children and 10 adults, causing severe hepatotoxicity with renal tubular damage (Byth, 1980; Hawkins et al., 1985; Ohtani et al., 1992). Fatalities in haemodialysis patients due to the reverse osmosis unit of the dialysis water supply has been reported (Carmichael, 2001). In vitro animal experimental studies in rodents fed with MCs contaminated environmental water samples and extracts, or purified toxins of CYN have shown toxic effects on the vascular system (Hawkins et al., 1985), lymphatic system (Hawkins et al., 1985), reproductive system (Chen et al., 2011), immune system (Lankoff et al., 2004), and other organ damage: liver (Hawkins et al., 1985; Manage et al., 2009c; Drobac et al., 2011), kidney (Hawkins et al., 1985; Piyathilaka et al., 2015; Abeysiri et al., 2018a; Abeysiri & Manage, 2018a), lungs, heart, stomach, and adrenal glands (Hawkins et al., 1985).

Thus, considering the toxic nature of CYN on human health, the World Health Organization (WHO) established a provisional guideline limit for CYN of 2 µg/L in drinking water (WHO, 2003; SLSI, 2013) and $0.02 \mu g/kg$ of body weight/day as the Tolerable Daily Intake (TDI) for humans (Guzman-Guillen et al., 2014). Recently, there have been increasing reports on the number and the intensity of blooms in reservoirs in the North Central, North East, and Uva Provinces of Sri Lanka (Madhushankha et al., 2013; Yatigammana & Perera, 2017; Manage, 2019). Further, scientific research has revealed contamination of the majority of drinking water reservoirs by toxin-producing cyanobacterium M. aeruginosa, Cylindrospermopsis sp., Anabaena sp. and a significant correlation between cyanobacterial cell density and the cyanotoxins in drinking water sources including dug wells (Hettiarachchi & Manage, 2014; Piyathilaka et al., 2016; Manage, 2019). Therefore, the present study was aimed at evaluating the effects of subchronic exposure to different doses of purified CYN and CYN contaminated well water on the liver of male Wistar rats through oral gavage over 90 days.

MATERIALS AND METHODS

Ethical approval was obtained from the Ethics Review Committee of the Faculty of Medical Sciences (An.E.17/18), University of Sri Jayewardenepura. The study was carried out following the relevant guidelines and regulations given by the Ethics Review Committee.

Animals

Eight-week-old male Wistar Rats (183.2 \pm 0.2 g) purchased from the Medical Research Institute (MRI) of Sri Lanka were acclimatized for one week at the Animal House, Faculty of Medical Sciences, University of Sri Jayewardenepura. The rats were provided with food and water *ad libitum* unless otherwise stated, and maintained on a 12 hour light/dark cycle at $28 \pm 2^{\circ}$ C. Water and food consumption were measured, and the behaviour of the animals was observed throughout the study.

Compound

CYN used in these investigations was Lot # BCBS9482v supplied by Sigma-Aldrich (USA). The purified toxin $(C_{15}H_{21}N_5O_7S;$ product no: 32087) had an estimated purity of >95%.

Experimental design

In this study, three CYN doses were prepared based on the WHO standard recommendation of CYN in drinking water: higher than the WHO standard (HW) 2.5 µg/L, equal to the WHO standard (W) 2.0 µg/L and lower than the WHO standard (LW) 1.5 μ g/L. The pure CYN doses, according to the mean initial body weights of the rats fed with the above concentrations, were HW = 0.175 μ g/kg/day, W = 0.140 μ g/kg/day, and LW = 0.105 µg/kg/day. A single well water sample collected from the Padaviya area of the North Central Province of Sri Lanka contained 2.3 µg/L CYN, which was above the WHO recommended level. This was used as the environmental sample. Accordingly, the environmental exposure group (EN) received 0.161 µg/kg/day of CYN. Distilled water was given to the control group. The CYN solutions, well water, and distilled water were fed for 90 days. The mouse groups were labeled as HW, W, LW, EN, and control according to the solutions received by them. Rats behaviour, and water and food consumption were recorded at intervals of every two days throughout the experiment. Nutritionally complete solid food (Saboudry, 1988) was supplied ad libitum to avoid contaminated urine samples with food dust. The solutions of CYN were formulated biweekly and adjusted to reflect weight changes in the treated groups before administration. After 90 days of dosing, the animals were euthanized, weighed (g), and the liver was resected out with minimum trauma.

Collection of blood

Venous blood samples were drawn from the lateral tail vein at 0, 7, 14, 28, 42, and 60 days for clinical chemistry. At the end of the experiment (90 days), when the rats were under anaesthesia prior to euthanization, blood was collected by cardiac puncture for haematological and chemical analysis. Blood for haematological analysis was transferred to a 1 mL tube containing EDTA to prevent coagulation.

Clinical chemistry

Serum concentrations of aspartate amino-transferase (AST) and aspartate alanine-transferase (ALT) were analyzed using Bialabo diagnostic kits (France) and a fully automated analyzer (Thermo Fisher Scientific, INDIKO, Finland). All animals were used to measure AST and ALT at 0, 7, 14, 28, 42, 60, and 90 days.

Haematology

Full blood count (haemoglobin (Hb), red cell indices, platelet count, total and differential white blood cell counts) and reticulocyte count were analyzed using a fully automated 3 part haematology analyzer (BCC-3000, China).

Histopathological evaluation

At necropsy, the liver was observed for gross pathological changes, and the weight was measured. The liver was serially cut, and the cut surfaces were examined for any difference in colour, areas of necrosis or fibrosis, and photographed with the reference number. Two random slices were selected and immediately fixed in ten times the volume of the slices in 10% formalin for routine histological assessment. If any abnormal areas were present, additional slices were fixed similarly for the histological evaluation. The changes observed were recorded in the datasheet. Following processing and dehydration, prepared slides were transferred to xylene for 5 min., absolute alcohol for 5 min., 90% alcohol for 2 min., 80% alcohol for 2 min., 70% alcohol for 2 min., and 60% alcohol for 2 min., and stained with hematoxylin and eosin; special stains used included periodic acid - Schiff stain (PAS) and silver special stains. Masson trichrome to assess fibrosis was not used due to the short duration of the experimental period.

For histological changes, the prepared sections were examined under the light microscope (Olympus CX31, magnification $\times 40$, $\times 100$, and $\times 400$). The scoring system consisted of evaluating the tissues and assigning a semi-quantitative score scheme, from 0 = no lesion to 3 = severe lesions, according to the changes observed in the liver architecture, hepatocytes, and the presence of inflammation. After an initial review, selected tissues were re-assessed.

Statistical evaluation

One-way ANOVA was used to statistically evaluate significant differences, by using MINITAB version 17 statistical software (MINITAB, State College, PA, USA), and differences were considered significant if p < 0.05.

RESULTS AND DISCUSSION

Clinical findings

The pure CYN doses received by the rats in different groups according to the mean initial body weights were $HW = 0.175 \ \mu g/kg$, $W = 0.140 \ \mu g/kg$, $LW = 0.105 \ \mu g/kg$ and $EN = 0.161 \ \mu g/kg$.

The external appearance and signs of toxicity were not observed in CYN-treated animals during the study. Similar findings were reported by Diez-Quijada *et al.* (2021); clinical and visual observations during the study period did not show abnormalities in the groups studied. In general, there were no changes in behaviour or locomotor activity during the study period due to repeated exposure to CYN. Gradual mean body weight increments were noted in both treated (190.4 \pm 3.8 - 300.3 \pm 2.2 g) and control $(198.4 \pm 1.6 - 317.7 \pm 3.5 \text{ g})$ groups of animals until the thirteenth week (Figure 2). However, there was a statistically significant reduction (p < 0.05) in body weights of rats in the W, HW, and EN groups given CYN when compared to the control group at weeks 12 and 13. Comparable results were reported by Chernoff et al. (2018) after the 90-day gavage of CYN in a mice experiment. Increased body weight with low toxin doses (30 and 60 μ g/kg/day) and decrease of body weight at high doses (432 and 657 µg/kg/day) of CYN in mice had been reported in a previous study (Humpage and Falconer, 2003). In another study, no statistically significant differences were observed in body weight between treated and control groups in males (Diez-Quijada et al., 2021). CYN is known to affect both hepatic and renal systems (Seawright et al., 1999; Humpage & Falconer, 2003; Bazin et al., 2012), and data presented here specify hepatic toxicity at four dose levels.



Figure 2: Mean body weight changes of Wistar rats in different treatment groups (HW = $0.175 \ \mu g/kg$, W = $0.140 \ \mu g/kg$, LW = $0.105 \ \mu g/kg$, EN = $0.161 \ \mu g/kg$) *p < 0.05

Table 1: Bodyweight (g) and relative percentages of the liver weight of male Wistar rats exposed to different doses of CYN for 90 days.
 Values are mean ± SD for 7 rats/group. Statistics ANOVA test; *p < 0.05

Parameters	Control	Pure CYN dose CYN envi		CYN environmental	
			(µg/kg)		dose (µg/kg)
	0	0.105 (LW)	0.140 (W)	0.175 (HW)	0.161(EN)
	N = 7	N = 7	N = 7	N = 7	N = 7
Body weight (g)	318 ± 0.98	$284\pm1.03~\text{*}$	$289\pm1.26\text{*}$	$300\pm2.42\texttt{*}$	$296\pm1.33\text{*}$
Absolute weight (g)	9.02 ± 0.63	8.62 ± 0.56	$10.41\pm0.70\texttt{*}$	9.27 ± 0.94	$9.27 \pm 1.30 \texttt{*}$
% Liver	3.19 ± 0.13	$2.87\pm0.03*$	$2.87\pm0.05\texttt{*}$	$2.75\pm0.02\texttt{*}$	$2.78\pm0.01\texttt{*}$

There were no gross pathological changes observed in the livers. The liver weights as a percentage of the body weight showed a significant decrease in treatments groups: 284 ± 1 g in LW (p < 0.05), 289 ± 1 g in W (p < 0.05), 300 ± 2 g in HW (p < 0.05), and 296 ± 1 g in EN (p < 0.05) in comparison to the control group (Table 1). No significant differences were found in the absolute and relative weights of males at any toxin concentration in the study by Diez-Quijada *et al.* (2021). However, Reisner *et al.*, 2004 found an increase in the relative weights of the livers in male mice exposed to CYN.

The AST level gradually increased up to the endpoint of the study and the differences were statistically significant in HW (p < 0.05), W (p < 0.05), LW (p < 0.05), and EN (p < 0.05) groups compared to the control at day 7, 14, 28, 42, 60, and 90 (Figure 3a).

Significant differences of the ALT activity was found in HW (p < 0.05), W (p < 0.05), LW (p < 0.05) and EN (p < 0.05) groups compared to the control at day 7, 14, 28, 42, 60, and 90 (Figure 3b). Increased serum levels of AST and ALT activity indicated hepatic toxicity, which was compatible with changes observed on histology. Elevated ALT levels in mice with a high dose of CYN exposure were reported by other investigators (Chernoff et al., 2011; 2014). The AST activity was also elevated after oral exposure at the intermediate and highest dose groups (37.5 and 75 μ g/kg CYN) to the control group (p < 0.001) (Diez-Quijada *et al.*, 2021). The data obtained from Dordevic et al. (2017) showed that the alterations in serum biochemical parameters did not produce significant liver damage in rats. Moreover, increased AST activity was found in females at all doses assayed (Chernoff et al., 2018).



Figure 3: (a) Mean serum aspartate amino-transferase (U/L), (b) Mean serum as partate alanine-transferase (U/L) concentrations changes of male Wistar Rats in different treatment groups (HW= 0.175 μg/kg, W= 0.140 μg/kg, LW= 0.105 μg/kg, EN=0.161 μg/kg) *p < 0.05</p>

Haematology

Haematological parameters measured at day 90 are summarized in Table 2.

Non-significant reductions in both haematocrit and haemoglobin levels were reported in the HW, W, and EN groups. In the EN group, a significant decrease in lymphocyte count (p < 0.05) and monocyte count

(p < 0.05) were observed in comparison to the control group. The absence of similar changes in the groups treated with pure CYN extracts makes it unlikely that the haematological changes can be attributed to the toxic effects of CYN. However, the rats in the EN group received the highest dose of CYN. Augmented haematocrit levels were reported as a response to exposure to low levels of CYN and morphological changes in the RBC of mice, which gradually developed into an acanthocyte-like form, cytoplasmic irregularly spaced projections, etc. (Reisner *et al.*, 2004). Similarly, elevated haematocrit levels in male and female mice after 16 weeks of exposure to variable concentrations of CYN in their drinking water (10–55 µg/kg/day during 42 weeks) were also found as the most critical effect observed (Sukenik *et al.*, 2006). By contrast, all the values remained within the normal range for the strain (Lira *et al.*, 2020). There were no significant alterations in the differential white blood cell counts of rats exposed to CYN in comparison to the untreated groups (Diez-Quijada *et al.*, 2021).

Table 2: Selected haematology parameters of Wistar Rats on the 90th day of the experimentStatistics ANOVA test; *p < 0.05

Parameters	HW (0.175 µg/kg)	W (0.140 µg/kg)	LW (0.105 µg/kg)	EN (0.161 µg/kg)	Control
Haematocrit (%)	23.59 ± 1.18	22.27 ± 0.71	24.58 ± 0.96	22.93 ± 1.20	24.00 ± 1.14
Haemoglobin (g/dL)	16.23 ± 1.02	15.68 ± 0.25	17.07 ± 0.68	16.13 ± 0.89	16.63 ± 0.97
Reticulocyte count (10 ¹² /L)	8.85 ± 0.35	8.19 ± 0.29	9.06 ± 0.23	8.55 ± 0.56	8.83 ± 0.50
Mean cell volume (fL)	106.54 ± 1.96	108.77 ± 3.09	108.47 ± 4.07	107.07 ± 1.41	108.93 ± 3.74
MCHC (g/dL)	69.11 ± 0.74	70.46 ± 1.38	69.43 ± 0.41	70.53 ± 0.61	69.27 ± 1.00
PLT (10 ⁹ /L)	689.1 ± 112.3	682.9 ± 82.6	754.3 ± 118.4	682.7 ± 97.2	622.3 ± 184.3
Lymphocyte count (10 ⁹ /L)	6.80 ± 2.58	6.06 ± 1.49	9.33 ± 2.81	$3.50 \pm 1.51 \texttt{*}$	7.03 ± 2.71
Monocyte count (10 ⁹ /L)	1.31 ± 1.65	0.74 ± 0.22	0.77 ± 0.27	$0.40\pm0.18\texttt{*}$	0.67 ± 0.24
Granulocyte count (10%/L)	1.51 ± 2.00	0.57 ± 0.21	0.83 ± 0.29	0.37 ± 0.15	0.47 ± 0.24

Table 3:Incidence and severity of histopathologic liver lesions in Wistar rats on the day 90 of exposure to cylindrospermopsin
Lesion severity was based on a 0 - 3 grading scale

Histological features	HW (0.175 µg/kg)	W (0.140 µg/kg)	LW (0.105 µg/kg)	EN (0.161 µg/kg)	Control
Hepatocytes	Ballooning (3)	Ballooning (2)	Ballooning (1)	Ballooning (3)	Normal (0)
Inflammation-lobular	Mild/spotty inflammation < 5/lpf (1)	Mild/spotty inflammation < 5/lpf (1)	Mild/spotty inflammation < 5/lpf (1)	Mild/spotty inflammation < 5/lpf (1)	Normal (0)
Hepatocyte cell death	Lobular/spotty (2)	Lobular/spotty (2)	Lobular/spotty (2)	Lobular/spotty (2)	Normal (0)
Hepatocyte regeneration	Bi-nucleation (2)	Normal (0)	Normal (0)	Bi-nucleation (2)	Normal (0)
Kupffer cells	Prominent (3)	Mild (1)	Mild (1)	Prominent (3)	Normal (0)
Nuclear pyknosis	Mild (1)	Mild (1)	Mild (1)	Mild (1)	Normal (0)
Haemorrhage	Mild (1)	Normal (0)	Normal (0)	Mild (1)	Normal (0)

Histopathology

Histopathological changes of the liver associated with CYN exposure are summarized in Table 3. These changes

include hepatocyte ballooning, focal inflammation nuclear pyknosis, hepatocyte cell death/necrosis, features of hepatocyte regeneration, Kupffer cell hyperplasia, and haemorrhages (Table 3). The hepatocytes showed ballooning degeneration (Figure 4b), Kupffer cell hyperplasia (Figure 4c), lobular haemorrhage and necrosis (Figure 4d) in EN $(0.161 \mu g/kg)$. Lobular haemorrhage and necrosis (Figure 5b), lobular inflammation (Figure 5c), lobular haemorrhage (Figure 5d), sinusoidal congestion (Figure 5e), and centrilobular haemorrhage and necrosis (Figure 5f) were prominent in HW (0.175 µg/kg) group. Lobular inflammation and necrosis were observed in W (0.140 μ g/kg) (Figure 6b). Lobular inflammation (Figure 7b) was significant in the LW (0.105 μ g/kg) group. All histological features were typical in the control group (Figure 4a, 5a, 6a, 7a). In this study, hepatic lesions were seen in all treated groups. The effects were more severe in the HW and EN groups. The well water fed to the EN group contained CYN well above the safe level recommended by the WHO. These results were similar to the findings in 50 µg/kg doses of CYN administered for 5 days which produced a widespread disruption of hepatic cords, hepatocellular degeneration, necrosis, and centrilobular inflammation, as reported by Chernoff (Chernoff et al., 2011; 2014). In CYNexposed cells in a culture, prominent features included necrosis and/or apoptosis of hepatocytes possibly

resulting from a disruption of the cytoskeleton (Fessard & Bernard, 2003). Rats exposed by gavage to pure CYN (7.5–75.0 μ g/kg b.w.) at 0, 24, and 45 h, showed irritation in the stomach and changes in hepatocytes (Diez-Quijada *et al.*, 2019).

Thus, Bernard et al. (2003) tested in vivo the toxicity of cell extracts of different C. raciborskii strains in mice and found fatal acute neurotoxicity, hepatotoxicity, and the absence of toxicity, depending on the toxins and the toxicological activity of unknown compounds detected. In the 2021 study by Leticia et al., at a dose of 75.0 µg/kg b.w, only scarce microscopic changes were observed in the livers of rats. Livers displayed a very mild hepatic sinusoidal ectasia and small numbers of hepatocytes with apoptotic features or spotty necrosis, mainly localized in the surroundings of the centrilobular vein. Occasionally, hepatocytes presented eosinophilic cytoplasmic inclusions (Leticia et al., 2021). The present study further confirmed the signs of hepatic injuries due to CYN. Liver/body weight ratios, the elevation of AST and ALT levels, and histopathological changes were important parameters of CYN-induced liver toxicity.



Figure 4: Environmental exposure (EN) Haematoxylin and eosin stain.

Figure 4a - Control ×400; Figure 4b - Ballooning degeneration of hepatocytes ×400; Figure 4c - Kupffer cell hyperplasia (arrow) ×400; Figure 4d - Lobular haemorrhage and necrosis (arrows) ×100



Figure 5: Above WHO level (HW) Haematoxylin and eosin stain
 Figure 5a - Control ×400; Figure 5b - Lobular haemorrhage and necrosis ×100; Figure 5c - Lobular inflammation (arrow) ×400; Figure 5d - Focus of lobular haemorrhage ×400; Figure 5e - Sinusoidal congestion ×400; Figure 5f - Centrilobular haemorrhage and necrosis shown by the arrow ×100



 Figure 6:
 WHO level (W) Haematoxylin and eosin stain

 Figure 6a Control ×400; Figure 6b - Lobular inflammation and necrosis (Arrow) ×400



 Figure 7:
 Below WHO level (LW). Haematoxylin and eosin stain

 Figure 7a - Control ×400; Figure 7b - LW - Lobular Inflammation (Arrow) ×400

CONCLUSION

The study results showed that the differences in liver toxicity were associated with the administered CYN dose. It provides substantial evidence that CYN contaminated water for drinking purposes may lead to mammalian liver cell injury.

Conflict of interest statement

The authors declare that they have no competing interests

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RESEARCH ARTICLE

Functional Analysis

On the construction of unitaries representing minimal inner toral polynomials

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Abstract: A polynomial $p = p(z, w) \in \mathbb{C}[z, w]$ is called an inner toral polynomial if the zero set of p, contained in $\mathbb{D}^2 \cup \mathbb{T}^2 \cup \mathbb{E}^2$ where \mathbb{D}^2 is the open bidisk, \mathbb{T}^2 is the two dimensional torus and \mathbb{E}^2 is the exterior of the closed bidisk in \mathbb{C}^2 . Such a polynomial is called a minimal inner toral polynomial if it divides all the other polynomials with same zero set as itself. The bidegree of p is defined as the ordered pair (n, m) whenever p has degrees n and m in variables z and w respectively. For a minimal inner toral polynomial p(z, w) of bidegree (n, m), there exist block unitary matrices, $\begin{pmatrix} A & B \\ C & D \end{pmatrix}_{((m+n)\times(m+n))}$, such that p(z,w) is a constant multiple of the determinant of $\begin{pmatrix} A - wI_m & zB \\ C & zD - I_n \end{pmatrix}$, where, blocks A, B, C and D are matrices with complex entries of sizes $(m \times m)$, $(m \times n)$, $(n \times m)$ and $(n \times n)$ respectively. Because the size of such unitaries solely depends on the bidegree of the considered minimal inner toral polynomial, the process of computing such unitary matrices is manageable when the bidegree is very small but become strenuous exponentially with the bidegree. In this work, we introduce a method of constructing unitary matrices representing reducible minimal inner toral polynomials in terms of the unitaries representing its factors and vice versa.

Keywords: Block matrices, distinguished varieties, inner toral polynomials, unitary matrices.

INTRODUCTION

Inner toral polynomials are polynomials in two complex variables, say in z and w, such that their zero sets lie in the open bidisk and exit the open bidisk to the exterior

of the closed bidisk through the two-dimensional torus. These polynomials became interesting, because the variety generated by them have this special form, and most importantly, a given pair of pure commuting isometries that satisfies an algebraic relationship contains an inner toral polynomial that annihilates the same pair of isometries. Therefore, studies related to such polynomials appear in many research articles in functional analysis as well as operator theory (Agler & McCarthy, 2002, 2005, 2006; Agler et al., 2006, 2012; Knese, 2010; McCarthy, 2012; Jury et al., 2012; Pal & Shalit, 2014; Wijesooriya, 2018; Bhattacharyya et al, 2022). An inner toral polynomial is called a minimal inner toral polynomial or simply minimal, if it divides all the polynomial with the same zero set as itself. In 2005, Agler and McCarthy have proved that a minimal inner toral polynomial can be represented with a unitary block matrix in a way that the determinant of a shifted version of the unitary block matrix is a constant multiple of the polynomial.

If the inner toral polynomial we consider has bidegree (n, m), then a unitary representing it, is a square matrix of size n + m. Due to the large size of these matrices, and the fact that the number of unknowns in the matrices increases significantly with the bidegree of the polynomial, it is not computationally efficient to come up with examples. In this study we paid attention to reducible minimal inner toral polynomials which are obviously factorizable into simpler irreducible minimal inner toral polynomials and we found a method to compute unitaries representing such reducible minimal inner toral polynomials in terms

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of unitaries representing their factors and vice versa. We define a special type of a direct sum for block matrices, which we call the diagonal wise direct sum, and we construct desired unitary matrices by performing the diagonal wise direct sum on them.

PRELIMINARIES

In this section we discuss necessary background materials to discuss the main result in the preceding section. Throughout this paper we consider \mathbb{D} to be the open unit disk, \mathbb{T} to be the unit circle and \mathbb{E} to be the exterior of the closed unit disk in the complex plane.

Definition 1: Let $p = p(z, w) \in \mathbb{C}[z, w]$ be a polynomial in *z* and *w*. We say that *p* has bidegree (n, m) if *p* has degree *n* in *z* and *m* in *w*.

Definition 2: A polynomial $\mathfrak{p} = \mathfrak{p}(z, w) \in \mathbb{C}[z, w]$ is called an inner toral polynomial if the zero set of \mathfrak{p} , $Z(\mathfrak{p})$, contained in $\mathbb{D}^2 \cup \mathbb{T}^2 \cup \mathbb{E}^2$.

In other words, for a two-variable polynomial $p = p(z, w) \in \mathbb{C}[z, w]$, in order to be inner toral, every zero (z_0, w_0) of p must satisfy the condition that either $|z_0|, |w_0| < 1$ or $|z_0| = |w_0| = 1$ or $|z_0|, |w_0| > 1$.

For example, $p_1(z, w) = z^n - w^m$ for $n, m \in \mathbb{N}$ is inner toral. However, $p_2(z, w) = z^n - 2w^m$ is not inner toral because $\left(2^{\frac{1}{n}}, 1\right)$ is a zero of p_2 but it is not in $\mathbb{D}^2 \cup \mathbb{T}^2 \cup \mathbb{E}^2$.

Definition 3: An inner toral polynomial $\mathfrak{p}(z,w) \in \mathbb{C}[z,w]$ is called minimal а inner toral polynomial, if it divides any $q \in \mathbb{C}[z, w]$ with Z(p) = Z(q).

A minimal inner toral polynomial is minimal in the sense that it has smallest possible bidegree, and it is square free (unique up to constant multiples) with the same zero set as the original one. Therefore, clearly the minimal polynomial is also inner toral. Given an inner toral polynomial p, its minimal version always exists as it is the product of all the irreducible factors (of algebraic multiplicity one) of p (up to constant multiples). For example, $(z - w)^2(z + w)$ is inner toral and (z - w)(z + w) is its minimal version.

Definition 4: For an inner toral polynomial \mathfrak{P} the set, $\mathfrak{V}(\mathfrak{p}) = Z(\mathfrak{p}) \cap \mathbb{D}^2$ is called a distinguished variety.

Definition 5: A square matrix U is called a unitary matrix if its conjugate transpose U^* is its inverse. That is, $UU^* = U^*U = I$ where I is the identity matrix.

Our work is based on the unitaries representing minimal inner toral polynomials. The Theorem given below explain the existence of such unitary matrices and how exactly they look like. This Theorem was first proved by Agler and McCarthy in 2005 and an alternative proof was given by Knese in 2010. This result is one of the breakthrough results in this setting as it eventually became a very helpful tool in the studies carried out by many mathematicians related to inner toral polynomials, distinguished varieties and pure algebraic iso-pairs (Knese, 2010; Jury *et al.*, 2012; McCarthy, 2012; Pal & Shalit 2014; Wijesooriya, 2018).

Theorem 1: (Knese, 2010) For a minimal inner toral polynomial $\mathfrak{p}(z, w)$ of bidegree (n, m) there exists an unitary matrix $U = \begin{pmatrix} A & B \\ C & D \end{pmatrix} \begin{pmatrix} m & \text{such that} \\ n \end{pmatrix}$

- $\det \begin{pmatrix} A wI_m & zB \\ C & zD I_n \end{pmatrix} = k p(z, w)$ where k is a constant and
- if det $(D \lambda I_n) = 0$, then $|\lambda| \neq 1$.

Here, A, B, C and D are matrices of sizes $(m \times m)$, $(n \times m)$, $(m \times n)$ and $(n \times n)$ respectively. Such unitary matrices, U, are called unitaries representing minimal inner toral polynomial p(z, w).

For example, let $U = \begin{pmatrix} a & b \\ c & d \end{pmatrix}_{2 \times 2}$ be a unitary representing the minimal inner toral polynomial z - w. That means the determinant of $\begin{pmatrix} a - w & zb \\ c & zd - 1 \end{pmatrix}$ is equal to k(z - w) for some constant $k \in \mathbb{C}$. By comparing the coefficients of the equation (ad - bc)z - dzw + w - a = k(z - w), and considering the fact that U is a unitary matrix, we have a = d = 0, and |b| = |c| = 1 with bc = 1. Therefore, unitaries representing z - w are of the form $\begin{pmatrix} 0 & a \\ \overline{\alpha} & 0 \end{pmatrix}$ where α is a unimodular constant. This computation can be done easily due to the fact that U has a relatively small size and the system of equations involved is simple. However, that is not the case all the time. For example, unitaries representing the minimal inner toral polynomial $z^4 - w^4$ are 8×8 matrices and it is not quite easy to compute such unitaries as it involved finding 64 unknowns.

Remark 1: It is evident from the above example that, given an inner toral polynomial one can find more than one unitary matrix representing the considered polynomial.

Before moving to the results and discussions, we prefer to discuss the invertibility of the block matrix zD - I to avoid any ambiguities that could appear in the proof of the main Theorem.

Theorem 2. (Bezout's Theorem) If two algebraic curves, say described by and p(z,w) = 0 and q(z,w) = 0 do not have any common components, then they have only finitely many points in common.

In fact, if p(z, w) is a minimal inner toral polynomial and if zeroes of p and q are common except for finitely many zeroes, then p must be a factor of q.

Remark 2: Let $U = \begin{pmatrix} A & B \\ C & D \end{pmatrix}$ be a unitary representing the minimal inner toral polynomial p(z, w) of bidegree (n, m). By Theorem 1, $\det\begin{pmatrix} A - wI_m & zB \\ C & zD - I_n \end{pmatrix} = k p(z, w) \text{ for some constant}$ k. Note that $\det \begin{pmatrix} A - wI_m & zB \\ C & zD - I_n \end{pmatrix} = \det(zD - I) \cdot$ $det[(A - wI) - zB(zD - I)^{-1}C]$. It is immediate that this equation is true only if the matrix zD - I is invertible. Since the matrix D is of size $n \times n$, the determinant of zD - I can be vanished only at finitely many values of z. Therefore, det $\begin{pmatrix} A - wI_m & zB \\ C & zD - I_n \end{pmatrix}$ vanishes at all except finitely many points in $Z(\mathfrak{p})$. By Theorem 2, and the minimality of \mathfrak{p} , det $\begin{pmatrix} A - wI_m & zB \\ C & zD - I_n \end{pmatrix}$ vanishes at all the points in Z(p). Eventually, this leads to the fact that $\det \begin{pmatrix} A - wI_m & zB \\ C & zD - I_n \end{pmatrix} = \det(zD - I) \cdot$ $det[(A - wI) - zB(zD - I)^{-1}C] = k p(z, w)[see page$ 16 of (Knese, 2010) for more details on this]. Therefore, the non-invertibility of zD - I at finitely many $z \in \mathbb{C}$, does not harm the determinantal representation of the unitary to the polynomial. In the proof of the main theorem presented in this paper, we say the fact that zD - I is invertible, with the fully understanding that zD - I is invertible at all except finitely many $z \in \mathbb{C}$.

Now we present the method we found on computing unitaries representing minimal inner toral polynomials.

RESULTS AND DISCUSSION

Let $m_i, n_i \in \mathbb{N}$ and suppose U_i is a block matrix of the form

$$U_i = \begin{pmatrix} m_i & n_i \\ A_i & B_i \\ C_i & D_i \end{pmatrix} \quad m_i$$

for i = 1, 2, ..., s with sizes given above.

$$U_1 \bigoplus_D U_2 \bigoplus_D \dots \bigoplus_D U_s =$$

$$\begin{pmatrix} A_1 & 0 & 0 & \cdots & 0 & B_1 & 0 & 0 & \cdots & 0 \\ 0 & A_2 & 0 & \cdots & 0 & 0 & B_2 & 0 & \cdots & 0 \\ 0 & 0 & A_3 & \cdots & 0 & 0 & 0 & B_3 & \cdots & 0 \\ \vdots & \vdots & \vdots & \ddots & \vdots & \vdots & \vdots & \vdots & \ddots & \vdots \\ 0 & 0 & 0 & \cdots & A_s & 0 & 0 & 0 & \cdots & B_s \\ C_1 & 0 & 0 & \cdots & 0 & D_1 & 0 & 0 & \cdots & 0 \\ 0 & C_2 & 0 & \cdots & 0 & 0 & D_2 & 0 & \cdots & 0 \\ 0 & 0 & C_3 & \cdots & 0 & 0 & 0 & D_3 & \cdots & 0 \\ \vdots & \vdots & \vdots & \ddots & \vdots & \vdots & \vdots & \vdots & \vdots & \ddots & \vdots \\ 0 & 0 & 0 & \cdots & C_s & 0 & 0 & 0 & \cdots & D_s \end{pmatrix}.$$

We call this sum the diagonalwise direct sum. For the ease of notation, let us denote the block matrix

$$\begin{pmatrix} A_1 & 0 & 0 & \cdots & 0 \\ 0 & A_2 & 0 & \cdots & 0 \\ 0 & 0 & A_3 & \cdots & 0 \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ 0 & 0 & 0 & \cdots & A_s \end{pmatrix}$$
by $\bigoplus A_i$ and likewise $\bigoplus B_i, \bigoplus C_i$

and $\bigoplus D_i$. Likewise, we will use the notation, \bigoplus , to denote the block diagonal matrices where it is necessary. Again, for the ease of using notations, given a block matrix $U = \begin{pmatrix} A & B \\ C & D \end{pmatrix}$, we denote $\begin{pmatrix} A - wI & Bz \\ C & zD - I \end{pmatrix}$ by U(z, w).

Let $\mathfrak{p}(z,w) \in \mathbb{C}[z,w]$ be a reducible minimal inner toral polynomial. Write $\mathfrak{p} = \mathfrak{p}_1 \mathfrak{p}_2 \mathfrak{p}_3 \dots \mathfrak{p}_s$ for some $s \in \mathbb{N}$. Note that each factor \mathfrak{p}_i is also a minimal inner toral polynomial. Let (n,m) and (n_i,m_i) be the bi-degrees of the polynomial \mathfrak{p} and \mathfrak{p}_i for $i = 1, 2, \dots, s$ respectively. Note that $n = \sum_{i=1}^{s} n_i$ and $m = \sum_{i=1}^{s} m_i$. Further, let U and U_i be square matrices of order (m + n) and $(m_i + n_i)$ for i = $1, 2, \dots, s$ respectively. Now, we present our main result:

Theorem 3

If U_i is a unitary matrix representing the minimal inner toral polynomial p_i for each, i = 1, 2, ..., s, then the

matrix $U = U_1 \bigoplus_D U_2 \bigoplus_D ... \bigoplus_D U_s$ is a unitary matrix representing the polynomial p. Conversely, suppose $U = \begin{pmatrix} \bigoplus A_i & \bigoplus B_i \\ \bigoplus C_i & \bigoplus D_i \end{pmatrix}$ is a unitary representing the minimal inner toral polynomial p. where i = 1, 2, 3, ..., s. Each matrix given by $U_i = \begin{pmatrix} A_i & B_i \\ C_i & D_i \end{pmatrix}$ is a unitary matrix representing the minimal inner toral polynomial p_i (up to re-labelling) for i = 1, 2, ..., s.

Proof: Let $U_i = \begin{pmatrix} A_i & B_i \\ C_i & D_i \end{pmatrix}$ for i = 1, 2, 3, ..., s with the sizes given as above. First, we must show that U is unitary. Since each $U_i = \begin{pmatrix} A_i & B_i \\ C_i & D_i \end{pmatrix}$ is a unitary, $U_i U_i^* = U_i^* U_i = I_{n_i+m_i}$ for i = 1, 2, ..., s. In fact,

$$\begin{pmatrix} A_i A_i^* + B_i B_i^* & A_i C_i^* + B_i D_i^* \\ C_i A_i^* + D_i B_i^* & C_i C_i^* + D_i D_i^* \end{pmatrix} = \\ \begin{pmatrix} A_i^* A_i + B_i^* B_i & C_i^* A_i + D_i^* B_i \\ A_i^* C_i + B_i^* D_i & C_i^* C_i + D_i^* D_i \end{pmatrix} = I_{n_i + m_i}$$

for i = 1, 2, ..., s. Clearly $U = \begin{pmatrix} \bigoplus A_i & \bigoplus B_i \\ \bigoplus C_i & \bigoplus D_i \end{pmatrix} m$, $UU^* = \begin{pmatrix} \bigoplus (A_iA_i^* + B_iB_i^*) & \bigoplus (A_iC_i^* + B_iD_i^*) \\ \bigoplus (C_iA_i^* + D_iB_i^*) & \bigoplus (C_iC_i^* + D_iD_i^*) \end{pmatrix}$ and $U^*U = \begin{pmatrix} \bigoplus (A_i^*A_i + B_i^*B_i) & \bigoplus (C_i^*A_i + D_i^*B_i) \\ \bigoplus (A_i^*C_i + B_i^*D_i) & \bigoplus (C_i^*C_i + D_i^*D_i) \end{pmatrix}$.

It easily follows that $UU^* = U^*U = I_{n+m}$. Therefore U is also a unitary matrix. It is left to prove that the unitary matrix U represents the polynomial \mathfrak{p} . Since U_i represent \mathfrak{p}_i , there exist $c_i \in \mathbb{C}$ such that $\det U_i(z,w) = c_i \mathfrak{p}_i(z,w)$ for all i = 1,2,3,...,s. In fact, for i =1,2,3,...,s, $\det \begin{pmatrix} A_i - wI_{m_i} & zB_i \\ C_i & zD_i - I_{n_i} \end{pmatrix} = c_i \mathfrak{p}_i(z,w)$, and hence

$$\det(zD_i - I_{n_i}) \cdot \det\left[\left(A_i - wI_{m_i}\right) - zB_i(zD_i - I_{n_i})^{-1}C_i\right]$$
$$= c_i \mathfrak{p}_i(z, w).$$

Note that

$$U(z,w) = \begin{pmatrix} \bigoplus (A_i - wI_{m_i}) & \bigoplus zB_i \\ \bigoplus C_i & \bigoplus (zD_i - I_{n_i}) \end{pmatrix}.$$

Since $(zD_i - I_{n_i})$ is invertible, $(\bigoplus(zD_i - I_{n_i}))$ is also invertible and for i = 1, 2, ..., s. Therefore,

$$\det [U(z,w)] =$$

$$\det (\oplus (zD_i - I_{n_i})) \cdot \det [\oplus (A_i - wI_{m_i}) - \oplus zB_i (\oplus (zD_i - I_{n_i}))^{-1} \oplus C_i]$$

$$= \det (\oplus (zD_i - I_{n_i})) \cdot \det [\oplus ((A_i - wI_{m_i}) - zB_i (zD_i - I_{n_i})^{-1}C_i)]$$

$$= \prod_{i=1}^{s} \det (zD_i - I_{n_i}) \cdot \det [(A_i - wI_{m_i}) - zB_i (zD_i - I_{n_i})^{-1}C_i]$$

$$= \prod_{i=1}^{s} c_i \mathfrak{p}_i(z,w)$$

$$= c \mathfrak{p}(z,w) \text{ where } c = c_1 c_2 c_3 \dots c_s.$$

Therefore, *U* represents \mathfrak{p} . Conversely, since *U* represent \mathfrak{p} , det $(U(z, w)) = c \mathfrak{p}(z, w)$ for some constant $c \in \mathbb{C}$. In other words,

$$\det(\oplus(zD_i-I_{n_i}))$$

$$\det\left[\bigoplus(A_i - wI_{m_i}) - \bigoplus zB_i(\bigoplus(zD_i - I_{n_i}))^{-1} \oplus C_i\right] = c p(z, w).$$

Since $(\bigoplus(zD_i - I_{n_i}))$ is invertible, $(zD_i - I_{n_i})$ is also invertible for i = 1,2,3, ..., s and

$$\left(\oplus (zD_i - I_{n_i})\right)^{-1} = \oplus (zD_i - I_{n_i})^{-1}.$$

Therefore,

$$\det\left(\bigoplus(zD_i - I_{n_i})\right) \cdot \det\left[\bigoplus\left(\left(A_i - wI_{m_i}\right) - zB_i(zD_i - I_{n_i})^{-1}C_i\right)\right]$$

= $c p(z, w).$

Consequently, $\prod_{i=1}^{s} \det(zD_i - I_{n_i})$.

$$\det\left[\left(A_{i}-wI_{m_{i}}\right)-zB_{i}\left(zD_{i}-I_{n_{i}}\right)^{-1}C_{i}\right]=c\,\mathfrak{p}_{1}\,\mathfrak{p}_{2}\ldots\mathfrak{p}_{s}.$$

That is, $\prod_{i=1}^{s} \det(U_i(z, w)) = c p(z, w) = c p_1 p_2 \dots p_s$. Note that each p_i is irreducible and distinct. Moreover, bidegree of $\det(U_i(z, w))$ is at most (n_i, m_i) and bidegree of $p_i(z, w)$ is (n_i, m_i) . Therefore, $\det(U_i(z, w))$ divides $p_i(z, w)$ up to relabeling of p_i 's. Therefore, unitary matrix U_i represents p_i , upto relabeling of p_i 's for $i = 1, 2, \dots, s$. Since p_i is a minimal inner toral polynomial, $\det(U_i(z, w)) = c_i p_i(z, w)$ up to relabeling of p_i 's. Therefore each U_i is a unitary representing p_i , up to relabeling.

For examples, consider the reducible minimal inner toral polynomial $p = z^4 - w^4 = (z - w)(z + w)$ (z - iw)(z + iw). Unitaries representing p are of the size 8 × 8 and obviously, it is difficult to evaluate all the 64 unknowns in such a matrix. However, we can easily compute the unitaries representing each factor of *p*. Unitaries representing z - w, z + w, z - iw and z + iw are of the form $\begin{pmatrix} 0 & \alpha \\ \overline{\alpha} & 0 \end{pmatrix}$, $\begin{pmatrix} 0 & \alpha \\ -\overline{\alpha} & 0 \end{pmatrix}$, $\begin{pmatrix} 0 & \alpha \\ i\overline{\alpha} & 0 \end{pmatrix}$ and $\begin{pmatrix} 0 & \alpha \\ -i\overline{\alpha} & 0 \end{pmatrix}$ respectively where $\alpha \in \mathbb{C}$ with $|\alpha| = 1$. Now using the diagonal wise direct sum, and the above Theorem, we can easily obtain a collection of unitary matrices representing *p*, which are of the form

/ 0	0	0	0	α	0	0	0\
0	0	0	0	0	α	0	0
0	0	0	0	0	0	α	0
0	0	0	0	0	0	0	α
$\overline{\alpha}$	0	0	0	0	0	0	0
0	$-\overline{\alpha}$	0	0	0	0	0	0
0	0	iα	0	0	0	0	0
/ 0	0	0	$-i\overline{\alpha}$	0	0	0	0/

Likewise, for the minimal inner toral polynomial $q(z,w) = (z^2 + zw) + i(zw + w^2) = (z + w)(z + iw)$, a collection of unitary matrices representing q can be obtained as:

 $\begin{pmatrix} 0 & 0 & \alpha & 0 \\ 0 & 0 & 0 & \alpha \\ -\bar{\alpha} & 0 & 0 & 0 \\ 0 & -i\bar{\alpha} & 0 & 0 \end{pmatrix}.$

In another example, a collection of unitaries representing

the polynomial $z^2 - w$ is given as $\begin{pmatrix} 0 & 0 & \lambda \\ \beta & 0 & 0 \\ 0 & \overline{\lambda\beta} & 0 \end{pmatrix}$ where

 $\lambda, \beta \in \mathbb{C}$ with $|\lambda| = |\beta| = 1$. Using the diagonalwise direct sum, we can construct a collection of unitary matrices representing the minimal inner toral polynomial $r(z, w) = z^3 - z^2w - zw + w^2 = (z^2 - w)(z - w)$ as

/0	0	0	λ	0\
0	0	0	0	α
β	0	0	0	0
0	0	$\overline{\lambda\beta}$	0	0
\0	$\overline{\alpha}$	0	0	0/

for $\alpha, \lambda, \beta \in \mathbb{C}$ with $|\alpha| = |\lambda| = |\beta| = 1$.

CONCLUSIONS

In this work, we have been able to introduce a method to construct unitaries representing reducible minimal inner toral polynomials in terms of unitaries representing its factors. This method of construction unitaries is useful as it reduces and simplifies the number of computations involve in computing unitaries representing reducible minimal inner toral polynomials of higher bidegree.

Conflict of interest

Authors hereby declare that there is no conflict of interest to disclose.

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RESEARCH ARTICLE

Applied Statistics

A coupled system of stochastic differential equations for probabilistic wind speed modelling

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Abstract: Wind-based electricity generation plays a vital role in the renewable energy industry. Benefits from its use can only be obtained if it is possible to accommodate its variability and limited predictability. Stochastic differential equation (SDE) based approaches have demonstrated an improved capability of predicting temporal wind speed patterns that have statistical properties that are similar to those observed in reality. However, no standard approach for deriving such models exist due to the wide variations in the temporal statistical properties that one observes in wind speed data measured from location to location. Wind speed data which have been recorded at coastal locations, exhibit non-stationary features, like diurnal effect, seasonal effects, and temporal trends. In this work, such effects are eliminated using standard smoothing techniques. A coupled system of second-order ordinary linear differential equations driven by a white noise forcing term was used for the probabilistic modelling of the residual data. The model was then used to predict the wind speed distribution and the corresponding autocorrelation function of wind speed data, recorded at the wind measurement centre of Kokkilai in northern Sri Lanka, from February 2015 to February 2016. Finally, the results of this novel approach were compared against the probabilistic modelling methods existing in the literature.

Key words: Autocorrelation function, probabilistic modelling, smoothing techniques, stochastic differential equation, white noise.

INTRODUCTION

Environmental concerns and supply uncertainties are forcing many countries to rethink their energy mix and develop diverse sources of clean, renewable energy. Cost-effective energy that can be produced without major negative environmental impact has become the goal worldwide. Wind energy, as a clean and renewable resource, has been under wide consideration around the world in the last decade. The benefits of wind energy are accompanied by several challenges: high variability, limited predictability, and non-storability. Thus wind can be identified as a highly volatile energy source and hence an efficient mechanism for modelling wind speed has become a mandatory requirement in power system regulations, control, and management of wind-based electricity generation (Zhu & Genton, 2012). How to reduce the uncertainty in predicting wind speeds has been the focus of studies over the last two decades. To smoothly integrate large-scale wind power into power systems, wind generation forecasting models have been developed to improve the accuracy of forecasts. Time series analysis-based methods, probability distributionbased approaches and applications of Kalman filters are found as the major statistical and mathematical modelling approaches used in modelling wind speeds. In Zhu & Genton (2012), it is pointed out that the accurate probabilistic modelling of wind speeds is more useful than point-forecasting for the power system operations of wind-based energy industry. Thus, in addition to the traditional mathematical and statistical tools for wind speed modelling, Stochastic Differential Equations (SDE) based models have become popular and have shown much promise (Zhu & Genton, 2012). Most of these SDE-based approaches have employed a single

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equation to produce forecasting trajectories of wind speeds (Bibby et al., 2005; Calif, 2012; Zarate-Minano et al., 2016). In addition, applications of those methods have been confined to modelling wind speed data that exhibit significant stationary features. Yet, from a practical point of view, wind speed data exhibit considerable nonstationary features. Therefore, applications of SDEs that generate stationary Ornstein-Uhlenbeck (O-U) processes are not sufficient to obtain reliable probabilistic forecasting for the wind energy industry. In this regard, the sums of diffusion method (Bibby et al., 2005) has demonstrated successful results for weakly stationary real process that have several time scale effects. However, when the influence of cyclic effects becomes intensive, the method becomes ineffective due to the increased nonstationariness. In addition, there are other sophisticated approaches based on non-linear filtering and particle filtering that can be successfully adapted to probabilistic modelling of wind speed data (Fernando & Hausenblas, 2018; Hausenblas et al., 2021).

In this paper, we propose a novel approach that first eliminates the non-stationary effects using a smoothing method and then uses two coupled oscillators driven by white noise to model the residual process. The approach was then used to successfully model empirical wind speed data, recorded at Kokkilai in the North Eastern coast of Sri Lanka from February 2015 to February 2016 under the Quantum Leap Wind Power Assessment Project of the Sustainable Energy Authority of Sri Lanka.

MATERIALS AND METHODS

In this section, we present a novel method for predicting wind speeds and validate it against the wind speed data measured at Kokkilai, a coastal point located on the North Eastern shore of Sri Lanka. The wind speed data that were originally recorded at 10 min intervals were converted to hourly averaged wind speeds. The empirical autocorrelation function of hourly wind speed data in Figure 1(a) exhibits a complex cyclic behaviour due to the influence of diurnal and seasonal effects (Brett & Tuller, 1991). As a coastal point, Kokkilai also comes under the influence of both North Eastern and South Western monsoonal wind patterns. In addition, due to the inter-monsoonal wind variations, short-scale temporal trends of wind speed variation were crucial factors that contributed to the non-stationariness. Figure 1(b) shows that the empirical wind speed distribution deviates significantly from a normal Gaussian distribution.

In order to eliminate the non-stationary features of wind speed data, the corresponding time plot and correlogram were carefully inspected to identify various cyclic patterns with distinct periods of temporal trends. Accordingly, the dataset was divided into distinct blocks based on the visual inspection of the time plot and correlogram. Regression models were then incorporated to eliminate the non-stationary features that were specific to each block.



Figure 1: (a): Empirical autocorrelation of hourly wind speeds and (b): empirical wind speed distribution approximated by a normal Gaussian distribution

When a cyclic variation of fixed period is apparent in a wind speed data set, it is eliminated with a regression model of the form,

$$u_r = b_0 + b_1 \cos\left(\frac{2\pi}{T}t\right) + b_2 \sin\left(\frac{2\pi}{T}t\right), \qquad \dots (1)$$

where $b_i(i = 0, 1, 2)$ are constants and T is the period of the cyclic variation (Chatfield, 2004; Shumway & Stoffer 2016). Here, b_0 captures the mean value of the data.

In the event where several cyclic effects are present a regression model of the following form can be used (Chatfield, 2004; Shumway & Stoffer, 2016).

$$u_r = b_0 + \sum_{i=1}^n \left(a_i \cos\left(\frac{2\pi}{T_i}t\right) + b_i \sin\left(\frac{2\pi}{T_i}t\right) \right). \quad \dots (2)$$

Here, $a_i, b_i (i = 0, 1, 2...)$ are constants and $T_i (i = 0, 1, 2...)$ is the period of the i^{th} cycle.

Furthermore, a polynomial regression model of the form shown below can be used to capture the temporal trend patterns included in a wind speed dataset (Chatfield, 2004; Shumway & Stoffer, 2016)

$$u_r = \sum_{i=0}^m c_i t^i, \qquad \dots (3)$$

where $c_i (i = 0, 1, 2...)$ are constants.

In the case, when both cyclic effects and temporal trends are present a sum of the regression models given by equation (2) and (3) can be used,

$$u_r = b_0 + \sum_{i=1}^n \left(a_i \cos\left(\frac{2\pi}{T_i}t\right) + b_i \sin\left(\frac{2\pi}{T_i}t\right) \right) + \sum_{i=1}^m c_i t^i.$$
...(4)

By visual inspection of the time plot of wind speed data and the corresponding correlogram, the dataset was partitioned into eleven blocks based on the observed different non-stationary features. The non-stationary features and the regression models incorporated are summarized in the following table (Table 1).

Then, the residual process (v) was obtained by eliminating the respective non-stationary component (u_r) from the empirical data (u) as in (5) and the corresponding time plots are given in Figure 2.

$$v = u - u_r \qquad \dots (5)$$

 Table 1:
 Periods of cyclic effects and the order of regression polynomial functions corresponding to each wind speed data block

Block numbers	Parameter values of regression models	Regression model
1 and 3	T = 24 hours	$b_0 + b_1 \cos\left(\frac{2\pi}{T}t\right) + b_2 \sin\left(\frac{2\pi}{T}t\right)$
2, 4, 5, 6, 7 and 9	$T_1 = 24$ hours,m = 2	$b_0 + \sum_{i=1}^n \left(a_i \cos\left(\frac{2\pi}{T_i}t\right) + b_i \sin\left(\frac{2\pi}{T_i}t\right) \right) + \sum_{i=1}^m c_i t^i$
8	$T_1 = 24$ hours, $T_2 = 350$ hours	$b_0 + \sum_{i=1}^n \left(a_i \cos\left(\frac{2\pi}{T_i}t\right) + b_i \sin\left(\frac{2\pi}{T_i}t\right) \right)$
10 and 11	$T_1 = 24$ hours, $T_2 = 144$ hours, $m = 2$	$b_0 + \sum_{i=1}^n \left(a_i \cos\left(\frac{2\pi}{T_i}t\right) + b_i \sin\left(\frac{2\pi}{T_i}t\right) \right) + \sum_{i=1}^m c_i t^i$



Figure 2: (a): Non-stationary component (u_r) of empirical wind speed data, (b): Time plot of empirical wind speeds and residua l process (v)



Figure 3: (a): Distributions of empirical wind speeds and residual process (v), (b): Correlograms of empirical wind speeds and residual process (v)

The correlogram of Figure 3(b) indicates that the residual process can be approximately treated as a wide sense stationary process (Chatfield, 2004).

In the rest of this section, we show how this approximately stationary residual process can be modeled by a process that is generated by a coupled pair of second-order linear SDEs.

Consider the general *n*-dimensional SDE,

$$\dot{\boldsymbol{X}} = \boldsymbol{A}(\boldsymbol{X}, t) + \boldsymbol{B}(\boldsymbol{X}, t)\boldsymbol{\eta}(t) \qquad \dots (6a)$$

$$y = CX \qquad \dots (6b)$$

where X is a $n \times 1$ matrix, A is a $n \times n$ matrix, B is a $n \times m$ matrix, and $\eta(.)$ is a $m \times 1$ white noise process.

Definition: White noise

A white noise process $\{\eta(t) | t \in [0, T]\}$ is a random process with the following properties (Gardiner, 1985; Särkkä & Solin, 2019):

1. For $t_1 \neq t_2$, $\eta(t_1)$ and $\eta(t_2)$ are independent.

2. $t \rightarrow \eta(t)$ is a Gaussian process with zero mean and

Dirac delta correlation: Mean: $m(t) = E[\eta(t)] = 0$ Correlation: $C_{\eta}(t,s) = E[\eta(t)\eta^{T}(s)] = \delta(t-s)Q$

where Q is the spectral density of the process.

From the above properties we can also deduce the following peculiar properties of white noise:

- a. The sample path $t \rightarrow \eta(t)$ is discontinuous almost everywhere.
- b. White noise is unbounded and it takes arbitrarily large positive and negative values at any finite interval.

White noise can be considered a weak derivative of the standard Wiener process (Särkkä & Solin, 2019). Therefore, the system (6a) - (6b) can be expressed as the following state-space model.

$$dX = A(X,t)dt + B(X,t)dw \qquad \dots (7a)$$

$$y = CX ...(7b)$$

Definition: Wiener process

The standard Wiener process $\{w(t) | t \in [0, T]\}$ is a continuous stochastic process that satisfies the following conditions (Gardiner, 1985; Särkkä & Solin, 2019).

- 1. Any increment $\Delta w_k = w(t_{k+1}) \cdot w(t_k)$ is a zero mean multivariate Gaussian random variable with covariance $T \Delta t_k$ where T is the diffusion matrix of the Wiener process and $\Delta t_k = t_{k+1} \cdot t_k$.
- 2. The increments are independent where the time spans of increments are not overlapped.
- 3. w(0) = 0 with probability one.

The solution of state-space model (7a) - (7b) is a random process (Gardiner, 1985; Allen, 2007; Särkkä & Solin, 2019). Since with a different realization of the noise process we get a different solution, the particular solutions of the equations are not often of interest, but instead, the aim is to determine the statistics of the solutions over all realizations. In the context of SDEs, the term A(X,t) of (7a) is called the drift function which determines the nominal dynamics of the system, and B(X,t) is the dispersion matrix which determines how the noise enters the system.

In our work, we consider a coupled system of differential equations which has a white noise driven forcing term of the following form (Särkkä & Solin, 2019).

$$\ddot{x}(t) + 2\xi_1 \omega_1 \dot{x}(t) + \omega_1^2 x(t) = \eta(t) \qquad ...(8)$$

$$\ddot{y}(t) + 2\xi_2 \omega_2 \dot{y}(t) + \omega_2^2 y(t) = \sigma \dot{x}(t)$$
 ...(9)

Here, σ is a constant and standard white noise is denoted by and represents the variation of wind speed. Then, following the relation between the white noise and the Wiener process in the context of SDEs, the above system of (8) - (9) can be expressed in the state space form of (7a) - (7b),

where

$$X = \begin{pmatrix} x \\ \dot{x} \\ y \\ \dot{y} \end{pmatrix}; A = \begin{pmatrix} 0 & 1 & 0 & 0 \\ -\omega_1^2 & -2\xi_1\omega_1 & 0 & 0 \\ 0 & 0 & 0 & 1 \\ 0 & \sigma & -\omega_2^2 & -2\xi_2\omega_2 \end{pmatrix};$$
$$B = \begin{pmatrix} 0 \\ 1 \\ 0 \\ 0 \end{pmatrix}; C = (0 & 0 & 0 & 1)$$

The transfer function of the above state space model is obtained by applying the Laplace transform with zero initial conditions.

$$H(s) = \frac{\sigma s^2}{(s^2 + 2\xi_1 \omega_1 s + \omega_1^2)(s^2 + 2\xi_2 \omega_2 s + \omega_2^2)} \qquad \dots (10)$$

By substituting $s = i\omega$ in the equation (10), the power spectral density function of the system is given by

$$S(\omega) = H(i\omega)H(-i\omega) = \frac{\sigma^2 \omega^4}{((\omega_1^2 - \omega^2)^2 + 4\xi_1^2 \omega_1^2 \omega^2)((\omega_2^2 - \omega^2)^2 + 4\xi_2^2 \omega_2^2 \omega^2)}$$
...(11)

According to the Wiener-Khinchin theorem stated as theorem 1 in Appendix *B*, the autocorrelation function is the Fourier inversion of the spectral density function. The Fourier inversion was obtained by substituting $p = i\omega$ (Lee,1970).

$$S(p) = \frac{\sigma^2 p^4}{\left[\left(\omega_1^2 + p^2 \right)^2 - 4 \xi_1^2 \omega_1^2 p^2 \right] \left[\left(\omega_2^2 + p^2 \right)^2 - 4 \xi_2^2 \omega_2^2 p^2 \right]}$$

By decomposing S(p) into partial fractions and taking the Fourier inversion, we have

$$\mathcal{F}^{-1}\left\{\frac{p}{p^2 + 2\xi_I \omega_I p + \omega_I^2}\right\} = u(t)e^{-\xi_I \omega_I t} \left(\cos\left(\omega_I \sqrt{I - \xi_I^2}\right) - \frac{\xi_I}{\sqrt{I - \xi_I^2}}\sin\left(\omega_I \sqrt{I - \xi_I^2}\right)\right)$$

$$\frac{\xi_I}{\sqrt{I - \xi_I^2}}\sin\left(\omega_I \sqrt{I - \xi_I^2}\right)$$
...(12)

$$\mathcal{F}^{-l}\left\{\frac{l}{p^2 - 2\xi_1 \omega_1 p + \omega_1^2}\right\} = -\frac{1}{\omega_1 \sqrt{1 - \xi_1^2}} u(t) e^{-\xi_1 \omega_1 t} \sin\left(\omega_1 \sqrt{1 - \xi_1^2}\right)$$
...(13)

$$\mathcal{F}^{-l}\left\{\frac{p}{p^2 - 2\xi_2 \omega_2 p + \omega_2^2}\right\} = -u(-t)e^{-\xi_2 \omega_2 t} \left(\cos\left(\omega_2 \sqrt{1 - \xi_2^2}\right) - \frac{\xi_2}{\sqrt{1 - \xi_2^2}}\sin\left(\omega_2 \sqrt{1 - \xi_2^2}\right)\right)$$
...(14)

$$\mathcal{F}^{-1}\left\{\frac{1}{p^2 + 2\xi_2\omega_2 p + \omega_2^2}\right\} = \frac{1}{\omega_2\sqrt{1 - \xi_2^2}}u(t)e^{-\xi_2\omega_2 t}\sin\left(\omega_2\sqrt{1 - \xi_2^2}\right)$$
...(15)

where u(t) is the Heaviside step function. Hence, the autocorrelation function of y, given by equation (7b), is found to be,

$$\kappa(t) = \frac{\sigma^2}{2\pi} (A_1 e^{-\xi_1 \omega_1 t} \cos\left(t\omega_{d_1} + \phi_1\right) + A_2 e^{-\xi_2 \omega_2 t} \cos\left(t\omega_{d_2} + \phi_2\right)) \qquad \dots (16)$$

Each term A_1 , A_2 , ϕ_1 and ϕ_2 is a function of the system parameters ξ_1 , ω_1 , ξ_2 and ω_2 . In order to estimate the system parameters, the normalized autocorrelation given below was used.

$$g(t) = \frac{(A_1 e^{-\xi_1 \omega_1 t} \cos(t\omega_{d_1} + \phi_1) + A_2 e^{-\xi_2 \omega_2 t} \cos(t\omega_{d_2} + \phi_2))}{(A_1 \cos\phi_1 + A_2 \cos\phi_2)}$$

...(17)

The system parameters were estimated by comparing the normalized autocorrelation function g(t) with the empirical autocorrelation function of the residual process v.

RESULTS AND DISCUSSION

To illustrate the validity of the developed model, the wind speed dataset measured at Kokkilai in the North Eastern coast of Sri Lanka was used. As described in section 2, the wind speed dataset was first filtered using (4) and (5) and then the residual was modelled using the coupled second-order SDEs (8) and (9). The theoretically constructed normalized autocorrelation function (17) that corresponds to (8) and (9) and the output was fitted to the normalized autocorrelation function of the residual data in order to estimate the model parameters. The empirical correlogram of Figure 3(b) exhibits rapid decay in the first 100 hours and this decides the time scale of the underlying process (Brett & Tuller, 1991; Zarate-Minano, Mele & Milano, 2016). Thus, the parameters of the normalized theoretical correlogram of equation (17) were estimated using the empirical autocorrelation values of first 100 hours. The least squares errors method was used to fit the parameters. Moreover, the variance of a stationary process is equal to the initial value of the corresponding autocorrelation function (Chatfield, 2004). Therefore, the value of the parameter σ , can be fitted using the following relationship.

$$Var(v) \approx \kappa(0)$$
$$Var(v) \approx \frac{\sigma^2}{2\pi} (A_1 \cos \phi_1 + A_2 \cos \phi_2) \qquad \dots (18)$$

where Var(v) is the variance of the residual process.

Using the estimated parameter values, the linear system of SDEs of equation (8) and (9) was simulated using the Euler-Maruyama numerical scheme. Then, the model-generated output was compared with the residual process using 30 sample realizations. Figure 5 shows that the probability density of the model matches that of the residual process v very closely.



Figure 4: Approximation of empirical correlogram of residual data: $\xi_1=0.9812$, $\omega_1=0.3321$, $\xi_2=0.9907$, $\omega_2=0.02845$ and $\sigma=0.4818$.



Figure 5: Comparison of probability distributions of SDE model generated stationary process and residual process



Figure 6: Comparison of final output of SDE model generated probability distribution of hourly wind speeds and that of empirical data

Next, the identified non-stationary component given by (4) was added to the model-generated outcome of the residual process by applying the inverse data transform corresponding to (5). Then the resulting wind speed distribution was compared with the empirical distribution of hourly wind speed distribution in Figure 6. It can be seen that our novel approach is capable of generating a wind speed distribution that is very close to those of the distribution of empirical data.

In addition to the comparison of the probability distributions, the comparison of the autocorrelation functions can be used to verify the level of accuracy. The comparison of autocorrelation functions is given in Figure 7 and shows excellent agreement between the model and empirical data.

Figure 8 shows the comparison of the probability distributions based on the proposed method with a method based on a single one-dimensional SDE. Probabilistic models based on a single one-dimensional SDE have been considered in many situations (Bibby *et al.*, 2005; Calif, 2012; Zarate-Minano *et al.*, 2016). Figure 8 demonstrates that the model proposed in this paper outperforms the models that use a single one-dimensional SDE.



Figure 7: Comparison of correlograms



Figure 8: Comparison of distributions of single SDE model and empirical distribution and the proposed model

CONCLUSION

SDE-based approaches for modelling statistical properties of wind speed data are comparatively new. In existing work reported in the literature, probabilistic models are developed based on a single SDE of the standard form. In contrast, in this work, we have developed a novel approach that uses two coupled linear oscillators to model the statistical properties of appropriately filtered hourly wind speed data. The coupled system of SDEs generates a strictly stationary Gaussian process. Since wind speed data usually exhibits complex non-stationary features, we propose the use of a filter to remove these effects from the original wind speed data. The correlogram of the residual process is then fitted to the normalized theoretical autocorrelation function of the coupled SDE system.

The coupled SDE model consists of five unknown parameters. Four of them were estimated by curve fitting the theoretical normalized autocorrelation function to the empirical autocorrelation values, using the least squares method, while the other parameter was computed using these estimated parameter values.

The developed model was used to predict the statistical properties of wind speed data measured at Kokkilai, a coastal point located in the North Eastern shore of Sri Lanka. The simulation results show that the statistical properties of the simulated wind sequence are very close to those of the measured wind sequence. In particular, the model-generated correlogram of our method successfully captures almost all the features of the empirical correlogram. This is a significant improvement in comparison to those predicted by existing methods.

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Appendix A

Linear time invariant stochastic differential equations

A time invariant SDE can be expressed in the standard form as follows,

dx = Ax(t)dt + Bdw(t) ...(A.1) Then the formal solution of the above equation (A.1) is written as,

$$x(t) = e^{At}x(0) + \int_{0}^{t} e^{A(t-\tau)}B(\tau)dW(\tau) \qquad ...(A.2)$$

where e^{At} is the matrix exponential function. Since the noise process is Gaussian forcing, a linear differential equation can be considered as a linear operator acting on the noise process (and the initial conditions) and the solution is also Gaussian (Gardiner, 1985; Allen, 2007; Särkkä & Solin, 2019).

• Expectation and correlation

Since the white noise process has zero mean, taking expectations from both sides of equation (A.2) gives,

$$m(t) = E(x(t)) = exp(At)E(x(t_0))$$
 ...(A.3)

which is the expected value of the SDE solutions over all realizations of noise. The corresponding covariance function is calculated by using the delta-correlation property of white noise as follows,

$$P(t) = E[(x(t)-m(t))(x(t)-m(t))^{t}]$$

$$P(t) = \exp(At)P_{0}\exp(A^{T}t) + \int_{t_{0}}^{t} \exp(A(t-\tau))BQB^{T}w(\tau)\exp(A^{T}(t-\tau))d\tau \dots(A.4)$$

where $P_0 = E[x(0)x(0)^T]$. By differentiating the above expressions for mean and covariance, the governing differential equations for the mean and covariance are given by

$$\frac{dm(t)}{dt} = Am(t) \qquad \dots (A.5)$$

$$\frac{dP(t)}{dt} = AP(t) + P(t)A^T + BQB^T \qquad \dots (A.6)$$

Moreover, it can be seen that the solutions of equation (A.1) generate a multivariate Gaussian distribution (Gardiner, 1985; Särkkä & Solin, 2019).

It should be noted when P(t) is a constant, (A.6) is reduced to a Lyapunov equation. Then, the steady covariance function can be obtained by following the context of a matrix Lyapunov equation (Mao, 2007; Särkkä & Solin, 2019).

Appendix **B**

• Fourier analysis of linear time invariant stochastic differential equations

One way to study linear time invariant SDEs is the Fourier domain (Lindgren *et al.*, 2002; Särkkä & Solin, 2019). In this case, the power spectral density which is the squared absolute value of the Fourier transform of the process is defined as follows,

$$\mathbf{S}_{\mathbf{X}}(\omega) = \mathbf{X}(i\omega)\mathbf{X}^{T}(-i\omega) \qquad \dots (B.1)$$

The power spectral density Q of a multi-dimensional white noise process is also defined accordingly.

$$\boldsymbol{Q} = \boldsymbol{W}(i\omega)\boldsymbol{W}^{T}(-i\omega)$$

The spectral density is an efficient tool for estimating the covariance function generated as a result of SDE. Suppose x(t) is a stationary multi-dimensional stochastic process with zero mean. Then the corresponding covariance function is defined as,

$$C_{\mathbf{x}}(\tau) = E[\mathbf{x}(t)\mathbf{x}^{T}(t+\tau)]$$

The covariance function of a wide sense stationary process is only dependent on the time lag (τ). The Wiener-Khinchin theorem (Lindgren *et al.*, 2002; Särkkä & Solin, 2019) states the Fourier inversion of the power spectral density function of a given wide sense stationary process provides the corresponding covariance function as follows,

Theorem 1 (Wiener-Khinchin theorem):

The covariance of a wide sense stationary random process has a spectral decomposition given by the power spectrum of that process, i.e., $C_x(\tau) = \mathcal{F}^{-1} \{ S_X(\omega) \}$.

Therefore, we have the autocorrelation function for a purely white noise process as follows,
• Covariance of the solution of linear time invariant stochastic differential equations

The spectral density function of resultant process $\mathbf{x}(t)$ of equation (A.1) can be given in the matrix version with the assumption of zero initial conditions. Note that the stationary stage can only exist if the matrix corresponds to a stable system, which means that all its eigenvalues have negative real parts. The Fourier transform of $\mathbf{x}(t)$ is,

$$X(i\omega) = ((i\omega)I - A)^{T} BW(i\omega)$$

where $W(i\omega)$ is the formal Fourier transform of the white noise w(t). It should be noted that the above Fourier transform does not strictly exist, as white noise is not square integrable. Then the spectral density of x(t) is given by the matrix,

$$S_X(\omega) = (A - (i\omega)I)^{-1}BQB^T (A + (i\omega)I)^{-T} \qquad \dots (B.2)$$

Hence the covariance function is,

$$\boldsymbol{C}_{\boldsymbol{x}}(\tau) = \mathcal{F}^{-1} \{ (\boldsymbol{A} - (i\omega)\boldsymbol{I})^{-1}\boldsymbol{B}\boldsymbol{Q}\boldsymbol{B}^{T} (\boldsymbol{A} + (i\omega)\boldsymbol{I})^{-T} \} \quad \dots (B3)$$

This provides useful means to estimate the autocovariance function of a solution to a SDE without explicitly solving the equation.

RESEARCH ARTICLE

Manufacturing and Industrial Engineering Failure prediction of solid resilient tyres due to kerb impact: A finite element modelling approach

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Abstract: Solid resilient tyres frequently impact on kerbs or obstacles when they are operated in the construction and transportation sectors. These sudden impacts can generate high stresses and thereby damage the tyre. The factors which cause such failures are not easy to capture experimentally due to their complexity and high experimental cost. Hence, this study focused on using the finite element method to model the solid resilient tyre, generate stresses, and identify the failures and regions that are susceptible to damage. Initially, a tyre static model was developed and validated using laboratory experimental data obtained from the industry. The validation results showed that the numerical results are in good agreement with the experimental data. Subsequently, the model was extended to incorporate impact simulation. The simulation considered tyre motion and impact on three different types of kerbs - angular, circular and square-shaped. Simulation results showed that high stresses occur mainly in the side walls of the solid tyre while high contact pressure and high in-plane shear stresses were observed at the tread layer especially when it moves on square-type kerbs. Hence, there is a high tendency for tyre failures to occur due to the side wall cracks at the base layer. Furthermore, sudden wear and damage due to chunking can be expected on the tread layer of the solid resilient tyre.

Keywords: Hyper-elastic models, impact simulations, nonlinear finite element model, solid resilient tyres.

INTRODUCTION

Solid resilient tyres are mainly utilized in forklift vehicles and often operate in harsh environmental conditions while bearing heavy loads. In these rugged environmental conditions, the tyres experience high stresses, which may lead to sudden wearing and damage due to chunking and rupturing that occur when the tyre impacts on kerbs or obstacles. High experimental cost and time are the main impediments to investigating such behaviour of solid tyres using experimental setups. In this study, finite element (FE) numerical modelling approaches have been utilized to model and predict the solid tyre impact behaviour. The main components of solid resilient tyres and their functions are illustrated in Figure 1. The base layer and embedded bead bundles of the solid tyre ensure good gripping contact with the tyre rim. The internal heat generation and traction of the tyre mainly occur in the cushion layer and tread layer, respectively.

Most of the literature has focused on the numerical analyses of pneumatic tyres (Han *et al.*, 2019; Azizi, 2020). Even for those analyses, the impact analyses were performed for rectangular cleats or kerbs only. Wei and Olatunbosun (2014) studied the impact analysis of a pneumatic tyre and their findings showed that when the height of the rectangular obstacles was increased the amplitude of the vertical force and longitudinal force on the tyre and the tyre deformation were also increased. Suvanjumrat and Rugsaj (2020) investigated impact force and vertical deformation of non-pneumatic tyres when impacted on rectangular cleats with three different heights. Moreover, impact forces on the axle of a pneumatic tyre against a rectangular kerb were analysed by including the suspension system with a full

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car model (Shiraishi et al. 2000). Furthermore, Reida et al. (2007) used a detailed truck numerical model to study the pneumatic tyre impacts on a rectangular kerb as well as its behaviour when it was moving over rocks and culvert grates. Kamoulakos and Kao (1998) analysed the similarities of numerical and experimental readings on vertical and horizontal spindle forces when a pneumatic tyre moved over a rectangular cleat. In Chang and Yang (2009), the generation of ductile fractures on a wheel has been observed numerically as well as experimentally, by impacting it on a solid rectangular cube. Their results showed that the damage to the wheel could be accurately predicted using maximum strain energy density based on the plastic work due to the impact. Duni et al. (2008) reported the contact pressure behaviour of a complete car numerical model with pneumatic tyres when it was moving on a pothole and fatigue reference track. Han et al. (2019) conducted a transient dynamic analysis to study the pneumatic tyre behaviour when it impacted on a rectangular cleat. The results showed that large contact pressure accumulated in the two edges of the tread layer where it made contact with the road. In a more recent study of Han et al. (2020), they utilized a quarter vehicle model with pneumatic tyres to investigate the structural vibration and acoustic radiation under rectangular cleat impact excitations. They observed localized contact pressure on the cleat impact area, and widely distributed contact pressure around the contact region in the tyre when the tyre was on the cleat and just before it passed



Figure 1: Components of solid resilient tyre

through the cleat. In Azizi (2020), a cleat impact experimental test was performed by using a pneumatic tyre and test drum. In this study, the frequency spectrum of the cleat impact was obtained as a function of tyre contact patch length, cleat size, and rolling speed.

Few studies have been performed to analyse static and dynamic behaviours of solid resilient tyres. Suripa and Chaikittiratana (2008), investigated the strain energy density and stress distribution of solid tyres under static conditions by using simplified 3D FE solid models. The results showed that the highest strain energy accumulated in the cushion layer of the solid tyre. Dechwayukul et al. (2010) also conducted laboratory tests to analyse solid tyre durability to evaluate tyre service life. Phromjan and Suvanjumrat (2018a) presented the development of a detailed 3D FE model to analyse the deformations and stress distributions of a solid tyre under different load levels in static conditions. In their results, high stress was observed in the base layer of the solid tyre. Phromjan & Suvanjumrat (2018b) conducted lab experiments to compare the contact forces and vibration analyses of a rolling solid tyre and a pneumatic tyre under different rolling velocities. It was identified that higher contact forces and vibration levels were generated in solid tyres than in pneumatic tyres. To the best of the authors' knowledge, there have been no dynamic numerical FE models developed to study the impact of solid resilient tyres on kerbs and predict their failures.

The structure of this paper is as follows: In the methodology section, the development procedures for static and impact models of a solid resilient tyre are presented. The next section presents the model validation, analysis of results, and discussion. The final section provides concluding remarks and possible future research directions.

MATERIALS AND METHODS

3D FE model of the solid tyre

A CAD model of a solid resilient tyre was first developed as shown in Figure 2. A detailed numerical model of the solid resilient tyre was then developed for the static analysis using Abaqus 6.14 software (ABAQUS Inc., 2014) and the procedure is shown in Figure 3.

In the detailed numerical model, the tyre constraints were applied on nodes which were in between the three rubber layers. Surface-to-surface contact was incorporated to assign interactions between contact surface and tread layer of the tyre. A mesh convergence analysis was performed to identify the suitable mesh size of the tyre, as shown in Figure 4. The mesh of all components in the tyre was modelled by using 8-node 3D hexahedral elements and the converged model contains



Figure 2: Static numerical model of solid resilient tyre



Figure 3: Methodology flow chart of static FE model development

in total 71685 solid elements. The inner surface of the base layer was assigned as a rigid body to represent the tyre rim. Loads were applied to the centre of the tyre.



Figure 4: Mesh convergence with vertical deformation

Selection of best-fitted hyper-elastic material models

Suitable hyper-elastic material models are required to develop the static model of the solid tyre. The best-fitted material models are utilized to describe the mechanical behaviour of rubber layers in the solid tyre. In this study, the best-fitted material models for base layer, cushion layer, and tread layer were selected by combining the curve fitting method and three standard measures of accuracy (MAPE: Mean Absolute Percentage Error; MAD: Mean Absolute Deviation; and MSD: Mean Squared Deviation). Based on the results of curve fitting, the Mooney-Rivlin, Ogden, and Yeoh hyperelastic material models were obtained as the best-fitted material models of the base layer, cushion layer and tread layer, respectively. Equations 1, 2, and 3 represent the general form of the Mooney-Rivlin material model, Yeoh material model, and Ogden material model, respectively (Mooney, 1940; Ogden, 1972; 1973; Yeoh, 1990; 1997).

$$U = C_{10}(\bar{I}_1 - 3) + C_{01}(\bar{I}_2 - 3) + \frac{1}{D_1}(J_{el} - 1)^2$$
...(1)

$$U = \sum_{i=0}^{3} C_{i0} (\bar{I}_{1} - 3)i + \sum_{i=1}^{3} \frac{1}{D_{i}} (J_{el} - 1)^{2i} \dots (2)$$

$$U = \sum_{l=1}^{N} \frac{2\mu_{l}}{\alpha_{l}^{2}} (\bar{\lambda}_{1}^{\alpha_{l}} + \bar{\lambda}_{2}^{\alpha_{l}} + \bar{\lambda}_{3}^{\alpha_{l}} - 3) + \sum_{l=1}^{N} \frac{1}{D_{1}} (J_{el} - 1)^{2l}$$
...(3)

Where:

 λ_1, λ_2 and λ_3 : Principle stretches of principle directions; C_{ij} and D_{ij} : Material constants which control the shear behaviour and bulk compressibility of the material; J_{el} : Elastic volume ratio; \bar{I}_n : Invariants; μ_i and α_i : Temperature-dependent material coefficients

Table 1 presents the best-fitted hyper-elastic material models and their coefficient values for each rubber layer of the solid tyre.

Dynamic impact simulation of the solid tyre

The validated static numerical model of the solid resilient tyre was utilized to implement the impact numerical model of the tyre. Figure 5 illustrates the steps that were followed to develop the tyre impact model. The *Abaqus/Explicit* solver has been utilized to generate this model. The *Abaqus* results transferring capability was employed to transfer results from *Abaqus/Std*. to *Abaqus/Explicit*. Furthermore, relevant boundary conditions and velocities were also applied to the tyre model. Since solid tyres usually move short distances at low velocities, 5 kmh⁻¹ and 10 kmh⁻¹ translational velocities were accounted as impact velocities of the tyre model. Finally, the generation of maximum stress in the rubber layers, maximum contact pressure, and shear stress due to friction were determined at the time of tyre impacts on three types of kerbs. In these impact models, the tread layer of the tyre was ignored to avoid convergence issues and to reduce the simulation time.

The relaxation property of each rubber layer was assigned by considering the viscoelastic behaviour of the rubber material. The corresponding viscoelastic properties were obtained by utilizing the dynamic mechanical analyzer (DMA) as shown in Figure 6 and frequency sweep test data.

 Table 1:
 Coefficients of best-fitted hyper-elastic material models for tyre base, cushion and tread layers

Rubber layer of the tyre	В	ase	Cushion				Tread				
Material model	Moone	y-Rivlin			C	Ogden				Yeoh	
Coefficients	<i>C</i> ₁₀	C_{01}	μ_1	μ_2	μ_2	α1	α2	α ₃	<i>C</i> ₁₀	C ₂₀	C ₃₀
	0.6	2.5	-9.3	8.9	0.9	1.5	2.1	-3.7	0.7	-0.1	0.1



Figure 5: Flow chart of impact FE model development



Figure 6: Dynamic mechanical analyser (Eindhoven University of Technology, 2021)

Table 2: Coefficients of Prony series for tyre base layer, cushion layer and tread layer

Rubber layer of		Tł	nree-term Pron	ny coefficient	s	
the tyre	g_{I}	ψ_{I}	g_2	Ψ_2	$g_{_3}$	Ψ_{3}
Base	0.2208	0.0003	0.1208	0.0146	0.0993	0.1890
Cushion	0.0016	0.0001	0.1099	0.0173	0.0603	0.2198
Tread	0.1545	0.0032	0.1223	0.0377	0.1253	1.8288

The corresponding storage modulus (G') and loss modulus (G'') data of the rubber materials were obtained using the frequency sweep test. Equations (4) and (5) express the relationship between Prony series coefficients related to storage modulus and loss modulus of the material, respectively.

$$G'(\omega) = G_0\{[1 - \sum_{n=1}^N g_n] + [\sum_{n=1}^N \frac{g_n \omega^2 \psi^2}{1 + \omega^2 \psi^2}]\} \qquad \dots (4)$$

$$G''(\omega) = G_0 \{ \sum_{n=1}^{N} \frac{g_n \omega \psi}{1 + \omega^2 \psi^2} \} \qquad ...(5)$$

Where:

 ω : Rolling frequency of the tyre

 G_0 : Instantaneous shear modulus

 g_n and ψ_n : Prony series coefficients

The obtained three-term Prony series coefficients are presented in Table 2. Subsequently, the tyre impact model was implemented using the validated static numerical model of the solid tyre as shown in Figure 7.



Figure 7: Impact numerical model of solid tyre

Selection of kerbs

According to the literature review, most of the impact analyses were performed for pneumatic tyres with square-type kerbs. Hence, it is necessary to select suitable kerb geometries to perform the impact analyses on solid tyres. Equal height and three different types of kerb geometries were introduced onto the contact surface as shown in Figure 8. The kerbs were selected based on their practical usage (Hills Kerbs PTY Ltd., 2021). The following kerb types are usually utilized on the edges of pavements, roads and parking areas.



Figure 8: Three types of kerbs: (a) circular type, (b) angle type and (c) square type



Figure 9: Experimental and numerical vertical deformation data of the solid tyre



Figure 10: Experimental and numerical horizontal deformation data of the solid tyre

RESULTS AND DISCUSSION

Validation of the solid tyre numerical model

The developed static model was validated by applying different loads on it and comparing corresponding deformation readings with laboratory experimental data. Figures 9 and 10 illustrate the vertical and horizontal deformations of the numerical tyre model and actual tyre.

Table 3: Readings of three measures of accuracy

Type of the deformation	MAPE (%)	MAD	MSD
		(%)	(%)
Vertical	2.91	0.66	0.46
Horizontal	3.00	0.29	0.12

Table 3 shows the values of three measures of accuracies calculated between experimental and numerical results. According to the Table 3 results, the Mean Absolute Percentage Error (MAPE) value for vertical and horizontal deformations are 2.19% and 3.00%, respectively. The readings of Mean Absolute Deviation (MAD) and Mean Squared Deviation (MSD) also show very small values (<1%). Hence, the values of the three measures of accuracy results confirmed that the behaviour of the developed tyre model is in good agreement with the actual tyre behaviour.

The maximum von Mises stress accumulation in the rubber components of the solid tyre under angle type, circle type, and square type kerb geometries is illustrated in Figures 11, 12, and 13, respectively. According to Figure 11, a maximum von Mises stress reading of 2.915 MPa was observed in the side wall rubber compound of the solid tyre when the tyre impacted on an angle-type kerb.

Furthermore, according to Figure 12, a maximum von Mises stress value of 2.861 MPa is detected in the side walls of the solid tyre when it impacted on a circle-type kerb.

Moreover, a maximum von Mises stress reading of 3.089 MPa is obtained in the side walls of the solid tyre when it impacted on a square type kerb, as shown in Figure 13.

Among these three types of kerbs, the solid tyre experiences a higher impact stress on the square-type kerb than on the other two kerb types. However, during the impact the fillet radius on the square-type kerb can significantly affect the generation of maximum stress in a solid tyre. Figures 14 and 15 illustrate the variation of maximum impact stress in the solid resilient tyre with 5 kmh⁻¹ and 10 kmh⁻¹ impact velocities, respectively. Figures 14 and 15 show that impact with the square-type kerb has resulted in a higher value of the maximum stress compared with angle and circle-type kerbs under both impact velocities.



Figure 11: Stress generation of the solid tyre when it impacted on an angle type kerb (load 19614 N and velocity 10 kmh⁻¹)



Figure 12: Stress generation of the solid tyre when it impacted on a circle type kerb (load 19614 N and velocity 10 kmh⁻¹)



Figure 13: Stress generation of the solid tyre when it impacted on a square type kerb (load 19614 N and velocity 10 kmh⁻¹)



Figure 14: Maximum impact stress history of solid tyre rubber components with 5 kmh⁻¹ velocity



Figure 15: Maximum impact stress history of solid tyre rubber components with 10 kmh⁻¹ velocity



Figure 16: Maximum impact stress history of bead wires at velocity 10 kmh⁻¹



Figure 17: Deformation history of solid tyre at velocity 10 kmh⁻¹

Furthermore, it can be observed that the maximum value of the impact stress is increasing with increasing impact velocity. According to the above stress analyses, it can be seen that the high impact stresses lead to sidewall crack propagations in the solid tyre rubber components. Figure 16 illustrates the maximum stresses obtained in the bead wires of the tyre when it travels over the three types of kerbs. It is noticed that there is no significant difference among three maximum stresses generated in the bead wires based on the three types of kerbs. In addition, their peak stress values are well below the material yield stress (1765 MPa) of the bead wires.

The vertical deformations of the solid tyre for the three types of kerbs are presented in Figure 17. In the

dynamic condition of the tyre, the maximum vertical deformations that can be observed are 80.1 mm for the square type kerb, 79.2 mm for the angle type kerb, and 68.6 mm for the circular type kerb. At 0.25 seconds, the tyre experiences the highest vertical deformation (80.1 mm) when it is moving over the square-type kerb.

Analysis of contact pressure generation of the solid tyre

In this section, the contact pressure of the solid tyre is examined to investigate its effect on tyre wear and tread layer damage during impact with the three different kerbs types, as shown in Figure 18.





tyre contact patch than the middle.

According to the contact pressure analysis, the highest contact pressure (3.438 Nmm⁻²) is observed when the tyre impacts on the square type kerb. Furthermore, the distribution of the contact pressure throughout the tyre contact patch varies with the type of impact kerb. Based on the contact pressure analysis, the highest wear rate and damage on tread layer can be expected from the tyre when it impacts on the square type kerb compared to the other two types of kerb. In the circle type kerb, damage in the tyre tread layer occurs closer to the edges of the

Analysis of in-plane shear stress generation of the solid tyre

The in-plane shear stress generation in the tyre circumferential direction is presented in Figure 19.

According to Figure 19, high in-plane shear stress (1.297 MPa and 1.314 MPa) is observed when the tyre impacts on the circular type and square type kerbs, respectively. This leads to sudden wear and damage due to chunking in the region of the tread layer. Based on



 Figure 19:
 In plane shear stress (CSHEAR1 in MPa) generation at tread layer of the solid tyre (load - 19614 N and velocity

 - 10 kmh⁻¹): (a) impact on angle type kerb; (b) impact on circle type kerb and (c) impact on square type kerb

above stress analyses and contact pressure analyses, it is clear that the sudden impacts of the solid tyre on squaretype kerbs have a relatively more adverse effect on the durability of the tyre.

CONCLUSIONS

This study was conducted to investigate the stress generation and failure regions of solid resilient tyres as they moved on angle, circle and square-type kerbs. The analyses were performed on stress histories, contact pressure and in plane shear stress generation of tyre rubber layers and side walls, under two different moving velocities. To obtain the behaviour of tyre impact stress histories, a validated numerical model of a tyre was developed. The numerical model was used for tyre impact analyses on the three different kerbs by importing static results into the Abaqus/Explicit interface. Based on the tyre impact analyses results, relatively higher stresses were observed in the tyre sidewalls (3.089 MPa) when it impacted on square type kerbs. The results show that there is a high tendency for the solid tyres to fail due to side wall crack propagation when the tyre moves on square-type kerbs. Furthermore, high contact pressure (3.438 Nmm⁻²) and high in-plane shear stress (1.314 MPa) were observed at the tread layer of the tyre when it moved on a square-type kerb. These findings highlighted that impact of solid tyres with square-type kerbs can significantly decrease the service life and durability of solid tyres compared to other kerb types. Therefore, this model can be reliably used to identify and analyse improvements to the solid tyre design in order to reduce the probability of failure due to kerb impact. This study can be further extended to observe the rolling resistance and heat generation of the solid resilient tyre as well as the behaviour of tyre impact due to changes of material properties and geometry parameters.

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RESEARCH ARTICLE

Bioremediation

Screening of potential aerobic denitrifying bacteria for nitrate removal from water

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Abstract: Nitrate pollution in groundwater is a common problem in areas where inorganic fertilizer is used to a large extent. This situation seriously affects communities that use ground water as their main source of drinking water and for many other purposes. Therefore, finding an efficient and cost effective system for the removal of nitrate from groundwater is an urgent necessity. The present study was aimed at identifying aerobic bacteria isolated from various soils and water sources and to test their potential for reducing nitrate in groundwater. The bacterial isolates (n = 128) were screened for nitrate reduction by various processes in nutrient broth and in mineral salt medium containing glucose and starch, using KNO, as the nitrate substrate. Liberated gases during nitrate reduction were analyzed using gas chromatography. Out of 128 morphologically different isolates, two strains, namely Paracoccus sp. (A2) and Bacillus sp. (A19), were selected for further analysis on the basis of their performance for water treatment. The nitrate reduction percentages of A2 and A19 were within the range of 59.63-100% and 86.67-100%, respectively. Gas chromatography results indicated that these two strains liberated a higher percentage of N₂ (68 - 90%) compared to N₂O (5-13%) and CO₂ (traces) while reducing the amount of nitrate. These results confirmed that A2 and A19 have the potential to be used in bioremediation of nitrate contaminated groundwater.

Keywords: Bioremediation, contamination, groundwater, isolates, nitrate, treatments

INTRODUCTION

Nitrate is a harmful pollutant that has become a common water contaminant in many parts of the world

(Rajta et al., 2019; Zhang et al., 2019b; Abascal et al., 2021). Excessive use of nitrogen-rich fertilizers used for agricultural purposes, discharge of poorly treated domestic and industrial wastewater, livestock manure, and leachate from landfill sites are the main anthropogenic sources for nitrate pollution of groundwater (Gutierrez et al., 2018; Tokazhanov et al., 2020). The World Health Organization (WHO) reconfirmed the safe level of nitrate as below 50 mg/L for drinking water, which was set to protect against methaemoglobinaemia (WHO, 2017). Excessive consumption of nitrates can cause health effects in humans and animals alike, especially methaemoglobinaemia (blue baby syndrome) in infants and gastrointestinal cancer in adults (Ren et al., 2018; Cotruvo, 2017). There are reports of other health disorders, including increased infant mortality, hypertension, central nervous system birth defects, diabetes, spontaneous abortions, respiratory tract infections, and changes to the immune system due to the consumption of high levels of nitrates (Kotopoulou et al., 2022).

Nitrate contamination of groundwater is a burning issue in many areas in the world such as UK (Neal *et al.*, 2006), Australia (Rasiah *et al.*, 2013), North America (Power & Schepers, 1989), Morocco, Changshu in China (Sadeq *et al.*, 2008), and Toyserkan in western Iran (Jalali, 2011). The problem is more severe in some regions of South East Asia. Karunanidhi *et al.* (2021) reported that synthetic fertilizers, cow dung and sheep manure, industrial discharge, septic tank leakage, and

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municipal solid waste disposal are the major sources of nitrate pollution in India. They reported that around 117.93 million people drink water contaminated with nitrate levels between 45-100 mg/L and 108.2 million people consume water with levels more than 100 mg/L in India. It is also reported that nitrate contamination in groundwater is a major concern in the coastal region of Bangladesh (Jannat *et al.*, 2022).

Similar to many parts of the seasonally dry tropical areas of the world, the Jaffna peninsula in Sri Lanka also experiences minimal periods of rain, while ground water is the only source of water for drinking (Prabagar *et al.*, 2020). The water quality in the area has been drastically reduced due to intensive inorganic fertilizer use, resettlement and urbanization within the last decade (Piyathilake *et al.*, 2022).

Due to the above reasons protection of groundwater quality is an important concern confronting much of the world's population. Though sophisticated technologies such as chemical denitrification (Xu *et al.*, 2017), ion exchange (Vandekerckhove *et al.*, 2018), reverse osmosis (Epsztein *et al.*, 2015), electro-dialysis and catalytic denitrification (Zhang *et al.*, 2016) can be used to remove nitrate from groundwater, proper, cost effective and environmentally friendly systems need to be adopted for remediation of nitrate from ground water in the developing countries. Therefore, finding appropriate treatment technologies for nitrate removal is critical. Biological denitrification is the most promising approach currently investigated for treatment of nitrate contaminated water.

Microbial denitrification has been proven to be an advanced, high performance, and highly selective method for complete nitrate elimination (Gomez et al., 2000b). Biological denitrification is the most important and widely used method to treat nitrate wastes as it enables the conversion of nitrogen compounds into harmless dinitrogen (N2) gas (Costa et al., 2018). A microbial consortium composed of Cellulosimicrobium sp., Aeromonas veronii, Lysinibacillus sphaericus, and Rhodococcus rhodochrous was found to be the most efficient bacterial consortium for reducing nitrate in rubber latex wastewater (Dey et al., 2019). Biological denitrification utilizes the anaerobic reduction of oxidized nitrogen compounds through the sequential activity of microbial reductase enzymes, finally converting them to harmless nitrogen gas. Four enzymes, namely, nitrate reductase, nitrite reductase, nitric oxide reductase, and nitrous oxide reductase, are responsible for the complete reduction of nitrate ion to dinitrogen gas (Pang & Wang,

2021). A variety of incomplete denitrification pathways exists. Some denitrifying bacteria reduce both nitrates and nitrites, while others reduce only nitrite. Some produce only dinitrogen, some produce a mixture of dinitrogen and nitrous oxide, while others produce only nitrous oxide.

Even within a single species such as Pseudomonas fluorescens, the biotypes differ in the end product of the pathway. Although nitrate reduction activity is exhibited by diverse microbial genera, with a range of heterotrophic and autotrophic metabolisms, the aerobic nitrate reducers belong to a variety of groups of heterotrophs (Guo et al., 2013). Aerobic denitrification gained attention due to its easier operation and higher nitrate reduction efficiency compared to anaerobic denitrification (Chen et al., 2012). The most predominant denitrifying bacteria that are reported in our environment belong to the genus Pseudomonas. There are reports on aerobic denitrifying species isolated from environmental samples such as ponds, canals, soils, and activated sludge (Patureau et al., 2000; Huang et al., 2020). Pseudomonas aeruginosa has been extensively studied genetically (Wu et al., 2013) and therefore is usually considered a favorable organism to be used in studies on denitrification in wastewater treatment plants. However, as it is an opportunistic pathogen, it could not be utilized for drinking water treatment processes.

The present study investigates the possibility of utilizing microorganisms isolated from the environment to recover NO_3^- contaminated ground water from Jaffna, Sri Lanka. We hypothesized that microorganisms isolated from different environments would be efficient and capable for this purpose. The expected result would be applicable for the remediation of groundwater resources in Sri Lanka and other countries in the same region of South Asia.

MATERIALS AND METHODS

Sources for the isolation of bacteria

Soil and water from submerged paddy fields and ponds, and wet soil enriched with partially decomposed manure from the Jaffna peninsula (9.6615°N, 80.0255°E) were used for the isolation of bacteria.

Isolation and screening of aerobic denitrifiers

Nitrate rich modified nutrient broth consisting of (g/L): Peptone 5.0, NaCl 5.0, KNO₃ 1.0, glucose 1.0, yeast extract 5.0, and beef extract 5.0 with pH 7.2 was used for the enrichment of nitrate reducers. The enrichment cultures were plated on modified bromothymol blue (BTB) medium containing 0.1% L-asparagine, 0.1% KNO,, 0.1% KH₂PO₄, 0.005% FeCl₂.6H₂O, 0.02% CaCl₂.2H₂O, 0.1% MgSO₄.7H₂O, 1 mL of BTB (1% in ethanol), 2% agar, and 0.5% glucose at pH 7.0, to isolate and screen the denitrifier under aerobic conditions (Takaya et al., 2003). Culture plates were incubated at 30 °C for 3 ds. Well defined bacterial strains on the basis of their colony and different morphological characteristics were selected for further screening. A nutrient broth with KNO, was used to determine denitrification activity at the initial screening stage. Each strain was inoculated into a 15 mL screw cap tube containing the sterile nutrient broth and Durham's tube, and was incubated at 30 °C for 48 hours (Guo et al., 2013).

Nitrate reduction in synthetic medium

Nitrate removal activity of five selected strains based on nitrate reduction in the nitrate broth medium was evaluated in a synthetic mineral salt medium (MSM) consisting of potassium dihydrogen phosphate (0.1 g/L), dipotassium hydrogen phosphate (1 g/L), ammonium chloride (0.5 g/L), calcium chloride (0.005 g/L), magnesium sulphate (0.1 g/L), and sodium silicate (0.05 g/L), the pH being adjusted to 7.2, with either glucose or starch as the carbon source in three different percentages, *viz;* 0.25 %, 0.5 % and 1.0 % (Ayyasamy *et al.*, 2007).

Analytical methods

Nitrate content of the bacteria inoculated sample was determined according to Anderson & Ingram (1993) via reaction with salicylic acid and sodium hydroxide followed by spectrophotometry at 410 nm. The amount of nitrite was measured through the reaction with sulphanilic acid and N,N-dimethyl-1-naphthylamine, followed by spectrophotometry at 520 nm, as described by Blaszczyk (1993). Ammonium ion content was determined by the method described by Guo *et al.* (2013).

Gas chromatographic analysis

Gases evolved during denitrification were analyzed for N_2 , N_2O , and CO_2 by a gas chromatograph equipped with Shimadzu GC 9 AM analyzer and a thermal conductivity detector with helium as the carrier (Green *et al.*, 2010). Column temperature, detector temperature and injector temperature were maintained at 50-200 °C, 200 °C and 175 °C respectively and the gas flow rate was kept at 30 mL/min. The mixture of gases evolved during nitrate reduction was collected and identified by comparing the retention time of the peaks with standards.

Identification of isolated bacteria

Standard physiological and biochemical characteristics, such as colony morphology, cell shape, gram reaction, catalase reaction, oxidase reaction, motility, nitrate reduction test, anaerobic growth, glucose acid test and starch hydrolysis were used for the identification of bacteria according to the methods described by Bergey (1994).

Evaluation of selected bacterial strains for removal of nitrate from contaminated well water with starch as the carbon source

Based on efficient nitrate removal in the carbon sources and negative growth on Maconkey agar medium, two bacterial strains (A2 and A19) were selected for further study with five nitrate contaminated water samples. Based on optimization of two different carbon sources and different percentages of the synthetic medium, 0.5% starch was selected for water treatment. Conical flasks with 100 mL of 0.5% (0.25 g) starch were sterilized by autoclaving at 120 °C for 20 min. A water sample of 50 mL was filter sterilized using 0.45 µm syringe filter and was added into each flask containing 0.5% (0.25 g) sterile starch and mixed well. Bacterial isolates were cultured on nutrient agar for 24 - 48 h and cell suspensions were prepared by suspending the cultures in 5 mL of sterile distilled water and the turbidity was adjusted to OD 0.5. Each culture suspension (0.5 mL) was added aseptically into each flask and incubated at 30 °C and kept at 120 rpm in a shaking incubator for 72 h. The same conditions were provided without the cultures as the control. Nitrate and nitrite concentrations were analyzed every 12 h in all samples.

Data analysis

All the experiments were carried out in triplicate and the results were analyzed by analysis of variance (ANOVA) using SAS statistical software version 9.1. Treatment means were compared using Duncan's multiple-range test at a significance level of 0.05.

RESULTS AND DISCUSSION

Screening of aerobic denitrifying bacteria

Out of 128 morphologically different strains isolated from different sources, 70 strains were capable of forming blue colonies on the BTB agar plates (Table 1) due to an increase of pH on the medium (Wu *et al.*, 2013).

Sources of isolation	Number of	Number of	
	strains isolated	nitrate reducers	
Municipal compost (GMC)	12	11	
Pond soil	8	1	
Fish waste (GFW)	15	1	
Municipal solid waste	17	4	
dumping place (MSW)			
Manure (swine, poultry)	17	6	
Paddy soil (KPS)	23	17	
Paddy water (KPW)	18	14	
Unutilized well water (PSW)	8	8	
Compost (COM)	10	8	
Total	128	70	

 Table 1:
 Number of isolates and nitrate reducers from different sources

Among the 70 strains, 38 showed gas bubble formation in Durham's tubes. After rescreening, the nitrate removal efficiency was examined in nutrient broth with nitrate under aerobic conditions. Out of 38 strains, 5 bacterial isolates, capable of reducing either nitrate or nitrite efficiently (more than 50%) were selected by quantitative screening. Since most of the denitrifiers were heterotrophs, they required carbon sources for energy consumption (Pang & Wang 2021). Although five strains had higher nitrate reduction capacity (more than 70 %) either with glucose or starch, three of them were grown on Maconky agar medium, hence, cannot be used for treatment of water samples. The other two, A2 and A19 were tested with five nitrate contaminated water samples for their nitrate removal efficiency.

 Table 2:
 Physical and biochemical identification of bacterial strains

Identification test	A2	A19
	(Paracoccus sp.)	(Bacillus sp.)
Shape	Round	Rod
Gram staining	Negative	Positive
Endospore formation	Negative	Positive
Motility test	Negative	Positive
Catalase test	Positive	Positive
Oxidase test	Positive	Positive
Anaerobic growth	Positive	Positive
Nitrate reduction test	Positive	Positive
Growth on Maconkey agar	Negative	Negative
Glucose acid test	Positive	Positive
Starch hydrolysis test	Positive	Positive

Identification of the selected bacterial strains

According to the physical and biochemical identification, strains A2 and A19 were identified as *Paracoccus* sp. and *Bacillus* sp. (Table 2).

Optimization of the percentage of the carbon sources for aerobic nitrate removal

The two carbon sources used, namely glucose and starch at the levels of 0.25%, 0.5%, and 1%, were tested for carbon source optimization for the strains A2 and A19. The optimum carbon percentage was decided based on efficient nitrate removal as well as lowest intermediate accumulation (nitrite) in the medium.

Figure 1 shows the effect of glucose (a) and starch (b) on nitrate removal and nitrite accumulation by the strain A2. The nitrate nitrogen concentration was lowered with both carbon sources at all three levels tested. Although nitrate reduction was observed with time, after 24 hours, nitrite began to accumulate with glucose at concentrations of 0.25%, 0.5%, and 1% up to 60 hours. However, when starch was used as the carbon source, nitrite accumulation was not observed at all three levels.

Strain A2, grown with 0.5% and 1% starch reduced nitrate to 11 mg/L after 36 hours of incubation (Figure 1). Although nitrate reduction was observed with glucose as the carbon source, nitrite accumulation also was recorded at all three levels. When the two carbon sources were compared at three different levels, 0.5% starch exhibited higher nitrate reduction and nitrite was not detected. Ammonium ion was not detected at any levels of the two carbon sources. Therefore, it could be concluded that the strain A19 has a higher potential to remove nitrate with 0.5% of starch, without accumulation of the nitrite intermediate. Further, the nitrate reduction efficiency and intermediate accumulation of strain A2 could be controlled by the organic carbon sources, nitrate concentration, and C/N ratio. A study conducted by Blaszczyk (1993) clearly stated that the denitrification performance of Paracoccus denitrificans strongly depended on the quality of the medium.

The carbon source can provide the energy for the aerobic denitrifier and electron donors for their growth and metabolism (Li *et al.*, 2020b). Nitrogen transformation of strain A19 with the carbon sources glucose (a) and starch (b) is shown in Figure 2. The results indicate that, after 60 hours of incubation, complete reduction of nitrate was observed with 0.5% and 1% of starch, without the accumulation of nitrite. However, with glucose,

nitrate was reduced to below the permissible level with high amount of nitrite accumulation. Intermediate accumulation mainly depends on the carbon sources. In a study, N_2O and NO accumulation in the presence of nitrite during denitrification was observed with acetate-fed denitrifying cultures, but not in methanol- or ethanol-fed denitrifying reactors with excessive carbon source supply (Lu *et al.*, 2014). At the end of 60 hours of incubation, almost all the nitrate was found to be in the form of nitrite with glulose as the carbon source. Therefore, it could be concluded that nitrite concentration also varied

with the types and levels of carbon sources for each strain. Among the two carbon sources, starch possessed significantly higher nitrate reduction efficiency for both strains. It may be due to the presence of amylolytic enzymes in the two strains (A2 and A19) and the ability to utilize starch as the carbon source. Ayyasamy *et al.* (2007) also stated that the amylolytic enzymes present in organisms can utilize the starch well. In another study conducted by Rajakumar *et al.* (2008), it is reported that the denitrification rate was higher for starch than glucose, acetic acid, cellulose, and sucrose.



Figure 1: Effect of glucose (a) and starch (b) on nitrate and nitrite concentration in MSM inoculated with strain A2 up to 60 hours of incubation

Therefore, the results indicate that the process of complete denitrification is not a stable process and depends on several factors such as bacterial strains and types and amounts of carbon sources. This unstable nature of the process is due to the influence of these factors, which have been reported in many previous studies (Li *et al.*, 2020; Yin & Yan, 2020). Furthermore, nitrate removal is a highly dynamic process that can

be affected by temperature, pH, C/N ratio, dissolved oxygen concentration, and bacterial population (Chen *et al.*, 2006; Olaya-Abri *et al.*, 2021; Zhou *et al.*, 2021). In another bacterial denitrification study conducted by Gomez *et al.* (2000), the nitrite accumulation was higher when sucrose was used as the carbon source, but it was not found when methanol and ethanol were used.



Figure 2: Effect of glucose (a) and starch (b) on nitrate and nitrite concentrations in MSM inoculated with strain A19

Aerobic nitrate removal in well water contaminated with nitrate

Five water samples having initial nitrate nitrogen concentrations of 57.37 mg/L (S1), 32.72 mg/L (S2), 26.70 mg/L (S3), 20.81 mg/L (S4) and 15.91 mg/L (S5) were tested with either A2 or A19. Changes in the nitrate nitrogen concentration of different water samples treated with A2 and A19 with 0.5% of starch are shown in Figures 3.

Figure (3a) shows the nitrate reduction profile of water sample S1, having an initial nitrate nitrogen concentration of 57 mg/L, treated either with A2 or A19. With the sample treated with A19, nitrate was found to be at a safe level after 60 hours and not detected at 72 hours. During nitrate reduction by A19, nitrite formation increased at a high rate up to 48 hours and declined thereafter. Although, nitrate was not detected at 72 hours, nitrite was not at a safe level (3.89 mg/L). Accumulation of nitrate at early stage by aerobic denitrifiers such as

Paracoccus denitrificans, which had 14.07 mg/L initially and reduced to zero at 40 hours (Zhang et al., 2020) has been reported. The accumulation of NO₂⁻-N in S1 treated with strain A19 is possibly due to high initial nitrate content of the water and the consecutive lag of nitrite reduction (Chen et al., 2020), which could be possibly removed with more time. Significant difference was observed in final nitrate concentration in the sample S1 treated with either A2 or A19. Further, A2 reduced nitrate nitrogen to the level of 9.1 mg/L at 72 hours without accumulation of nitrite. Paracoccus sp. (A2) expressed greater aerobic nitrate removal capacity similar to numerous aerobic bacteria. For instance, Paracoccus denitrifcans strain removed 90.00% of NO₂-N in a 250 mg/L initial NO₃^{--N} medium (Medhi et al., 2018). Similar aerobic nitrate removal (87.63%) was achieved by Paracoccus denitrifcans strain Z195 (Zhang et al., 2020). Paracoccus thiophilus strain LSL 251 had an aerobic denitrification rate of 5.90 mg/L/h in nitrate rich medium (Chen *et al.*, 2020).



Figure 3: Bacterial reduction of nitrate nitrogen in well water in the presence of 0.5 % starch a) S1, b) S2, c) S3, d) S4 and e) S5

Both strains reduced nitrate with time without nitrite accumulation in sample S2 (Figure 3b). Reduction of the nitrate nitrogen concentration from 32.71 mg/L to 7.30 mg/L was observed in 72 hours with A2, while 4.36 mg/L was observed with A19 at the same time. In another water sample having an initial nitrate nitrogen concentration of 26.7 mg/L, the nitrate level was reduced below the safe level (11 mg/L) at 24 hours by the strain A2, while it reached 1.09 mg/L with A19 at 36 hours, without nitrite being detected (Figure 3c). Moreover, when the initial nitrate concentration was about 20.81 mg/L both strains A2 and A19 lowered the nitrate concentrations of water sample S4 and S5 treated with the strains are shown in Figures

3d and 3e respectively. Initial concentrations in water samples S4 and S5 were 20.81 mg/L and 15.91 mg/L, respectively.

During incubation, fluctuation was observed in the nitrate nitrogen concentration with the treatment of either A2 or A19, however, no nitrite was detected in either sample with strain A2. This might be due to the oxidization of the negligible amount of nitrite, resulting from the brief exposure to air during sampling (Zhang *et al.*, 2011) or the simultaneous nitrification and denitrification capability of the strains (Kim *et al.*, 2008; Khardenavis *et al.*, 2007). The denitrification process can also be influenced by metal ions such as Fe³⁺ and Mo⁶⁺ (Pintathong *et al.*, 2009). As can be observed from the

figures, both strains of A2 and A19 with 0.5% of starch lowered the nitrate nitrogen concentration from various initial levels of nitrogen in water. Nevertheless, the time needed to attain the permissible level or below varied with strains and the well water samples. Although the difference is not significant, it might be due to the effect of initial nitrate concentration, pH, and availability of other nutrients in the water sample (Körner & Zumft, 1989). Denitrification of synthetic waste water having a high nitrate level was inhibited at the pH values of 6.5 and 7.0. Although, higher nitrate reduction was achieved with the increased pH values of 7.5, 8.0 and 9.0, accumulation of nitrite increased significantly (Glass & Silverstein, 1998). A Bacillus pumilus strain removed 99.7% of NO₂ – nitrogen in a 70 mg/L initial NO₂⁻ containing medium (Elkarrach et al., 2021). In another study, 89.4% of nitrate removal was reported by Bacillus sp. after a 48-h cultivation in a sole N-source medium with initial nitrogen approximately 20 mg/L (Huang et al., 2017). Bacillus sp. PB8 showed excellent aerobic denitrifying ability (0.25 mg/L/h) both in artificial media and real wastewater treatment (Barman et al., 2018).

Gas chromatographic analysis

During denitrification by A2, a higher percentage (90.6%) of N2, lower percentage of N2O (5.7%) and traces of CO2 were released, while A19 released 68.5% of N2 and 12.5% of N2 O. Considering the composition of gases released, A2 would be a better strain than A19.

CONCLUSION

Among the 70 nitrate reducing bacteria strains isolated, Paracoccus sp. A2 and Bacillus sp. A19 exhibited a high nitrate reduction potential either with glucose or starch as the carbon source. However, with glucose, in all three levels (0.25%, 0.5% and 1%), nitrite accumulation was observed. Among the three percentages of starch (0.25%, 0.5% and 1%), 0.5% was the optimum level for efficient nitrate removal. Further, Paracoccus sp. A2 and Bacillus sp. A19 were capable of removing the nitrate nitrogen content in the range of 15 mg/L to 57.37 mg/L in groundwater to safe levels within 72 hours with 0.5% of starch. To the best of our knowledge, this is the first report on nitrate reduction by bacterial strains in Sri Lanka and this finding may provide useful information for the potential use of these two bacterial species. Further studies are required to validate nitrate removal efficiency of the strains under different physicochemical conditions.

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Data availability statement

The data that support the findings of this study are available from the corresponding author upon reasonable request.

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RESEARCH ARTICLE

Dairy Technology

Physicochemical and sensory attributes of inulin incorporated set yoghurt prepared using cow and buffalo milk

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Abstract: The demand for probiotic food products has been increased over the last few decades. The current study aimed to evaluate the microbiological and physicochemical properties of two set yoghurt types prepared using cow milk (CM) and buffalo milk (BM) with three different concentrations of inulin (1%, 2% and 3% w/v). Two types of starter cultures BB12 (Bifidobacterium bifidum - BB) and YC-X11 (Streptococcus thermophilus - ST and Lactobacillus bulgaricus subspp. Bulgaricus - LB) were used as probiotic sources. Probiotic viable cell counts, syneresis, total acidity, pH, organoleptic properties, and viscosity were evaluated during 21 days of storage at 4 °C. Both inulins incorporated CMY and BMY showed significantly high (p < 0.05) ST and LB counts at the end of the enumeration period (~8.0 log cfu/mL) while control samples of each group showed significantly lower (~7.0 log cfu/mL) probiotic counts. 3% CMY and BMY reported the highest probiotic survivability. Significantly (p < 0.05) high viscosity was observed in 2% inulin incorporated yoghurts which were highly acceptable and showed better stability during storage. This study concludes that the addition of inulin improves the overall probiotic survivability of the tested microorganisms while ensuring consumer acceptance without drastically altering the physicochemical properties of yoghurt.

Keywords: *Bifidobacterium*, buffalo milk, inulin, *Lactobacillus*, probiotic, yoghurt.

INTRODUCTION

The consumer approach towards eating healthy foods has evolved during the last decade, enhancing the potential to develop various functional foods that provide more benefits than simply nutritional value. Among the functional foods, much attention has been focused on probiotic products. Probiotics are health-promoting viable microorganisms such as Lactobacillus acidophilus, Bifidobacterium bifidum, and Bifidobacterium longum that exhibit a beneficial effect on human health, by improving the intestinal microbial balance (Martin-Diana et al., 2003). In addition, probiotics improve lactose utilization by producing lactase enzyme which helps to digest lactose, preventing intestinal infections by producing organic acids and other antibacterial agents, reducing cancer risk, improving immunity, lowering cholesterol and blood pressure, and protecting against diarrhoea, constipation and osteoporosis (Ranadheera et al., 2010). Several studies have demonstrated the modes of actions of probiotics as antioxidants, antimicrobial compounds, and agents that improve bowel functions (Beena et al., 2012).

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In order to ensure the survival of probiotic strains, most probiotic-containing foods also include prebiotic sources. A prebiotic is defined as a 'non-digestible food ingredient that beneficially affects the host by selectively stimulating the growth and/or activity of one or a limited number of bacteria in the colon, thus improving the health of the host' (Gibson et al., 2004). Ingestion of prebiotics significantly enhances the proliferation of colon microflora by acting as a substrate for probiotics, while improving the colonization rate of bacterial strains such as Lactobacillus and Bifidobacterium which have high probiotic activity. The prebiotics can balance the composition of microflora in the human gut (Ranadheera et al., 2010). In many foods, non-digestible oligosaccharides and fructooligosaccharides are used as prebiotics. The most utilized prebiotic in yoghurt manufacturing is inulin. It is a non-digestible and fermentable fructan (Ehsani et al., 2016) that stimulates the growth and activity of probiotic bacteria. Inulin consumption significantly increases faeces loading and improves faecal excretion in humans. Thus, inulin is effective against colon disorders such as constipation.

In the food industry, inulin is often used as a lowcalorie sweetener and a non-digestible fibre that forms gels which increase viscosity and enhance organoleptic properties (Meyer et al., 2011). Considering these beneficial health effects imparted by both prebiotic compounds and probiotic microbial strains, current food manufacturers tend to produce symbiotic foods. Yoghurt is such a food that is widely consumed and produced by incorporating prebiotics with the probiotics Lactobacillus acidophilus, Lactobacillus casei, and Bifidobacterium to enhance their therapeutic value. The viability of the probiotic bacterial count should be as high as 106-107 CFU/mL at consumption to achieve the aforementioned benefits (de Souza Oliveira et al., 2011). Hence, it is essential to ensure sufficient and viable probiotic bacteria throughout the shelf life of the yoghurt. Recent studies have shown that most probiotic foods have a low population of probiotic bacteria at the time of consumption. The main dairy source of yoghurt is fresh cow milk. When compared to cow milk, buffalo milk has a higher content of fat, protein, lactose and total solids (Abesinghe et al., 2020). These properties of buffalo milk give a rich flavour and taste and make it very suitable for the manufacture of a wide variety of milk products (Han et al., 2012b). Moreover, recent studies on the shelf life of buffalo milk yoghurt have indicated that it could be a good vehicle for probiotics (Han et al., 2012a). As a result, the potential of utilizing buffalo milk to produce a symbiotic yoghurt was examined. The present study

aims to explore and compare the physicochemical and probiotic activity of cow and buffalo milk set yoghurt to identify the best probiotic carrier with a suitable level of prebiotic inulin addition.

MATERIALS AND METHODS

Preparation of set type yoghurt

Cow milk set yoghurt (CMY) and buffalo milk set yoghurt (BMY) were prepared according to the following methodology. Milk was pre-heated at 65 °C and inulin powder (Orafti® HPX, High-Performance Inulin Powder, Mannheim, Germany) was added in three concentrations of 1%, 2% and 3% (w/v). Then, skim milk powder (SMP) was added at concentrations of 3% (w/v) for the control, 2% (w/v) SMP for 1% inulin treated sample and 1% (w/v) SMP for 2% and 3% inulin treated samples respectively. The mixtures were homogenized for 15-20 min at 5 Mpa. The milk was then pasteurized at 90-95 °C for 5-10 min followed by cooling down to 36 °C. A starter culture containing Streptococcus thermophilus (ST) and Lactobacillus delbrueckii subsp. bulgaricus (LB) (Chr. Hansen YC-X11, Hoersholm, Denmark) and Bifidobacterium bifidum (BB) (Chr. Hansen, Hoersholm, Denmark) were added in 0.002% (w/v) concentration. The yoghurts were incubated in cups at 42 °C for 4 h. Finally, the samples were stored at 4 °C pending analyses.

Preparation of microbiological isolates

The pour plate method was used to evaluate the growth and viability of yoghurt bacteria, and bifidobacterial (Shori & Baba, 2012) MRS media (M641, HIMEDIA, Mumbai, India) was used to isolate Lactobacillus. Commercial M17 agar base (HIMEDIA, Mumbai, India) was used for isolation of the S. thermophilus strain, and commercial Bifidobacterium agar (HIMEDIA, Mumbai, India) was used to isolate Bifidobacterium strains of the samples. A dilution series was prepared by using 0.1% (w/w) peptone (CDH, India) solution for all microbiological analyses. All culture media were incubated at 35-37 °C for 48-72 h in the laboratory incubator (IN-601/GEMMYCO) in anaerobic conditions. Colony counts were obtained at 1, 7, 14, and 21 ds by hand enumeration through a colony counter (Witeg Colony Counter Galaxy 330, Germany). For each microbial strain, three nearby dilutions were prepared and for each dilution, duplicate plates were prepared and enumerated. The microbial counts were expressed as log colony forming units per millilitre (log CFU/mL).

Physicochemical analyses

pH measurement

Changes in the pH during the storage period were measured using a multi-parameter pH meter (35 Series, Eutech Instruments, Singapore). The pH probe was immersed in 25 mL of homogenized sample and the readings were taken in relation to room temperature $(27\pm2 \ ^{\circ}C)$ in triplicate. Readings were collected on the 1st, 7th, 14th, and 21st ds after manufacturing.

Titratable acidity

A slurry of each sample was made by mixing 10 g of yoghurt with 5 mL of distilled water. Samples were titrated against 0.1 N NaOH solution, in the presence of 1% phenolphthalein until a persistent pink colour was observed. Acidity is expressed as a percentage of lactic acid (Jayarathna *et al.*, 2020). Analysis was carried out in triplicate.

Syneresis

Twenty grams of each sample was placed on a Whatman No.1 filter paper and the weight of the filtrate was measured for 2 h with 30 min time intervals in triplicate. Syneresis is expressed as the percentage of the initial sample (Amal *et al.*, 2016).

Texture

Texture analyses were completed using a programmable digital viscometer (Brookfield DVE Digital Viscometer, Massachusetts, USA) at 4 °C, using spindle numbers 6 and 7 at speed of 10 rpm. Readings were recorded after 30 s and expressed in millipascal-second (mPa.s) on the 7th day after manufacturing.

Sensory evaluation

Sensory evaluation of inulin incorporated cow milk and buffalo milk yoghurt was carried out with 30 untrained panelists (aged 22-55 years, women 60%, men 40%) according to a randomized design in order to find out the degree of likeability of the probiotic yoghurt with the most preferable level of inulin. The study was carried out at least 2 h after a main meal and smokers, betel chewers, or alcohol consumers were not recruited as panelists. Each panelist evaluated the final yoghurt product according to a five-point hedonic scale (1-5, from strongly disliked to strongly liked, respectively) using a simplified and structured sensory evaluation ballot at 25 °C.

Statistical analysis

Physicochemical data of the yoghurt were analysed by using one-way ANOVA of Minitab 18 software (Minitab[®] 17.1.0, UK) with a 95% confidence interval (significance at the p < 0.05) while the Friedman test was used to analyze sensory data. Microbiological data were analysed via one way ANOVA of the SAS program (Version 9.0, SAS Inc, North Carolina, USA).

RESULTS AND DISCUSSION

Changes in the microbial count during the storage

In consideration of variations of viable ST counts in CMY during the end of storage, there was a significant difference (p < 0.05) of viability between 2% and 3% inulin incorporated yoghurt samples in contrast to the control and 1% inulin incorporated samples (Table 1). The highest ST count throughout the storage period in both CMY and BMY was recorded from 3% inulin incorporated samples (9.11 ± 0.07 log CFU/mL to 8.72 ± 0.14 log CFU/mL). There was no significant difference in initial viable counts among the samples, although there was a significant reduction of the viable counts in both 2% and 3% CMY on the 21st day of storage. The viable counts in CMY followed a gradual decline towards the end of the storage period, yet they were remaining in the range of critical microbial count which is >10⁶ CFU/mL.

It is evident that ST has dominantly grown in both cow milk and buffalo milk set yoghurts. Oliveira *et al.* (2002), reported that ST predominated in all products, showing higher counts than 9 log CFU/mL in fermented milk prepared with mixed culture.

The yoghurt samples prepared with buffalo milk showed a significant difference (p < 0.05) in microbial viability of ST among inulin incorporated samples and the control sample at the end of storage. The highest viable count on the 21st day of storage was reported from the 3% inulin incorporated BMY sample (8.23 \pm 0.12 log CFU/mL). Considering the BMY, they showed no significant difference (p > 0.05) compared to their initial viable count throughout the storage up to the 14th day, yet they showed a significant reduction (p < 0.05) when comparing the initial stage and the final day of storage. However, the viable ST counts were still above the critical count. This prominent and better viability of ST is likely due to its enhanced proteolytic activity combined with higher lactose utilization, compared to the other two probiotic strains (Özer & Kirmaci, 2010).

Treatment	Period of storage in days (Mean \pm SD)				
Cow milk yoghurt	1	7	14	21	
Control	$8.34\pm0.14^{\rm Ba}$	$8.27\pm0.17^{\rm Ca}$	$8.13\pm0.15^{\rm Ba}$	$7.93\pm0.12^{\rm Ca}$	
1% Inulin	$8.46\pm0.1^{\rm Ba}$	$8.49\pm0.06^{\rm BCa}$	$8.41\pm0.12^{\rm Ba}$	$8.27\pm0.17^{\rm Ba}$	
2% Inulin	$8.82\pm0.14^{\rm ABa}$	$8.75\pm0.13^{\rm ABab}$	$8.65\pm0.09^{\rm Bab}$	$8.49\pm0.06^{\rm ABb}$	
3% Inulin	$9.11\pm0.07^{\rm Aa}$	$9.08\pm0.08^{\rm Aa}$	$9.04\pm0.08^{\rm Aab}$	$8.78\pm0.12^{\rm Ab}$	
Buffalo milk yoghurt					
Control	$8.13\pm0.15^{\scriptscriptstyle Ba}$	$8.08\pm0.09^{\rm Ca}$	$7.78\pm0.33^{\rm Bab}$	$7.54\pm0.16^{\rm Bb}$	
1% Inulin	$8.38\pm0.09^{\rm ABa}$	$8.31\pm0.10^{\rm BCa}$	$8.13\pm0.15^{\rm ABab}$	$7.93\pm0.12^{\rm Ab}$	
2% Inulin	$8.63\pm0.12^{\rm ABa}$	$8.58\pm0.11^{\rm ABa}$	$8.43\pm~0.15^{\rm Aab}$	$8.13\pm0.15^{\rm Ab}$	
3% Inulin	$8.72\pm0.14^{\rm Aa}$	$8.65\pm0.10^{\rm Aa}$	$8.55\pm0.14^{\rm Aab}$	$8.23\pm0.12^{\rm Ab}$	

 Table 1: Viable counts of Streptococcus thermophilus (ST) in cow and buffalo milk set yoghurt stored at 4 °C during 21 days of storage (log CFU/mL)

Means in the same row without a common simple letter superscript significantly (p < 0.05) differ for each treatment; means in the same column without a common capital letter superscript significantly (p < 0.05) differ among the treatments per microbial analysis.

Table 2:	Viable counts of Bifidobacterium bifidum (BB) in cow and buffalo milk set yoghurt stored at
	4 °C during 21 days of storage (log CFU/mL)

Treatment	Period of storage (Days) (Mean \pm SD)				
Cow milk yoghurt	1	7	14	21	
Control	$8.31\pm0.10^{\rm Ba}$	$8.23\pm0.12^{\rm Ca}$	$8.13\pm0.15^{\rm Ba}$	$7.69\pm0.21^{\rm Bb}$	
1% Inulin	$8.63\pm0.12^{\rm ABa}$	$8.49\pm0.06^{\rm Ca}$	$8.49\pm0.14^{\rm ABa}$	$8.31\pm0.10^{\rm Aa}$	
2% Inulin	$8.75\pm0.13^{\rm Aa}$	$8.67\pm0.07^{\rm ABab}$	$8.60\pm0.08^{\rm Aab}$	$8.32\pm0.1^{\rm Ab}$	
3% Inulin	$8.91\pm0.15^{\rm Aa}$	$8.86\pm0.13^{\rm Aa}$	$8.78\pm0.12^{\rm Aa}$	$8.65\pm0.04^{\rm Aa}$	
Buffalo milk yoghurt					
Control	$8.00\pm0.21^{\rm Ba}$	$8.67\pm0.10^{\rm Bab}$	$8.46\pm~0.10^{\rm Bb}$	$7.39\pm0.21^{\rm Bab}$	
1% Inulin	$8.35\pm0.04^{\rm Aa}$	$8.31\pm0.10^{\rm Aa}$	$8.28 \pm 0.05^{\rm Aa}$	$8.17\pm0.21^{\rm Aa}$	
2% Inulin	$8.58\pm0.05^{\rm Aa}$	$8.46\pm0.10^{\rm Aa}$	$8.41\pm0.12^{\rm Aa}$	$8.35\pm0.04^{\rm Aa}$	
3% Inulin	$8.67\pm0.10^{\rm Aa}$	$8.46\pm0.10^{\rm Aa}$	$8.39\pm0.21^{\rm Aa}$	$8.27\pm0.17^{\rm Aa}$	

Means in the same row without a common simple letter superscript significantly (p < 0.05) differ for each treatment; means in the same column without a common capital letter superscript significantly (p < 0.05) differ among the treatments per microbial analysis.

According to Table 2, it is evident that the addition of inulin, the duration of storage, and their collective interaction have had significant (p < 0.05) effects upon the BB viability in both CMY and BMY samples. Especially, considering the 1st and 21st days of CMY samples, the control yoghurt sample has reported the lowest BB viability (8.31 ± 0.10 log CFU/mL and 7.69±0.21 log CFU/mL, respectively) in contrast to other

samples, although the viable BB count has not remarkably varied among the inulin incorporated samples. A recent study by Kamel *et al.* (2021) concludes that increasing inulin concentrations helped to amplify the viability of BB significantly (p < 0.05) during storage, as well as acting as an antifungal agent of probiotic yoghurt, suggesting that inulin can be used as a suitable and a successful nutrient for BB.

Table 2 shows that significantly lower (p < 0.05) viable BB counts were observed with time in each BMY sample, compared to their initial counts, though the viable counts were reported above the critical microbial count up to the 21st day. Similarly, Han *et al.* (2012a) reported that *Bifidobacterium* spp. remained viable throughout 10 weeks of storage when incorporated in

low-fat symbiotic yoghurt. The highest viable counts observed in 3% inulin incorporated yoghurt sample on 1st and 21st days of storage (8.67 \pm 0.1 log CFU/mL to 8.27 \pm 0.17 log CFU/mL) suggesting that BM yoghurt could provide a suitable environment for the growth of probiotic microbes and maintain therapeutic levels until the end of the storage period.

 Table 3:
 Viable counts of Lactobacillus bulgaricus (LB) in cow and buffalo milk set yoghurt stored at 4 °C during 21 days of storage (log CFU/mL)

Treatment	Period of storage (Days) (Mean \pm SD)					
Cow milk yoghurt	1	7	14	21		
Control	$8.28\pm0.05^{\rm BCa}$	$8.08\pm0.09^{\rm Ca}$	$7.93\pm0.12^{\rm Bab}$	$7.69\pm0.21^{\rm Bb}$		
1% Inulin	$8.54\pm0.06^{\rm ACa}$	$8.43\pm0.15^{\rm BCa}$	$8.39\pm~0.21^{\rm Aa}$	$8.27\pm0.17^{\rm Aa}$		
2% Inulin	$8.79\pm0.10^{\rm ABa}$	$8.82\pm0.07^{\rm ABa}$	$8.52\pm0.09^{\rm Aab}$	$8.39\pm0.21^{\rm Ab}$		
3% Inulin	$8.96\pm0.16^{\rm Aa}$	$8.86\pm0.08^{\rm Aab}$	$8.74\pm0.11^{\rm Aab}$	$8.59\pm0.13^{\rm Ab}$		
Buffalo milk yoghurt						
Control	$8.08\pm0.09^{\text{Ca}}$	$7.93\pm0.12^{\rm Cab}$	$7.80\pm0.70^{\rm Bb}$	$7.54\pm0.39^{\rm Bc}$		
1% Inulin	$8.45\pm0.07^{\rm Ba}$	$8.38\pm0.09^{\rm Ba}$	$8.27\pm0.17^{\rm Aa}$	$8.04\pm0.27^{\rm ABa}$		
2% Inulin	$8.64\pm0.02^{\rm ABa}$	$8.44\pm0.07^{\rm ABab}$	$8.31\pm0.10^{\rm ABbc}$	$8.13\pm0.15^{\rm Ac}$		
3% Inulin	$8.73\pm0.09^{\rm Aa}$	$8.59\pm0.13^{\rm Aab}$	$8.50\pm0.16^{\rm Aab}$	$8.31\pm0.10^{\rm Ab}$		

Mean values in the same row without a common simple letter superscript significantly (p < 0.05) differ for each treatment; means in the same column without a common capital letter superscript significantly (p < 0.05) differ among the treatments per microbial analysis.

According to Table 3, there is a significant difference (p < 0.05) in LB viability between inulin incorporated samples and control samples made with cow milk. The viable counts of LB ranged from $8.28 \pm 0.05 \log$ CFU/mL to 7.69 \pm 0.21 log CFU/mL on the 1st and 21st day in storage, while the highest viable LB counts were observed in 3% inulin incorporated yoghurt sample, on the 21st day of storage ($8.59 \pm 0.13 \log \text{CFU/mL}$). However, the viability of LB counts significantly declined with time compared to their initial values yet remained in the critical microbial count even on the final day of storage. A similar study (Saccaro et al., 2009) revealed that the viability of LB was reduced by 1.51 log CFU/mL when grown in co-culture with ST and in association with probiotic microorganisms, which explains the receding nature of LB viability in these mixed culture CMY samples.

The viability of LB in BMY samples also shows a significant difference between inulin incorporated and control samples, where the viable counts of LB in the control sample gave values of $8.08 \pm 0.09 \log \text{CFU/mL}$

and 7.54 \pm 0.39 log CFU/mL on 1st and 21st day of storage, respectively. However, a significantly different (p < 0.05) value for the highest viability of LB was reported from the 3% inulin incorporated sample on the 21st day, which was 8.31 \pm 0.1 log CFU/mL. Significantly low viable counts were observed in each treatment compared to their initial counts and the rate of reduction of viable counts was higher in the control sample though the counts remained in the range of critical microbial count up to the 21st day (>10⁶ CFU/mL).

The results of the study showed the increase of probiotic bacterial counts with increasing inulin concentrations for both milk types. Inulin is a fructan that can be effectively used as a nutrient for microbial growth (Srisuvor *et al.*, 2013). Inulin enhances probiotic proliferation by acting as dietary fibre (Ramirez-Santiago *et al.*, 2010). In addition, lactic acid bacteria (LAB) are capable of producing exopolysaccharides (EPS) during their growth, and these EPS are essential for LAB-host interaction mechanisms such as bacterial tolerance to gut conditions and their immune-modulatory activities,

which helps them to survive in the host's gastrointestinal tract (Caggianello *et al.,*, 2016), The increase in inulin levels may lead to high concentrations of EPS amounts with a positive effect on microbial growth.

Lower viability was observed for LB bacteria in both voghurt types, which may be due to the post acidification (Sendra et al., 2008) leading to lower pH and low availability of bacteria in the commercial starter culture (Chr. Hansen YC-X11, Hoersholm, Denmark). The results of the present study showed that the viability of microbes decreased with time, yet the decrease of LAB viability was less with high concentrations of inulin. These results agree with Ehsani et al. (2016), who also observed that counts of probiotic strains declined over four weeks of storage. Donkor et al. (2007) suggested that ultimate pH and accumulation of organic acids had a notable impact on the survival of LB and BB in yoghurt. Therefore, considering the results obtained in this study, reduction of energy source with time would be another reason for the declining survival rate of microbes with time, apart from previously mentioned factors.

Changes in pH, titratable acidity, syneresis and texture

Variations of pH in different yoghurts

According to the pH variation data of CMY, there were no significant differences (p > 0.05) among the samples throughout the storage period and a decrease of pH during storage was observed in all the treatments with time (Figure 1a). At the end of the experiment, the control sample recorded the lowest pH (4.13 \pm 0.01) while the 1% inulin incorporated sample recorded the highest pH value (4.24 \pm 0.01).

When considering the pH variation of BMY, there were significant differences (p < 0.05) in pH among the treatments throughout the storage period (Figure 1b). The pH varied from 4.86 ± 0.01 to 4.96 ± 0.00 from the 1st day to the 21st. On the 14th day, 1% CMY recorded a significantly higher (p < 0.05) pH value compared to other samples (4.97 ± 0.01). However, the pH at the end of the storage period of yoghurts manufactured with inulin addition showed no significant difference (p > 0.05) among samples.



Figure 1: a) Variation in pH of CMY with different inulin concentrations stored at 4 °C during 21 days of storage b) Variation in pH of BMY with different inulin concentrations stored at 4 °C during 21 days of storage.

ABC Means with different uppercase letters are significantly different between each treatment.

^{abc} Means with different lowercase letters are significantly different between days of storage.

In yoghurt processing, lactic acidification is the result of lactose fermentation by the associative growth of the two thermophilic, homofermentative, lactic acid bacteria, ST and LB. Increasing H⁺ concentration in the yoghurt due to acid formation leads to low pH. Similar results were reported by Tiwari et al. (2015). The accumulation of organic acids contributes to the continuous reduction of pH throughout the storage. Results of the present study also showed that pH decreased with increasing inulin levels. Continued acidification during the storage can be linked to a progressive transformation of lactose into lactic acid. However, in yoghurts containing high soluble fibre components, other mechanisms could also promote a further increase in acidity level (Ramirez-Santiago et al., 2010). According to Sendra et al. (2008), fibre addition itself contributes to a decrease in pH but the increased metabolic activity of the probiotics would also contribute to drastic pH reduction. Increasing inulin levels lead to high bacterial growth, consequently increasing acid production, which tends to lower the pH over time.

Changes in titratable acidity of yoghurts

The titratable acidity (TA) of CMY showed significant differences (p < 0.05) between control and inulin added yoghurt samples while there were no significant differences in acidity among treatments throughout 21 days of storage period (Figure 2a). Throughout the storage, the control sample reported the highest titratable acidity levels whereas, the 1% sample reported the lowest titratable acidity values.

According to data of BMY, there was no significant difference in TA between the control sample and inulin added yoghurt samples throughout the storage period. A significant (p < 0.05) increase of TA during storage was observed in all the treatments. The TA values were ranged from 0.94 - 0.95% and 0.96 - 0.97% during the storage period of CMY and BMY, respectively. The 1% inulin incorporated BMY sample showed a lower TA throughout the storage period while the 3% inulin incorporated BMY sample showed the highest TA. However, there



Figure 2: a) Variation in titratable acidity of CMY with different inulin concentration levels stored at 4 °C during 21 days of storage. b) Variation in titratable acidity of BMY with different inulin concentration levels stored at 4 °C during 21 days of storage.

^{ABC} Means with different uppercase letters are significantly different between each treatment. ^{abc} Means with different lowercase letters are significantly different between days of storage. was an overall increase in titratable acidity while the pH dropped marginally throughout the storage period. These phenomena indicate continuously ongoing enzymic and metabolic activities of mixed starter cultures in low-temperature storage. These results agree with the findings of Canbulat and Ozcan (2015) who also reported a similar type of pH drop and proportional increase in titratable acidity with inulin incorporated set yoghurts. These results confirmed that there is no drastic change in both pH and titratable acidity with the incorporation of inulin into CM and BM yoghurts.

Variations of syneresis in yoghurt during storage

The results of CMY showed a significant (p < 0.05) difference in syneresis among treatments throughout the storage period (Figure 3a). On the 1st day, the highest syneresis was observed in the 3% CMY sample (32.51 ± 0.50%) and the lowest value in the 2% CMY (30.37 ± 0.38%). On the 21st day, the lowest syneresis was observed with the yoghurt produced with the 2% inulin CMY (14.3 ± 0.28%), while the highest value was observed in the 3% inulin CMY (16.3 ± 0.0%).

With BMY, the control sample showed significantly (p < 0.05) lowest syneresis (23.81 \pm 0.08%) while the 3% inulin BMY sample gave the highest syneresis value (26.12 \pm 0.10%) on the 1st day of storage. At the end of the storage period, the highest syneresis of 14.3 \pm 0.28% was observed in the 3% inulin inclusion, while the lowest syneresis value of 9.27 \pm 0.28% was observed in the 2% inulin incorporated BMY sample. A significant (p < 0.05) decrease of syneresis during the storage time was observed in all the BMY yoghurts throughout the storage period.

Inulin, when incorporated into the protein matrices, forms micro-crystals that interact with each other to form small aggregates. These aggregates are capable of encapsulating water, decreasing the whey removal (syneresis), while creating a smooth and creamy mouthfeel (Kip *et al.*, 2006). It was observed that the 2% inulin incorporated yoghurt samples displayed a similar pattern of behaviour to that of the control sample when compared with the other two concentrations. The 1% and 3% inulin incorporated CMY and BMY samples had higher syneresis levels than the 2% inulin yoghurts.



Figure 3: a) Variation in syneresis (%) of CMY with different inulin concentration levels stored at 4 °C during 21 days of storage. b) Variation in syneresis (%) of BMY with different inulin concentration levels stored at 4 °C during 21 days of storage.

^{ABC} Means with different uppercase letters are significantly different between each treatment. ^{abc} Means with different lowercase letters are significantly different between days of storage.
Perhaps, the 1% inulin concentration was insufficient to from a strong tridimensional network of of insoluble submicron crystalline inulin particles that entraps large amounts of water, hence the rate of syneresis was higher. In contrast, it has been observed that gel stiffness becomes lower when the inulin concentration is increased. Therefore, the highest syneresis levels observed by 3% inulin incorporated CMY and BMY samples can be explained in terms of the weak network resulting from decreased amounts of casein-casein bonds. It was observed that 2% inulin incorporation level has the ability to retain the strength and rigidity of the yogurt gel matrix without affecting its structure or the rate of whey removal in a drastic manner. Amatayakul et al. (2006) observed a reduction in whey separation when using exopolysaccharide (EPS) producing LAB starter cultures. YCX-11 the commercial starter culture used during this study can produce EPS which can facilitate the water binding capacity and improve the yoghurt microstructure (Han et al., 2016). The resulting increase in inulin level could enhance the EPS production reducing the syneresis process compared to the control sample.

Table 4: Changes in the viscosity of CMY and BMY withrefrigerated storage (4 °C) for 21 days.

Treatment	Viscosity (x10 ³ mPa.s) (Mean \pm SD)		
Cow milk yoghurt			
Control	$49.850\pm0.07^{\rm C}$		
1% inulin	$46.75\pm0.07^{\rm D}$		
2% inulin	$55.85\pm0.07^{\rm A}$		
3% inulin	$54.05\pm0.07^{\rm B}$		
Buffalo milk yoghurt			
Control	$70.85\pm0.07^{\rm B}$		
1% inulin	$62.85\pm0.07^{\rm C}$		
2% inulin	$86.45\pm0.07^{\rm A}$		
3% inulin	$86.40\pm0.06^{\rm A}$		

Means in the same column without a common capital letter superscript significantly (p < 0.05) differ for each treatment.

Viscosity changes in yoghurt

Overall, there seems to be an increased viscosity in BMY yoghurts when compared to CMY (Table 4). This phenomenon can be linked to the differences in the two types of milk. The average amount of total solids in buffalo milk is 16.3%, whereas in cow milk it is 12.8%. In addition, buffalo milk has high amounts of protein, ash, fat and lactose content (Abesinghe *et al.*, 2020).

Viscosity is directly related to the curd stability of voghurt. According to Srisuvor et al. (2013), the curd stability of the yoghurt is one of the important physical properties in voghurt matrices. Parameters such as total solids and protein content, the heating process, homogenization, the acidity of yoghurt, and the activity of starter culture bacteria generally influence curd stability. The structure and the chemical composition of the added ingredients mainly affect the strength of the casein aggregate. The generated branching structure enhances the protein and carbohydrate interaction giving the gel better curd stability. According to the results, 2% inulin incorporated voghurt samples had a much higher viscosity when compared to 1% inulin incorporated yoghurts, though there was only a slightly lower viscosity at the 3% incorporation level compared to the 2% incorporation level. Viscosity reduction may be due to the incompatibility of gel structure formed with the interactions of inulin. Increasing inulin levels developed the casein aggregates only up to a certain limit, if inulin addition exceeded the critical value, layer separation could happen, and leading to lower viscosity. Similar observations were made by Helal et al. (2018), where the addition of 1% and 2% inulin increased the viscosity of a low-fat yoghurt by 14.7 % and 17.9% respectively, yet the addition of 3% inulin only showed an 8% increase in the viscosity of the yoghurt samples.

Sensory evaluation

According to Figure 4a), 2% CMY gave the highest preference for the odour, colour, texture, taste and overall acceptability while the control sample showed the lowest acceptability. Figure 4b) shows that panelist perception for odour and taste was higher in 3% IBM while colour, texture and overall acceptability were higher in 2% IBM. The control yoghurt sample had the lowest acceptability. However, the sensory results of the present study directly implied inulin addition improved the organoleptic properties of set yoghurt without affecting negatively the taste, odour and colour of the yoghurt. The addition of inulin influences the perception of 'creaminess.' Therefore, among these samples, with increasing inulin concentrations, the perception of 'creaminess' has increased as well. Increasing inulin levels lead to high acceptability via taste and odour while excess addition of inulin lowers the acceptability due to texture degradation. The acceptability may increase due to inulin addition. Inulin enhances the taste (due to fructose), odour, and colour of the yoghurt, though it adversely affects improving the texture by adding additional total soluble solids to yoghurt. Staffolo et al. (2004) stated that yoghurt supplemented with inulin maintained its colour stability during storage, and Seydim *et al.* (2005) stated that yoghurt made with inulin addition had more perceptive flavour and a smooth texture.



Figure 4: a) Variation of sensory properties in CMY with varying levels of inulin incorporation. CCM, 1% ICM, 2% ICM and 3% ICM samples contain varying levels of inulin, w/w = 0% (Control), 1%, 2%, and 3%, respectively. b) Variation of sensory properties in BMY with varying levels of inulin incorporation. CBM, 1% IBM, 2% IBM and 3% IBM samples contain varying levels of inulin, w/w = 0% (Control), 1%, 2%, and 3%, respectively.

CONCLUSION

The finding of the present study concludes that incorporation of inulin enhances the survival of *S. thermophiles*, *L. bulgaricus*, and *B. bifidum* in set type cow milk and buffalo milk yoghurts during refrigerated storage for 21 days. Cow milk was shown to be a better probiotic carrier than buffalo milk during the storage. It confirms that the addition of 2% inulin into cow milk and buffalo milk yoghurts results in low syneresis without altering the textural properties of the yoghurt matrices. The sensory evaluation results showed that additional inulin concentrations help to increase the 'creaminess' in yoghurts, hence indicating the 2% inulin incorporation as the optimal condition for yoghurt with both cow milk and buffalo milk.

Conflict of interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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RESEARCH ARTICLE

High Voltage Engineering

Fractal characteristics of creeping discharges propagating on nano-epoxy composite insulators immersed in coconut oil

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Abstract : A solid/liquid dielectric interface is considered to be one of the weakest points in a composite insulation system, as it facilitates creeping discharges on the interface when the electric field strength exceeds a threshold value. This paper presents a study on the use of nano epoxy composite to minimize the effect of damage which occurs due to creeping discharge activity, when it was immersed in coconut oil. A point-plane electrode arrangement-based test apparatus, energized by a high voltage supply, was used for analysing the propagation of creeping discharges over solid/liquid interfaces using visual observation. An algorithm was used to determine the fractal dimension of creeping discharges propagating over various solid/liquid insulating interfaces. In particular, it focused on the variation in creeping discharge patterns with the use of pure epoxy and nano-composite epoxy samples. The results showed that the use of nanofillers can increase the dielectric breakdown strength of epoxy by 7%. The results also showed nanofillers of 20 nm with 1% concentration do not affect the tortuosity of the creeping discharge pattern on the solid material immersed in coconut oil. However, it can increase the possibility of flashover due to creeping discharge propagation over epoxy insulating material immersed in coconut oil by around 12%.

Keywords: Coconut oil, creeping discharges, final discharge length, fractals, nanotechnology, propagation.

INTRODUCTION

A quality electricity supply has become one of the basic necessities in modern society, demanding an efficient

and uninterrupted power supply. The failure of critical oil-filled high voltage (HV) equipment such as cables, transformers, capacitors, and circuit breakers with many solid/liquid interfaces may lead to catastrophic events, resulting in economic losses and injury to humans. Therefore, it is important for the power utilities to reliably operate and maintain their high voltage equipment (Kebbabi & Beroual, 2006; Ediriweera et al., 2018a). Dielectric liquids are used in the power industry together with solid insulating material to serve as an insulation, impregnation, and cooling medium. In some cases, the solid barrier materials are used to improve the dielectric strength of gas and oil gaps (Liu et al., 2016; Ediriweera et al., 2018b). Dielectric liquid and solid barriers can increase the dielectric strength of oil-filled HV equipment. However, the arrangement of the composite insulation system is also one of the key points that has to be considered. When the electric field strength exceeds a threshold value, the tangential electric field over the solid/liquid interface initiates surface discharges (Yi, 2012). These discharges are commonly called creeping discharges. This discharge can cause irrecoverable damage to the insulation system and cause flashover which undermines the use of a composite insulation system (Yi, 2012).

A lot of work has recently been devoted to improving the dielectric properties of insulating materials by adding specific nano-fillers to polymer-based materials, changing the dielectric and mechanical properties. Nano-

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composite polymer materials show better performance when it comes to the dielectric properties of the materials (Yanashima *et al.*, 2017; Hornak *et al.*, 2020). In addition to the electrical characteristics, they also show higher tensile strength, thermal stability, and improved chemical properties over conventional insulating materials (Liang & Wong, 2017).

Petroleum-based mineral oil has been used as the dielectric liquid in high voltage equipment for more than seven decades due to its better performance. However, due to some of its disadvantages such as low fire point, limited biodegradability, and high toxicity, attention has been paid to use alternative vegetable oils such as soya bean oil, coconut oil (Abeysundara et al., 2001; Lucas et al., 2002), and sunflower oil, even though their properties such as viscosity, conductivity, acidity, and water solubility are still to be improved. Coconut oil is produced on a commercial scale in Sri Lanka and it is rich in saturated fat with medium and short chains. Most of its physical, chemical, and dielectric properties are at satisfactory levels as an alternative for the conventional mineral oil for high voltage equipment (Abeysundara et al., 2001; Matharage et al., 2013), and it offers better oxidation stability compared to other vegetable oil due to the higher percentage of saturated fatty acids. However, the studies have not progressed to the analysis of creeping discharge over nanocomposite insulating materials immersed in coconut oil, which can affect the performance of the nanocomposite insulation system. Fractal analysis of creeping discharges can be used to compare the amount of discharge branch formation on the advanced solid dielectric materials with that of conventional materials.

The main objective of this study was to perform a fractal analysis of creeping discharges propagating over nanocomposite polymer insulating materials immersed in coconut oil. In particular, the discharge length and the fractal dimension are determined using a test apparatus based on a point-plane electrode arrangement. An experimental setup based on a point-plane electrode system is used to initiate the discharges in a laboratory environment. Pure epoxy and nanocomposite samples are synthesized for testing purposes and a statistical method is used to analyse the breakdown of prepared solid material samples.

MATERIALS AND METHODS

Experimental test setup

In the current study, a point-plane electrode system based test apparatus was adopted in order to initiate creeping discharges and compare their propagating characteristics on solid/liquid interfaces. Needle plane electrode systems are usually selected to experimentally investigate the creeping discharge development (Kebbabi & Beroual, 2006; Dang *et al.*, 2012). The tangential field component of the divergent electric field drives the propagation of creeping discharge as observed inside actual high voltage apparatus (Ediriweera *et al.*, 2018a).



Figure 1: Schematic of the test setup

Figure. 1 shows the schematic view of the test cell and this is similar to what other researchers adopted in their studies (Kebbabi & Beroual, 2006). The test cell consists of a cylindrical core with a height of 100 mm and 130 mm inner diameter, made of clear acrylic to visualize the discharge, and a square-shaped brass plate as its lower cover. The test cell is mounted on a structure made of wood. The test cell has a point-plane electrode arrangement, where the vertical point electrode is held using a movable bearing system to permit vertical movement. The point electrode, made of tungsten, has a diameter of 1 mm and the tip of the point is hemispherical so that the field between the two electrodes would be divergent (Ediriweera et al., 2018a). The gap between the two electrodes can be varied from zero to 20 mm. Thin flat insulating material samples can be maintained horizontally between the two electrodes and the point electrode is configured in such a way that the tip of the point electrode nearly touches the solid material sample. The samples are immersed in an insulating liquid by filling the test cell with oil. A CCD camera, connected to a high-performance video card is mounted over the test cell and the integrated images taken by it are used for the optical observation of the discharge. The plane electrode is grounded and the point electrode is energized by a partial discharge (PD) free HVAC test transformer.

Sample preparation

There are various types of epoxy resins that are used for many engineering applications. Bisphenol - A type epoxy resins do not fall in to the category of cycloaliphatic epoxy resins. The epoxy used consists of diglycidyl ether of bisphenol-A (DGEBA) type Epon@Resin 828. The curing agent is Modified Cyclo-aliphatic Amine Adduct type 3388. The chemical structure of the epoxy resin is shown in Figure. 2. The considered epoxy type has a lower viscosity compared to other types and is commonly used in the high voltage industry. SiO₂ with a particle size of 20 nm is chosen as the filler material of the epoxy-based nano-composite. Both pure epoxy and nano-composite material samples are synthesized and a block diagram representing the synthetic procedure for the nano-composite material is shown in Figure 3 (Kochetov, 2012).

Dielectric samples having silicon dioxide nanoparticle compositions (weight to weight) of 1 % are prepared in

this study for testing purposes. The synthetic procedure is similar to what other researchers adopted in their studies (Kochetov, 2012; Karunarathna *et al.*, 2019). As the first step, required volumes of epoxy and curing agents are measured using a pipette and measuring cylinders. Sonication is a process used to disperse the agglomerate filler particles in an ethanol medium. The mechanical mixing process mixes the epoxy and the filler medium making the fillers transfer into the epoxy. The evaporation process removes the ethanol remaining in the mixture. Finally, in the mechanical process, it mixes the medium after the curing agent is added to the mixture. The fabrication process of pure epoxy consists of simple mechanical mixing of the measured volumes of epoxy and curing agents.

Breakdown voltage of solid samples

The chemical and physical properties of dielectric insulators can affect their breakdown voltage, and breakdown voltage becomes a statistically distributed



Figure 2: Chemical structure of bisphenol-A type epoxy resin



Figure 3: Fabrication process of nano-composite material

quantity. Therefore, statistics are being considered to analyse the breakdown voltage (Gupta & Das, 2017). In the current study, a two-parameter Weibull distribution was used to model and analyse the distribution of the breakdown voltage of the prepared solid insulating samples. The probability density function (PDF) and the cumulative distribution functions are given by equation 1 and equation 2, respectively (Gupta & Das, 2017; Yuan *et al.*, 2017).

$$f(x) = \frac{\beta}{\alpha} \left(\frac{x}{\alpha}\right)^{\beta-1} \exp\left\{-\left(\frac{x}{\alpha}\right)^{\beta}\right\} \qquad \dots (1)$$

$$f(x) = 1 - \exp\left\{-\left(\frac{x}{\alpha}\right)^{\beta}\right\} \qquad \dots (2)$$

where β and α are the shape and scale parameters of the experimental data respectively.

The shape parameter is a measure of the range of experimental data and the scale parameter is a characteristic breakdown voltage at which the probability of failure is 63 %.

Skewness can be used to identify the symmetric or asymmetric nature of breakdown voltage values using equation 3 (Gupta & Das, 2017).

$$skw = \frac{E(x-\mu)^3}{\sigma^3} \qquad \dots (3)$$

Experimental procedure

Flat square shaped solid materials of prepared pure epoxy and silica/epoxy nano-composite of side length 9 cm and thickness 3 mm were used to analyse the creeping discharge propagation over the solid/liquid interface. A type of commercially available copra coconut oil with a relative permittivity of 2.9 was selected as the insulating oil in the study, as it is an emerging alternative liquid insulating material (Matharage et al., 2013). The electric field in the experimental setup used in this study was not uniform. In actual power system assets, a uniform electric field cannot be expected due to the electrodes and dielectric configurations in the equipment. Therefore, tests were carried out to make a comparison between propagation patterns on different interfaces that were subjected to the same electrode arrangement. According to the IEC 60156 standard, a test cell with 2 spherical electrodes separated by a gap of 2.5 mm was used to measure the breakdown voltage of coconut oil samples at the University of Moratuwa, Sri Lanka. The mean value of the breakdown voltage given for five oil samples under ambient conditions was considered as the average breakdown voltage. The average breakdown voltage of 'as received' oil samples was measured at 20 kV.

The test cell was filled with coconut oil and solid materials prepared in square shape were inserted between the two electrodes, such that the point electrode touches the surface of the sample. If the flashover is observed, the test cell was refilled with new oil from the same test oil sample. The samples were kept in the laboratory in a darkened place without any UV exposure in order to avoid contamination. Therefore, the material and electrical properties of the test volumes were assumed to be the same in all the samples in the experiments. The height of the oil volume was kept at 5 cm throughout the testing procedure.

Dielectric strength tests were performed according to the ASTM-D149 standard at the University of Moratuwa, Sri Lanka, and five samples of epoxy and nanocomposite were tested. Parameters of Weibull distribution were estimated using Minitab software.

Fractal dimension

Dimension identifies the way the object fills the space. Dimensions of a point, a line, a square, and a cube are 0, 1, 2, and 3 respectively. Such dimensions are called the Euclidian dimensions. Likewise, fractal patterns have fractal dimensions. As a parameter of characterization of a complex degree of natural phenomena, a fractal dimension is used to describe the surface topography of a fracture and to give a quantitative value to each pattern (Zhou et al., 2014). Several methods can be adapted to calculate the fractal dimension of creeping discharges (Beroual et al., 2014; Ediriweera et al., 2018a). In the current study, the box-counting method was used. Researchers have used this method for purposes of analysis of surface discharges in their findings (Kebbabi & Beroual, 2005). A greater fractal dimension means a more tortuous fracture surface.

One of the common methods called box-counting had been adopted to calculate the dimension of fractal patterns in our previous studies (Kebbabi & Beroual, 2005; Ediriweera *et al.*, 2019). The same method was used in the current study and basic steps were recalled. The original image shown in Figure 4a, obtained in a creeping discharge, was converted into a binary image as shown in Figure 4b. Before the conversion process, all the important traces of the discharge were extracted using photo editing software. The camera did not depict the entire discharge, since the needle was in the centre of the image. However, the pattern in the middle has the least amount of unclear information due to the point electrode, when compared to the visible area as the pattern was recorded to its full extent.



Figure 4: Application of the box-counting method



Figure 5: Application of the box-counting method

However, with the glow around the tip of the point electrode and thick streamers in the centre, individual streamers cannot be distinguished because of the noise captured by the camera. So, information in the middle was ignored in any case, and the box-counting algorithm is applied considering that the morphological variation of the pattern has the same self-similarity throughout the pattern. Fractals reflect the self-similarity of the morphological variation of the pattern. Therefore selfsimilarity can be regarded as implying that at all scales or in all areas of the pattern, it should show the same relative variation of morphology or the fractal dimension. Then a code was applied to the binary image which generates a grid of square boxes over the image.

While changing the side length of the boxes at each step, the number of boxes having any parts of the discharge pattern was counted. Figures 5a and 5b show the binary images covered by squares of side l = 32 pixels and l = 64 pixels, respectively. The slope of the log-log plot of the number of boxes vs. box size gives the fractal dimension as shown in Figure 6.



Figure 6: Total number of boxes N versus the side length l of the boxes obtained from the analysis of an example of a discharge

Discharge length

When it comes to creeping discharges, the discharge may have several mainstreamers starting from its starting point. There also can be side branches starting from mainstreamers. Therefore, each streamer may have different total discharge lengths. What is defined as the final discharge length is the maximum discharge length out of all the streamers of the discharge pattern, and the voltage required to get a streamer of a particular length is assessed. In the study, the camera setup provides the images of the propagating pattern continuously and an algorithm was used to identify the image with maximum extension to measure the final discharge length. It should be noted that each discharge length value presented in the study is an average value measurement taken from four solid dielectric samples



Figure 7: Probability plot for breakdown strength of nano - composite and pure epoxy

Table 1: Parameters of	of Weibull distribution
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Material	Shape parameter	Scale parameter (kV/mm)	Mean (kV/mm)	Skewness
Pure epoxy	40.061	36.551	36.046	0.03
Nano-composite	12.034	40.667	38.975	-0.53

RESULTS AND DISCUSSION

The maximum likelihood estimator is used to estimate the parameters of the distribution of breakdown voltages. Figure 7 shows the probability plot for the breakdown voltages of the material samples.

The shape, scale, mean, and skewness of the distribution are listed in Table 1. The shape parameters are 40 and 12 for pure epoxy and nanocomposite respectively. The shape parameter gives an idea about the failure. A shape parameter lower than one means early failure and when it equal one, then it represents a constant failure rate after commissioning (Gupta & Das, 2017). The skewness of the nanocomposite is a little bit higher than that of epoxy material and the asymmetrical nature of the data points can be seen with non-zero screw values for both of the material samples.

The mean breakdown voltage of epoxy materials has been increased slightly by nano-particles as shown in Table 1. It accounts for around a 7% increment. The behavior of nano-fillers is somewhat different even though microparticles decrease the breakdown strength, acting as centres of charge concentration. The interfacial area of nano-particles is considerably higher and their interparticle distance is quite small. Therefore, they act as barriers to the flow of current through the composite increasing the breakdown strength. Figure 8 shows the development of discharge patterns propagating over epoxy samples and Figure 9 on nanocomposite samples. The basic characteristics of the propagating patterns are similar to the studies with patterns propagating on the solid/mineral oil interface (Kebbabi & Beroual, 2006). Discharge patterns propagating on the interfaces have curved streamers as a result of the electro-hydrodynamic motion of oil in the vicinity of the point electrode and the electric field distorted by space charges on the insulators under AC voltage.

Therefore, streamers try to follow these contours. Observation shows that the discharges propagating on epoxy/coconut oil and nano-composite/coconut oil interfaces have similar tortuous behavior, and, however, with longer discharges on a nano-composite/ glass interface. The kind of the solid material can have a significant effect on the amount of surface discharge.

The most likely reasons for the promotion effect on discharges of the solid insulation materials considered would be the influence of hetero-charges, the permittivity of the material (Yi & Wang, 2015), and nanoparticles (Jang, 2017). Once the surface of a solid material is charged with space charges, it takes some time for charges to dissipate. The dissipation time depends on the characteristics of the material such as surface



Figure 8: Stages of creeping discharge development on an epoxy surface



Figure 9: Stages of creeping discharge development on a nano-composite surface.

conductivity. The effect of both homo-space charges and hetero-space charges, which promote the development of subsequent discharge, should be considered in this scenario, because deposited space charges can last until the negative cycle of the supply voltage (Yi, 2012). The capacitance of the solid insulation affects the discharge inception voltage (Jang et al., 2016). When a flat solid material sample is modeled as a capacitor, its value depends on the permittivity of the material. The increase in the capacitance of the solid material sample enhances the capacitive charges. Also, the permittivity mismatch between the liquid and solid affects the electric field at the tip of the point electrode, which has a direct impact on the ionization probability due to the added effect of change in total polarizability and interfacial polarization, which in turn will affect the propagation of the creeping discharge. The enhanced electric field and charges make the discharge propagate more on the surface of the material. Therefore, as the relative permittivity of the solid material increases, it increases the tendency of creeping discharges to propagate. The dielectric constants of pure epoxy and nano-composite samples are 4.89 and 5.25, respectively. The parallel plate method is used to measure the relative permittivity of the material samples at 100 Hz using a LCR meter, as in previous studies (Mandric et al., 2018; Karunarathna et al., 2019). Therefore, the promotional effect on the nano-composite/ glass interface can be expected.

The final discharge length holds the quasi-linearity with the voltage similar to the previous experiments with mineral oil, as shown in Figure 10. It increases with the dielectric constant of the solid dielectric material. For a given value of the voltage, discharge lengths are higher with nano-composite than with pure epoxy samples. The capacitive effect is more significant over the pattern propagation on the nano-composite material samples immersed in coconut oil.

Nanoparticles have increased the final discharge length corresponding to the nano-composite samples by 4% to 17%, while having a 7.3 % increment in the dielectric constant. So there is a positive correlation between the dielectric constant and corresponding discharge length curves, with coconut oil being the dielectric liquid. Therefore, more damage on the nanoepoxy solid insulator can be expected due to the traces of the discharges.

Table 2: Fractal dimension of creeping discharges

	Epoxy	Nano- composite
Fractal dimension	1.26	1.26



Figure 10: Variation of the final discharge length of the creeping discharges



Figure 11: Variation of the breakdown voltage of coconut oil with the temperature.

Fractal dimension is more related to the amount of the divergence of the streamers and branches of a discharge pattern into smaller ones. Previous studies have shown that when the relative permittivity increases, dimension increases (Beroual *et al.*, 2014). However, according to Table. 2 it can be seen that the nanoparticles considered do not have a significant effect on the amount of ramification on the epoxy/coconut oil interface, as the fractal dimension of patterns propagating on nanocomposite and epoxy material samples have the same values. Therefore, the same amount of tortuousness is found in both cases.

According to previous studies (Abeysundara *et al.*, 2001; Matharage *et al.*, 2013), moisture absorption can reduce the breakdown characteristics of coconut oil.

Therefore, a domestic heater was used to heat the oil samples, and at different temperature values, samples are taken and the dielectric breakdown voltage is measured. After the oil is heated up to 100 °C, it was then allowed to cool down back to room temperature. While the sample was cooling, its dielectric breakdown voltage was measured at several intermediate temperatures. An improvement of dielectric strength was observed after heating the oil sample to over 100 °C.

Figure 11 shows that the breakdown voltage increases gradually up to 60kV when the oil is heated. The water content in the sample affects the dielectric strength of the oil. Heating evaporates water in the oil sample. When the temperature of the sample is near to the boiling point of water (100°C) the water content is very low, hence it shows a higher dielectric strength. When the oil sample is kept open to the air and allowed to cool back, a reduction in dielectric strength was observed as a result of moisture absorption as shown in Figure 11. Therefore, poor dielectric properties can affect the discharge propagation and hence the accuracy of the results. Therefore, the condition of the dielectric liquid on the creeping discharge propagation is an under-researched topic that requires further investigation.

CONCLUSION

This paper presented a study of creeping discharge propagation over epoxy/coconut oil and nanocomposite/ coconut oil insulation interfaces. Experiments were carried out with a point plane electrode arrangement and an image recorder in order to investigate creeping discharges along surfaces of pure epoxy and nanocomposite epoxy samples under an AC divergent field. Results on discharge length, breakdown strength, and fractal dimension have shown that the propagation of creeping discharges depends on the properties of the solid insulator such as dielectric constant and properties of fillers.

Silicon dioxide nanoparticles with 1% concentration were used to increase the dielectric properties of the epoxy material. According to the experimental results, it was evident that it increases the dielectric breakdown voltage of epoxy by 7%. The relative permittivity of the nanocomposite was higher than that of pure epoxy. Therefore, further studies should be carried out with different filler sizes to increase the dielectric properties, especially creeping discharge performance significantly. It was shown that there is a positive correlation between the dielectric constant and corresponding discharge length curves, with coconut oil as the dielectric liquid. Final discharge length increases quasi-linearly with the applied voltage even on materials immersed in coconut oil. According to the discharge length results corresponding to the nano-composite material, it can be concluded that even though nanoparticles can increase the dielectric strength of the epoxy insulators, the capacitive effect is more significant in discharge propagation, and final discharge length has been increased by around 12 %. However, the considered nanoparticle concentration of 1% does not have a significant effect on the amount of ramification of the creeping discharges, because the fractal dimension of 1.26 was observed in creeping discharges propagating on both nanocomposite and epoxy material samples. Overall, it can be concluded that the concentration and the particle size of silicon dioxide

can affect the nano dielectric properties of nano-epoxy composite, such as an increase in breakdown strength and dielectric constant by 7%. A nanocomposite with high breakdown strength is not always satisfactory when it is immersed in coconut oil as the electric field at the tip of the point electrode depends on the permittivity mismatch. The research had shown the fabricated material samples enhance the breakdown strength by 7 % while increasing the possibility of damage to the solid insulator due to creeping discharges by 4% to 17%. Therefore, the materials should be fabricated to minimize the creeping discharge propagation while keeping the dielectric properties within an acceptable level.

Conflict of interest statement

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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RESEARCH ARTICLE

Meteorology

Forecasting the track and the intensity of the cyclone Burevi using WRF

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Abstract: The cyclone 'Burevi' was recorded in the North Indian Ocean in the month of December 2020. In this study a numerical weather forecasting model WRF-ARW was used to forecast the track, intensity, and landfall location and time of the cyclone. The forecasting was performed as 16 consecutive model runs. Each forecast was initiated with different model initializations with a 12 hour time interval. For such initializations, the lateral boundary condition was set by the global forecast system data with a horizontal resolution of $0.25^{\circ} \ge 0.25^{\circ}$ at 3 hour time intervals. The forecasts were conducted up to 120 hours and evaluated using the data obtained from an international best track archive. The forecasted track was evaluated using root mean square error, direct positional error, along-track error and cross-track error. The intensity of the cyclone was evaluated by comparing the minimum sea level pressure and maximum forecasted 10-m wind speed obtained from the positions of the forecasted cyclone track. The WRF-ARW model was successful in forecasting the track and the intensity of the cyclone up to a lead time of 48 hours having a mean direct positional error of less than 106 km. This cyclone recorded three landfalls in Sri Lanka. The accuracy of the forecasted first landfall position and time were within the acceptable range, but the second and third require further investigation.

Keywords: North Indian Ocean, track forecast, tropical cyclone, WRF-ARW.

INTRODUCTION

Weather research and forecasting (WRF) is a numerical weather prediction system used for research and

operational forecasting (Skamarock et al., 2019). Advanced Research WRF (ARW) is a configuration of the WRF system having ARW as the dynamic solver. WRF has been used by a number of researchers in the past to track cyclones (Islam et al., 2015; Kotal & Bhattacharya, 2020; Nadimpalli et al., 2020; Mohanty et al., 2021). Islam et al. (2015) used WRF to track the cyclone 'Haiyan' recorded over West Pacific Ocean. They managed to forecast the track of the cyclone prior to 40 hours with a track error less than 100 km. Although they were successful in forecasting the track of the cyclone, the intensity of the cyclone was underestimated. Nadimpalli et al. (2020) carried out an assessment of WRF and hurricane weather research and forecasting (HWRF) by forecasting tropical cyclones in the Bay of Bengal. They have shown that the results obtained using WRF and HWRF are comparable up to 36 hours. For longer forecasts beyond 36 hours HWRF has given better results. Hon (2020) has forecasted 13 North Indian Ocean (NIO) cyclones. Mean positional errors for those forecasts are 38 km, 69 km and 107 km for 0, 24 and 48 hour forecast lead times respectively.

The forecast error of rainfall depth due to a cyclone was subjected to experiment using ARW by Osuri *et al.* (2020). They reported higher rainfall error with the long-range (48 - 96 hour) forecast than the short-range (24 hour). The forecast of wind field displacement and speed of the cyclone 'Vardah' formed over the Bay of

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This article is published under the Creative Commons CC-BY-ND License (http://creativecommons.org/licenses/by-nd/4.0/). This license permits use, distribution and reproduction, commercial and non-commercial, provided that the original work is properly cited and is not changed in anyway. Bengal is improved by applying the following two methods. The two methods are relocation of the NCEP GFS forecast wind field and modification of the wind speed at each grid point over the cyclone influenced area (Kotal & Bhattacharya, 2020). Severe cyclonic storm 'Fani' over the Bay of Bengal was forecasted using WRF and HWRF by Mohanty *et al.* (2021). They concluded that the landfall time and position of the cyclone can be forecasted well with a 60 hour lead time. The tropical cyclone tracking and verification techniques used in weather prediction models by the UK Met Office is described elsewhere (Heming, 2017).

The cyclonic storm 'Burevi' was recorded in the Bay of Bengal in December 2020 (Farzan, 2020b). It is the fifth cyclone recorded in the North Indian (NI) basin during year 2020 (Knapp et al., 2018). Three of those five are recorded in the Bay of Bengal subbasin. Those cyclones were named 'Amphan,' 'Nivar,' and 'Burevi' (Knapp et al., 2018). It is worth noting that all those three cyclones have affected Sri Lanka, and cyclone 'Burevi' made a landfall in the east coast (Farzan, 2020a). Figures 1a and 1b are meteorological satellite observation images of cyclone 'Burevi' recorded at 0600 Coordinated Universal Time (UTC) on 02 December 2020. Those two images consist of Day Convective Storms RGB and Day Microphysics RGB, issued by the Japan Meteorological Agency (JMA) using their satellites. The yellow areas shown in Figure 1a are the cumulonimbus clouds with strong updrafts. Those clouds produce severe wind, tornadoes and heavy rainfall (Cotton et al., 2011). In Figure 1b the red and brown colour area shows the thick clouds of cyclone 'Burevi' with deep precipitation.



Figure 1: Satellite images captured at 0600 UTC on 02 December 2020

According to the report issued by the Disaster Management Center (DMC) of Sri Lanka there were three deaths due to cyclone 'Burevi' (Priyantha, 2020). The number of people affected in the Northern and Eastern provinces due to the cyclone are 102,098 and 30,830 respectively. In those two provinces, 103 houses were fully damaged and 3,598 were damaged partially. There were 1,054 small and medium scale enterprises were also damaged due to the cyclone (Priyantha, 2020). Prediction of the track and the intensity of the cyclone is important for the disaster management teams to take early mitigation measures to minimize the damage.

The main objective of this work is to forecast the track and the intensity of cyclone 'Burevi' that affected Sri Lanka, using WRF. Another objective is to identify the best possible forecast lead time in forecasting the track and the intensity of the cyclone. The final objective of this work is to identify the forecasted landfall position and time of the cyclone.

MATERIALS AND METHODS

In this study the track and the intensity of the cyclone "Burevi" is forecasted using WRF. WRF consist of two dynamical cores namely 'ARW' and 'NMM'. In this study the dynamical core 'ARW' was selected. The lateral boundary condition of the selected domain was set by the National Centers for Environmental Prediction (NCEP) Global Forecast System (GFS) data with a horizontal resolution of 0.25° x 0.25° at 3 h time intervals. The dataset was released four times a day in real-time and in this study 16 such datasets were used having 12 h time intervals, starting from 0000 UTC 27 November 2020 to 1200 UTC 04 December 2020. The domain was selected to cover the entire Bay of Bengal in the NI basin. It was centred over the point 12°N, 83°E having 240 x 180 grid points with 15 km grid spacing, covering a 3600 km x 2700 km area. The terrain height of the selected domain is shown in Figure 2. In the model configuration the number of vertical levels were set to 33. Based on the literature the pressure at the top of the model was set to 5000 Pa for all the forecasts (Baki et al., 2022). However with the support of the dataset used to set the lateral boundary condition, it is possible to use a lower model top pressure (Hon, 2020). Adaptive time step was used to automatically set the time step to a maximum value while keeping the model numerically stable (Hutchinson, 2007; Skamarock et al., 2019).

The physics options used in this study were WRF Single-moment 6-class Scheme (WSM6) for microphysics (Hong & Lim, 2006), New Tiedtke Scheme (Zhang & Wang, 2017) for the cumulus parameterization option, Dudhia shortwave Scheme (Dudhia, 1989) for the shortwave radiation, Rapid and Accurate Radiative Transfer Model (RRTM) Longwave Scheme (Mlawer *et al.*, 1997) for the longwave radiation and University of Washington Boundary Layer Scheme (Bretherton & Park, 2009) for Planetary Boundary Layer (PBL). Added to those options MM5 Similarity Scheme (Beljaars, 1995) is used as the Surface Layer option and the Unified Noah Land Surface Model (Tewari *et al.*, 2004) is used as the Land Surface Option.



Figure 2: Terrain Height of the selected domain



Figure 3: Absolute Vorticity at 0600 UTC 02 December 2020

The forecasting of the cyclone 'Burevi' is performed as 16 consecutive model runs. Each forecast is initiated a with different model initialization, from 0000 UTC 27 November 2020 to 1200 UTC 04 December 2020, with

a 12 h time interval. The initialization times and the forecast durations are shown in Table 1. The forecast durations vary from 30 to 120 h. The forecast positions of the cyclone are identified in the literature using the forecasted minimum sea level pressure (MSLP) and relative vorticity (RV). The locations of maximum RV at 850 hPa and MSLP are close in a well developed cyclone over the open ocean (Heming, 2017). In this research the local MSLP locations are identified as possible positions of the cyclone. The absolute vorticity (AV) at 850 hPa is considered instead of RV for a search radius of 2° having local MSLP locations as the centres. The combination of local MSLP and the AV at 850 hPa is considered in selecting the forecasted position of the cyclone. The forecasted AV at 0600 UTC on 02 December 2020 is shown in Figure 3. All the forecasts conducted in this study are evaluated using the International Best Track Archive for Climate Stewardship (IBTrACS) (Knapp et al., 2018). In that dataset the track records for the cyclone 'Burevi' are available from 1200 UTC 01 December 2020 to 1800 UTC 05 December 2020, and it is shown in Figure 4.



Figure 4: The track of cyclone "Burevi" (Knapp et al., 2018)

The forecast leads vary from 0 to 120 h with a 3 h time interval. Since the track record of the cyclone 'Burevi' provided by IBTrACS is from 1200 UTC 01 December 2020 to 1800 UTC 05 December 2020 the number of verifiable forecasting positions for different forecast leads are different. The root mean square error (RMSE) is calculated to evaluate the track of the cyclone 'Burevi' using IBTrACS dataset as a reference. The calculation is done according to the Equation 1. In that equation 'simlat_n' and 'simlon_n' are simulated latitude and longitude coordinates of the cyclone. The values for 'btlat_n' and 'btlon_n' are the latitude and longitude of the cyclone track recorded in the IBTrACS dataset.

No	Initialization time	Forecast duration (h)	Verifiable positions	Track RMSE (km)
1	00 UTC 27 November 2020	120	5	340.1
2	12 UTC 27 November 2020	120	9	293.9
3	00 UTC 28 November 2020	120	13	127.0
4	12 UTC 28 November 2020	120	17	166.7
5	00 UTC 29 November 2020	120	21	59.0
6	12 UTC 29 November 2020	120	25	164.9
7	00 UTC 30 November 2020	120	29	215.7
8	12 UTC 30 November 2020	120	33	202.8
9	00 UTC 01 December 2020	114	35	146.4
10	12 UTC 01 December 2020	102	35	90.5
11	00 UTC 02 December 2020	90	31	245.1
12	12 UTC 02 December 2020	78	27	84.7
13	00 UTC 03 December 2020	66	23	195.7
14	12 UTC 03 December 2020	54	19	72.3
15	00 UTC 04 December 2020	42	15	87.9
16	12 UTC 04 December 2020	30	11	88.3

Table 1: RMSE of the selected forecasts with different initialization times

The great-circle distance between those two points was calculated using the 'haversine' formula (Sinnott, 1984). N is the number of verifiable forecasting positions for each forecast as shown in Table 1.

$$RMSE = \sqrt{\frac{\sum_{n=1}^{N} (haversine(simlat_n, btlat_n, simlon_n, btlon_n))^2}{N}}$$
...(1)

The verification of the tracks for different forecast leads is done using direct positional error (DPE), along-track error (ATE), and cross-track error (CTE) as mentioned by Heming (2017). The DPE is the greatcircle distance between the observed and forecast positions considered at the same time instant in UTC. The DPE does not provide any information about speed and the direction of the forecast (Heming, 2017). The ATE was calculated by extrapolating the line joining observed position and the position observed 3 h earlier and drawing a perpendicular line from the forecasted point to it. The distance between the point of intersection of those two lines and the observed point is considered as the ATE. If the ATE is positive, the forecast is ahead of the observed position in the observed direction of motion having a faster movement, and if the ATE is negative, it is having a slower one. The CTE is the perpendicular distance from forecasted point to the extrapolated line joining observed position and the position observed 3 h

earlier (Heming, 2017). Since the ATE and CTE require one observed position recorded 3 h prior to the observed position, it is not possible to calculate the ATE and CTE for the first observed position. In each of the DPE, ATE, and CTE calculations, the 'haversine' formula is used to calculate the great-circle distance.

The forecasted intensity of the cyclone was evaluated by comparing the sea level pressure and 10-m wind speed obtained in the positions of the forecasted track with the corresponding pressure and maximum sustained wind speed values recorded in IBTrACS dataset (Knapp *et al.*, 2018). The forecasted positions have the MSLP of the considered domain. The forecasted position of the cyclone 'Burevi' at 0600 UTC on 02 December 2020, having an MSLP of 1001.27 hPa, was rounded to 1001 hPa and marked in Figure 5. The absolute MSLP error was calculated by using such forecasted MSLP vales and using the pressure values recorded in the IBTrACS dataset as a reference.

The wind speed forecasted at 0600 UTC 02 December 2020 of the cyclone 'Burevi' is shown in Figure 6. A search radius of 2° is considered form the point of MSLP to obtain the maximum forecasted 10-m wind speed as used by Heming (2017). The absolute wind speed error of the forecasted maximum wind speed was calculated by using the maximum sustained wind speed of IBTrACS dataset (Knapp *et al.*, 2018) as a reference.





Figure 5: Forecasted sea level pressure at 0600 UTC on 02 December 2020.

The cyclone 'Burevi' produced three landfall positions. Those positions were calculated based on the observed cyclone positions recorded in IBTrACS dataset (Knapp et al., 2018). The latitude and longitude coordinates of such points before and after the landfall are used to find the equation of the straight line joining those two. Once the coefficients of the equation are obtained, it is possible to find all the latitude and longitude coordinates between the two points. The terrain elevation of all those points were obtained using the calculated latitude and longitude coordinates. The difference in terrain elevation between a location on sea and a location on land is used to identify the latitude and longitude coordinate of the observed landfall location. The same process described above was applied to the forecasted cyclone positions to obtain the forecasted landfall location. Landfall error was obtained by calculating the great-circle distance between the observed and forecasted landfall positions using the 'haversine' formula. Landfall time was calculated by assuming that the cyclone had moved at a constant speed between the two cyclone positions considered for a period of 3 h. It was calculated according to the distance ratio between the cyclone positions and the landfall location. Landfall time error was calculated by getting the time difference between the forecasted landfall time and the observed one.

Figure 6: Forecasted 10-m wind speed at 0600 UTC on 02 December 2020.

RESULTS AND DISCUSSION

The track RMSE of the selected forecasts with 16 different model initializations are shown in Table 1. From the model initialization time 0000 UTC on 27 November 2020 to 0000 UTC on 29 November 2020, the track RMSE was reduced from 340.1 km to 59.0 km. For the model initialized at 1200 UTC on 01 December 2020, the track RMSE was 90.5 km.

In Figure 7 the forecasts of up to 72 hours are plotted in blue colour and the reference provided by IBTrACS dataset (Knapp et al., 2018) is plotted in red colour. The cyclone started at 1200 UTC 01 December 2020 from the location 7.80°N, 84.80°E and moved towards Sri Lanka according to the IBTrACS dataset (Knapp et al., 2018). It is worth noting that all those forecasts shown in Figure 7 are initiated before the starting time of the cyclone. The observed track of the entire cyclone is plotted in red from Figure 7a to Figure 7d as a reference. According to the observed track of the cyclone, it had got trapped between Sri Lanka and India and was weakened at the end (Knapp et al., 2018). It had also made its second and third landfall on Sri Lanka during that final stage. That event is not forecasted and the cyclone points have moved away from Sri Lanka at the final stage as shown in Figure 7.



Figure 7: Forecast tracks up to 72 hour duration with different model initialization times. Observed and forecasted tracks are plotted in red and blue colours, respectively.

The DPEs of the forecasted tracks for different forecast leads are verified using the IBTrACS data (Knapp et al., 2018). The mean DPEs for different forecasting times up to 72 hours are shown in Table 2. The mean DPE is less than 61 km up to a 24 hour lead time. It is less than 106 km and 131 km for lead times up to 48 and 72 hours, respectively. These results are comparable with the results obtained by Hon (2020) for their forecasts of 13 cyclones in the NIO. The ATE is calculated for different forecast leads using IBTrACS data (Knapp et al., 2018) as observed positions. The ATE gets positive and negative values depending on the forecast position. The absolute value of ATE is considered to calculate the mean ATE, and mean absolute ATE records up to 72 hours are shown in Table 2. The mean absolute ATE is less than 48 km up to a 24 hour forecast lead time. It is less than 84 km and 122 km for lead times up to 48 and 72 hours respectively.

The CTE is calculated for the forecast leads based on the observed positions recorded in IBTrACS data (Knapp *et al.*, 2018). The CTE is considered positive if the forecast position of the cyclone is recorded to the right (left) of the extrapolated line joining observed position and the position observed 3 hour earlier in the northern (southern) hemisphere. The absolute value of

CTE is considered to calculate the mean CTE, and mean absolute CTE records up to 72 hours are shown in Table 2. Mean absolute CTE is less than 42 km up to a 24 hour forecast lead time. It is less than 47 km and 64 km for lead times up to 48 and 72 hours respectively. The mean DPE, mean absolute ATE, and mean absolute CTE are further illustrated in Figure 8. According to that figure the mean DPE, mean absolute ATE, and mean absolute CTE shows higher values for 0 and 3 hour forecast lead times compared to a 6 hour forecast lead time. The main reason is that, during that period, the model is in the spin-up phase to develop the waves and the clouds (Skamarock et al., 2019). It is possible to provide a pre-forecast period to the model, which will provide the required time to adjust to the topography, and produce cloud fields (Skamarock et al., 2019). Such a pre-forecast period is not configured in this model setup and will be considered in future work.

The use of AV with MSLP in selecting the forecasted positions of the cyclone has reduced DPE, ATE, and CTE compared to using only MSLP. When the forecasted MSLP is considered for the entire domain, there exist some MSLP points that do not belong to the cyclone. Those low pressure points may belong to another

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cyclone that is forecasted in the same simulation domain. If MSLP is the only parameter considered in selecting the cyclone position it would provide an erroneous point resulting in larger DPE, ATE, and CTE. Considering AV with local MSLP points avoids such erroneous points that do not belong to the forecasted track of the cyclone. It is possible to extend this method to forecast the tracks of multiple cyclones of the same simulation domain by using the MSLP and AV.

MSLP error is calculated based on the forecasted sea level pressure along the cyclone positions and the pressure values recorded in IBTrACS data (Knapp *et al.*, 2018).

The mean absolute MSLP error up to a 72 hour lead time is shown in Table 2. The mean absolute MSLP error is less than 2.93 hPa up to a 72 hour forecast lead time. The forecasted MSLP is shown in Figure 9 for the corresponding observed MSLP values of cyclone 'Burevi' recorded in IBTrACS data (Knapp *et al.*, 2018). The correlation coefficient was calculated using linear regression based on the observed and forecasted MSLP. The MSLP forecast has a bias of 1.18, an RMSE of 3.07 hPa, and a correlation coefficient of +1.00 compared to the observations. In Figure 9 it is evident that the forecasted MSLP is increased based on the observed MSLP values, with a positive correlation coefficient of 1.00.

 Table 2:
 Mean DPE, mean absolute ATE, mean absolute CTE, mean absolute MSLP error and mean absolute wind speed error for different forecasting lead times.

Forecast lead	Mean DPE	Mean absolute	Mean absolute	Mean absolute	Mean absolute wind
time (h)	(km)	ATE (km)	CTE (km)	MSLP error (hPa)	speed error (ms ⁻¹)
T + 0	58.2	45.4	23.8	1.15	2.71
T + 3	60.2	39.1	41.8	2.03	2.43
T + 6	46.4	27.9	36.1	2.14	1.73
T + 9	47.9	36.8	28.6	1.94	2.76
T + 12	50.4	38.7	25.6	2.85	3.02
T + 15	54.4	47.3	21.4	2.93	3.45
T + 18	54.9	39.6	27.8	2.25	3.27
T + 21	53.0	34.9	32.5	2.39	3.81
T + 24	50.2	38.6	24.5	2.54	3.85
T + 27	53.5	42.4	23.3	2.53	3.32
T + 30	68.0	53.5	33.2	2.11	3.09
T + 33	48.4	40.6	16.5	1.77	3.54
T + 36	62.1	44.0	24.7	1.85	3.92
T + 39	62.4	49.2	29.1	2.36	3.35
T + 42	83.3	60.8	36.8	2.00	3.57
T + 45	90.2	71.1	46.2	1.86	3.50
T + 48	105.7	83.1	42.9	2.20	4.13
T + 51	124.9	117.7	27.4	2.89	4.30
T + 54	130.9	121.7	33.4	2.17	3.65
T + 57	129.8	106.3	57.2	2.27	4.49
T + 60	104.1	92.4	49.6	2.58	4.51
T + 63	125.9	110.8	42.2	2.88	4.24
T + 66	124.2	92.9	63.1	1.89	3.79
T + 69	104.3	82.8	53.5	1.91	4.50
T + 72	128.8	112.9	53.2	2.24	4.55



Figure 8: Mean DPE, mean absolute ATE and mean absolute CTE for different forecast lead times of the cyclone 'Burevi'.



Figure 10: Scatter diagram of maximum forecasted 10-m wind speed, line indicates the straight line fit to the data

The forecast error of maximum wind is calculated based on the 10-m wind speed and the maximum sustained wind speed recorded in IBTrACS data (Knapp *et al.*, 2018). The mean absolute wind speed error up to a 72 hour lead time is shown in Table 2. The mean absolute wind speed error is less than 4.55 ms⁻¹ for all the forecasts. The forecasted 10-m wind speed is shown in Figure 10 for the corresponding observed MSLP values of cyclone 'Burevi' recorded in IBTrACS data (Knapp *et al.*, 2018). The correlation coefficient was calculated using linear regression based on the observed and forecasted 10-m wind speed. The 10-m wind speed forecast has a bias of -2.92, an RMSE of 3.96 ms^{-1} and a correlation coefficient of +0.73 compared to the observations. In Figure 10 it is evident that the forecasted 10-m wind speed is increased



Figure 9: Scatter diagram of forecasted MSLP, line indicates the straight line fit to the data

based on the observed wind speed values with a positive correlation coefficient of 0.73.

The first landfall of the cyclone 'Burevi' was observed at the 9.35°N, 80.74°E latitude and longitude coordinate at 0041 UTC on 03 December 2020. The method used to calculate the observed landfall location and time is described in Section 2. The observed cyclone positions in the IBTrACS dataset (Knapp et al., 2018) are used in that calculation. The same process is used to calculate the forecasted landfall location and time. The landfall location and time error for the first landfall were calculated and the results are shown in Table 3. The forecasted minimum landfall position error is recorded as 18.1 km with a lead time of 48 hours. The minimum forecasted landfall time error of the first landfall of the cyclone is 4 minutes with a lead time of 36 hours. Obtaining such information regarding the landfall location and time of the cyclone is an advantage when it comes to disaster management.

The intensity of the cyclone was reduced after the first landfall. This is clearly visible from the data recorded by Knapp *et al.* (2018), where the maximum sustained wind speed was 20 ms⁻¹ at the first landfall, 14 ms⁻¹ at the second landfall, and 10 ms⁻¹ at the third landfall. The second landfall was observed at 9.08°N, 80.02°E at 0300 UTC on 05 December 2020. In all the forecasts conducted in this study the second landfall was only forecasted with a landfall error of 62 km in a forecast initialized at 0000 UTC on 03 December 2020. In that forecast, the landfall time error is minus 40 hours and 30 minutes. The third landfall was observed at 8.9°N, 79.9°E at 1800 UTC on 05 December 2020. This landfall is not

recorded in any of the forecasts conducted in this study. The intensity of the cyclone was reduced when it reached its final stage. This is the reason for not forecasting the third landfall accurately.

Initialization time	Lead time (h)	Landfall position	Landfall time
		error (km)	Error
00 UTC 29 November 2020	102	35.0	6 h 49 m
12 UTC 29 November 2020	96	21.0	11 h 49 m
00 UTC 30 November 2020	63	51.0	-5h 11m
12 UTC 30 November 2020	48	18.1	- 11 h 33 m
00 UTC 01 December 2020	42	27.8	4 h 26 m
12 UTC 01 December 2020	36	48.8	4 m
00 UTC 02 December 2020	15	45.0	-7h 49m
12 UTC 02 December 2020	0	58.8	-9h 56m

 Table 3:
 Landfall position and time error of the first landfall

CONCLUSIONS

The cyclone considered in this study has affected Sri Lanka by having its first landfall at 0041 UTC on 03 December 2020. Its track and intensity were forecasted with a maximum forecast duration of 120 hours using WRF. The track RMSE was reduced from 340.1 km to 59.0 km for model initializations from 0000 UTC on 27 November 2020 to 0000 UTC on 29 November 2020. This clearly shows the increase of accuracy of the forecasted track when the forecast lead time is reduced. For forecast lead times up to 24 hours the DPE, ATE, and CTE are less than 61 km, 48 km, and 42 km, respectively. When it comes to a forecast lead time up to 48 hours the DPE, ATE, and CTE are less than 106 km, 84 km, and 47 km, respectively. DPE, ATE, and CTE are higher for 0 and 3 hour forecast lead times compared to a 6 hour forecast lead time. The authors believe that this is due to the model spin-up phase. Providing the model with a preforecast period will be considered in future to minimize the effect of the spin-up phase. The forecasted mean absolute MSLP is less than 2.93 hPa up to a 72 hour forecast lead time. The forecasted MSLP has a positive correlation coefficient of 1.00 with its corresponding observations. The mean absolute wind speed error is less than 4.55 ms⁻¹ for all the forecasts. The forecasted wind speed has a positive correlation coefficient of 0.73 with its corresponding observations. With all the above mentioned results, it is possible to conclude that WRF is capable of forecasting the track and the intensity of a cyclone that affects Sri Lanka with a forecast lead time up to 48 hours.

The first landfall of the cyclone 'Burevi' was forecasted with a minimum landfall position error of 18.1 km with a forecast lead time of 48 hours. The minimum forecasted landfall time error of the cyclone is 4 minutes with a forecast lead time of 36 hours. Obtaining such information is an advantage when it comes to disaster management. The intensity of the cyclone was reduced after the first landfall. The second landfall was forecasted by only one forecast. That had a landfall error of 62 km and a landfall time error of minus 40 hours and 30 minutes. The third landfall of the cyclone was not recorded in any of the forecasts conducted in this study. In conclusion, the first landfall position forecast of this cyclone is acceptable with the WRF configuration used in this study. The position and time forecasts for the second and third landfall require further investigation to improve their accuracy.

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RESEARCH ARTICLE

Marine Microalgae

Anti-inflammatory activity of nonyl 8-acetoxy-6-methyloctanoate, isolated from the cultured marine diatom, *Phaeodactylum tricornutum* : mediated via suppression of inflammatory mediators in LPS-stimulated RAW 264.7 macrophages

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Abstract: In this study, the anti-inflammatory effects of nonyl 8-acetoxy-6-methyloctanoate (NAMO), isolated from the cultured marine diatom, Phaeodactylum tricornutum Bohlin, against LPS-induced RAW 264.7 macrophages were evaluated. NAMO has indicated the strongest inhibitory effects against nitric oxide (NO) and prostaglandin E₂ (PGE₂) production, dosedependently, on lipopolysaccharide (LPS)-induced RAW 264.7 cells. The 50% NO production inhibitory concentration (IC_{50}) of NAMO was 24.8 µM with the least cytotoxic effect in both LDH and MTT assays. Inflammatory stimulators such as LPS, which induce cytokines in the process of macrophage activation and mediate tissue response in different phases of inflammation, were studied. NAMO showed significant and strong suppression of pro-inflammatory cytokine and interleukin-1ß (IL-1ß) production, but no significant inhibitory effect on the production of cytokines including tumor necrosis factor- α (THF- α) and interleukin-6 (IL-6) at the tested concentrations against LPS treatment on RAW macrophages. Western blot analysis was carried out to determine the protein expressions of inducible nitric oxide synthase (iNOS) and cyclooxygenase-2 (COX-2) proteins which mediate the suppression effect of NAMO on NO and PGE, production. The Western blot assay confirmed the suppression of iNOS and COX-2 protein expressions against LPS-stimulated RAW264.7 cells. Collectively, NAMO isolated from the cultured marine diatom, P. tricornutum exhibited a

profound anti-inflammatory effect *in vitro*, suggesting that the compound might have a beneficial effect during the treatment of inflammatory diseases and can be used in functional food applications.

Keywords: Anti-inflammatory activity, cultured marine diatom, fatty alcohol esters, *Phaeodactylum tricornutum*, RAW 264.7 cells

INTRODUCTION

Marine microalgae, with the associated broad spectrum of secondary metabolites, have stimulated a vast interest due to biochemical and ecological diversity (Borowitzka, 1995). Exploration of new bioactive natural products from marine bio-resources like marine microalgae for pharmacological applications has been described as a promising aspect of this. The interest in the isolation of pharmacologically active metabolites from marine microalgae has increased throughout the world. Microalgae are rich sources of nutrients including a high content of protein, polyunsaturated fatty acids (PUFA), carbohydrates, minerals, vitamins, pigments, and secondary metabolites (Becker, 2007).

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Cultured marine microalgae have been widely used as a food source for aquaculture. Diatoms are abundant components of the phytoplankton, representing major nutritional sources. Phaeodactylum tricornutum, a marine diatom, has been characterized due to containing different forms of PUFAs and lipids, including free sterols, and esterified and glycosylated conjugates (Veron et al., 1996; Yang et al., 2019). A few reports have investigated the quantification of lipids that are available in P. tricornutum. The distribution of lipid constituents in marine microalgae is dependent on the growing conditions, that include different temperatures, light spectral quality, salinity, and nutrients provided in the aquaculture (Samarakoon et al., 2013). The possibility of changing the lipid concentration of microalgae is a very important characteristic that can be used to determine the physiological state of a microalgae population, as a population biomarker (Fabregas et al., 1997).

Inflammation is a physiological process initiated by pathogenic invasion or cell and tissue injury (Wadleigh et al., 2000). Inflammation can be caused by the activation of various immune cells such as macrophages, lymphocytes, and neutrophils. Overproduction of proinflammatory cytokines including interleukin (IL) and tumor necrosis factor (TNF)- α can be stimulated in macrophages as per the inflammatory disease. In addition, inflammatory mediators such as nitric oxide (NO) and prostaglandin (PGE₂) are also activated due to inducible nitric oxide synthase (iNOS) and cyclooxygenase-2 (COX-2) (Lee et al., 2006). Therefore, finding a better insight into anti-inflammatory agents is gaining interest, in order to regulate immune-related diseases, inflammatory disorders, and infections (Jung et al., 2009). Assessing the anti-inflammatory effect of the isolated lipid against LPS-stimulated macrophages from cultured marine microalgae was of great interest. Hence, in this study, we examined whether the cultured marine diatom, P. tricornutum is capable of producing novel compounds with pharmaceutical value. In our previous publication, the anticancer activity of a novel fatty alcohol ester; nonyl 8-acetoxy-6-methyl octanoate (NAMO) isolated from the cultured marine diatom P. tricornutum has been revealed (Samarakoon et al., 2014). In addition, in this study, we evaluate the in vitro anti-inflammatory activity of the isolated novel compound from the hexane fraction of the extract from the cultured marine diatom, P. tricornutum, against LPSstimulated RAW macrophages.

MATERIALS AND METHODS

Chemicals and reagents

Dulbecco's Modified Eagle's Medium (DMEM), foetal bovine serum (FBS), and penicillin-streptomycin were purchased from Gibco/BRL (Burlington, ON, Canada). 3-(4,5-Dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT), and dimethyl sulphoxide (DMSO) were purchased from Sigma–Aldrich (St. Louis, MO, USA). All other chemicals and reagents used in these investigations were of analytical grade.

Culturing of Phaeodactylim tricornutum

The marine diatom *Phaeodactylum tricornutum* Bohlin was kindly provided by the Korea Marine Microalgae Culture Center (KMMCC). The alga was inoculated in 30 L plastic cylinders at 20 °C after pre-cultivation in 5 L glass vessels (medium 4 L), and the air was continuously supplied at 5 L min⁻¹ by air-lift. The light was provided by 60 W fluorescent lamps at an intensity of 2,500 lx (Light : Darkness = 24 : 0). The alga was cultured in Conway medium (Walne, 1966) prepared from filtersterilized seawater, and the cultures were continuously active during the 8-10 ds after inoculation. The cells were flocculated with 200 ppm Al₂ (SO₄)₂ (v/v) (Ilshin biochemical, Magicpool-99) and then recovered with centrifugation at 2,000 rpm using a basket centrifuge (Hanseong Co., Ansan, South Korea). The harvested diatom biomass was frozen at -25 °C and preserved until freeze drying.

Extraction and isolation

The material (38 g) was extracted three times with 80% methanol by sonication for 90 min each time, at 25 °C. After being concentrated by evaporating off the solvent under reduced pressure using a rotary evaporator, the crude methanol extract was further subjected to solvent-solvent partition chromatography. Then, four different fractions with varying polarity, namely, *n*-hexane, chloroform, ethyl acetate, and aqueous extracts were separated. The *n*-hexane fraction (550 mg) was found to be active and subjected to fractionation using a solid-liquid phase chromatography (normal phase silica) column (3 cm \times 22 cm) using hexane and ethyl acetate as an increasing hydrophilic solvent system to

give 14 fractions, labelled F1~F14. Using the thin layer chromatography (TLC) results, visualized by heating after spraying with ethanol (90%) - sulphuric acid (10%) staining system, the desired fraction was identified as eluting with hexane (10%): ethyl acetate (90%), (F7: 19 mg), with enough purity to determine the molecular structure. Therefore, the molecular mass was measured using a high-performance liquid chromatography - diode array detector coupled with an electrospray ionization mass spectrometer (HPLC-DAD-ESI/MS) (Hewlett-Packard, Waldbronn, Germany), and NMR studies using a JEOL JNM-ECX400 spectrometer (¹H-NMR/400 MHz and ¹³C-NMR/100 MHz) used for structure elucidation.

Nonyl 8-acetoxy-6-methyloctanoate (NAMO)

Liquid (CHCL₃): $C_{20}H_{38}O_4$ in the ESI (negative mode) as (M-H) 341.14 m/z. The calculated molecular mass is 342.28 m/z (M). ¹H-NMR (400 MHz, CDCl₃): δ / ppm: 0.87 (3H, t, J = 6.72 Hz, H-9'), 0.96 (3H, s, H-11), 1.25 (2H, m, H-4, 5, 5', 6'), 1.29 (2H, m, H-4', 7'), 1.31 (2H, m, H-8'), 1.43 (2H, m, H-3'), 1.50 (2H, m, H-2'), 1.53 (2H, m, H-7), 1.65 (H, m, H-6), 1.68 (2H, m, H-3), 2.21 (3H, s, H-10), 2.32 (2H, t, J = 7.53 Hz, H-2), 4.08 (2H, dd, J = 7.4,14.6 Hz, H-8), 4.13 (2H, dd, J = -7.4,14.6 Hz, H-1'). ¹³CNMR (100 MHz, CDCl₃): δ 14.3 (q, C-9'), 21.2 (q, C-10), 22.8 (q, C-11), 22.9 (t, C-8'), 24.6 (t, C-3'), 24.7 (t, C-4), 24.9 (t, C-3), 29.2 (t, C-2'), 29.4 (t, C-7), 29.6 (t, C-4'), 29.9 (t, C-5'), 29.9 (t, C-6'), 32.1 (t, C-7'), 32.9 (d, C-6), 34.1 (t, C-2), 37.5 (t, C-5), 60.6 (t, C-1'), 63.3 (t, C-8), 171.3 (s, C-9), 179.2 (s, C-1).

Cell culture

The murine macrophage cell line RAW 264.7 was purchased from the Korean Cell Line Bank (KCLB, Seoul, South Korea). RAW 264.7 cell line was cultured in Dulbecco's modified eagle medium (DMEM) supplemented with 100 U mL⁻¹ of penicillin, 100 μ g mL⁻¹ of streptomycin, and 10% fetal bovine serum (FBS). The cells were incubated and maintained in an atmosphere of 5% CO₂ at 37 °C. The cells were sub-cultured every 2 days and exponential phase cells were used throughout the experiments (Kim *et al.*, 2013).

Determination of nitric oxide (NO) production

RAW 264.7 cells $(1 \times 10^5$ cell mL⁻¹) were placed in a 24-well plate and after 24 h the cells were pre-incubated

with various concentrations of the sample at 37 °C for 1 h. Then further incubation was done for another 24 h with LPS (1 µg mL⁻¹) at the same temperature. After the incubation, the quantity of nitrite accumulated in the culture medium was measured as an indicator of NO production (Lee *et al.*, 2006). Briefly, 100 µL of cell culture medium were mixed with 100 µL of Griess reagent (1% sulfanilamide and 0.1% naphthyl ethylenediamine dihydrochloride in 2.5% phosphoric acid), and the mixture was incubated at room temperature for 10 min, and the optical density at 540 nm was measured using an enzyme-linked immunosorbent assay (ELISA) microplate reader (Sunrise, Tecan Co. Ltd., Australia). The fresh culture medium was used as a blank in every experiment (Kim *et al.*, 2013).

Lactate dehydrogenase (LDH) cytotoxicity assay

RAW 264.7 cells $(1.5 \times 10^5$ cells mL⁻¹) were plated in a 96-well plate and after 16 h the cells were pre-incubated with various concentrations of the sample for 1 h at 37 °C. Then the cells were further incubated for another 24 h with LPS (1 µg mL⁻¹) at the same temperature. After the incubation, the LDH level in the culture medium was determined using an LDH cytotoxicity detection kit (Promega, Madison, WI, USA) according to the manufacturer's instructions. Briefly, 50 µL of the reaction mixture were added to each well, and the incubation was done for 30 min at room temperature in the dark. Then, 50 µL of stop solution were added to each well, and absorbance was measured at 490 nm using a microplate reader (Sunrise, Tecan Co. Ltd., Australia) (Kim *et al.*, 2013).

Cytotoxicity assessment using MTT assay

The cytotoxicity of NAMO against the RAW 264.7 cells was determined using the colorimetric MTT assay. Cells were seeded in a 24-well plate at a concentration of 1×10^5 cells/mL⁻¹. After 24 h, the seeded cells were treated with sample. Then, all of the cells were incubated for an additional 24 h at 37 °C. MTT stock solution (50 µL; 2 mg/mL in PBS) was added to each well to a total reaction volume of 250 µL. After incubating for 3 h, the supernatants were aspirated. The formazan crystals in each well were dissolved in 200 µL of dimethylsulphoxide (DMSO). The resulting absorbance was measured with an ELISA plate reader set at 540 nm (Samarakoon *et al.*, 2014).

Determination of pro-inflammatory cytokines, tumor necrosis factor- α (TNF- α), interleukin-1 β (IL-1 β), and interleukin-6 (IL-6) production.

The inhibitory effect of the sample on the production of pro-inflammatory cytokines from LPS-stimulated RAW 264.7 cells was determined according to a previously described method (Cho *et al.* 2000). Briefly, RAW 264.7 cells $(1 \times 10^5 \text{ cells mL}^{-1})$ were pretreated with the sample for 2 h and then treated with LPS $(1 \mu \text{g mL}^{-1})$ to allow the production of pro-inflammatory cytokines for 24 h. Supernatants were used for the assay using an ELISA kit (R & D Systems, Minneapolis, MN, USA) according to the manufacturer's instructions.

Determination of prostaglandin-E₂ (PGE₂) production

RAW 264.7 cells (1×10^5 cells mL⁻¹) were pretreated with the sample for 2 h and then treated with LPS ($1 \mu g m L^{-1}$) to allow the production of pro-inflammatory cytokines for 24 h. The PGE₂ levels in the culture medium were quantified using a competitive enzyme immunoassay kit (R & D Systems, Minneapolis, MN, USA) according to the manufacturer's instructions. The release of PGE₂ was measured relative to that of the control value (Cho *et al.*, 2000).

Western blot analysis

RAW 264.7 cells $(1 \times 10^5 \text{ cells mL}^{-1})$ were pretreated for 16 h and then treated with LPS (1 μ g mL⁻¹) in the presence or absence of the sample. After incubation for 24 h, the cells were harvested, washed twice with icecold phosphate-buffered saline (PBS), and the cell lysates were prepared with lysis buffer (50 mmol L⁻¹ Tris-HCl (pH 7.4), 150 mmol L⁻¹ NaCl, 1% Triton X-100, 0.1% SDS and 1 mmol L⁻¹ ethylenediaminetetraacetic acid (EDTA)) for 20 min on ice. Cell lysates were centrifuged at 14,000 x g for 20 min at 4°C. Then the protein concentrations in the supernatants were measured using a BCATM protein assay kit. Cell lysates containing 30 µg of protein were subjected to electrophoresis using sodium dodecyl sulphate-polyacrylamide gels (SDS-PAGE) at 12%, and the separated proteins were transferred onto nitrocellulose membranes (Bio-Rad, Hercules, CA, USA). The membranes were incubated with a blocking solution (5% skim milk in Tris-buffered saline containing Tween -20) for 90 min at room temperature. Then the membrane was incubated with anti-mouse iNOS (1:1000; Calbiochem, La Jolla, CA, USA) and anti-mouse COX-2 (1:1000; BD BiosciencPharmingen, San Jose, CA, USA) overnight at room temperature.

After washing, the blots were incubated with horseradish peroxidase-conjugated goat anti-mouse IgG secondary antibody (1: 5000; Amersham Pharmacia Biotech, Little Chalfont, UK) for 90 min at the ambient temperature. Signals were developed by exposure to X-ray films and visualized according to the described procedure, using a Western Lightning-ECL detection kit (PerkinElmer, MA, USA).

Statistical analysis

All the data are expressed as mean \pm standard deviation of three determinations. Statistical comparison was performed via a one-way analysis of variance followed by Duncan's multiple range test (DMRT). P values of less than 0.05 (p < 0.05) were considered as significant.

RESULTS AND DISCUSSION

Inhibitory effect of 8-acetoxy-6-methyloctanoate (NAMO) from P. tricornutum against LPS-induced nitric oxide production and cytotoxicity on RAW macrophages

The isolated compound (Figure 1), NAMO from the hexane fraction of the cultured marine microalga, *P. tricornutum*, was studied to assess the anti-inflammatory activity using LPS-induced RAW macrophages, and the NO production inhibitory and cytotoxic effects were determined. Five different concentrations (0, 5, 12.5, 25, and 50 μ g mL⁻¹) of NAMO were pre-incubated with LPS for 24 h and the significant inhibitory effect on NO production level was observed dose-dependently. The 50% inhibitory concentration (IC₅₀) value of 8.50 μ g mL⁻¹ (24.8 μ M) was determined. Nitric oxide (NO) is a signaling molecule. The inflammatory mediator, which is stimulated by iNOS. The effects of various



Figure 1: Structure elucidation of nonyl 8-acetoxy-6methyloctanoate (NAMO) isolated from *Phaeodactylum tricornutum*

120

100

80

60

40

20

0

120

100

80

40

20

0

Control

0

Cell viability (%) 60

NO production (%)



25

LPS 1 µg/mL + NAMO (µg/mL)

50



12.5

5

concentrations of effect of NAMO isolated from the cultured marine diatom P. tricornutum on NO production in LPS-activated RAW 264.7 cells was evaluated in vitro. LPS stimulates the cells to form NO molecules and causes inflammation. The increase of NO concentration in the medium was dose-dependently suppressed with the incubation of NAMO (Figure 2). Moreover, the cell viability assay indicated that NAMO did not show a cytotoxic effect on RAW 264.7 cells at all the concentrations. The cytotoxic effects on RAW 264.7 cells with the tested concentrations of NAMO were determined by MTT assay, and a 5.0 µg/mL concentration of NAMO gave the highest cell viability, 100%, compared to the negative control (Figure 2B). It is also confirmed that the LDH release assay indicated little or no cytotoxicity on macrophages at the tested concentrations (Figure 2A). Therefore, NAMO is marked as a potential agent for attenuating NO production without cytotoxic effects.

Inhibitory effect of NAMO on LPS-induced PGE_2 production

In this study, NAMO inhibited the LPS-induced PGE₂ production in a dose-dependent manner. PGE, production

was 30% of normal at 25 μ g mL⁻¹ concentration of NAMO, and 100 % without NAMO treatment. Therefore, NAMO might induce anti-inflammatory effects by strongly inhibiting PGE₂ production when incubated with RAW macrophages (Figure 3 A).



Figure 3: Inhibitory effect of the isolated NAMO compound from cultured marine diatom *Phaeodactylum tricornutum* on LPSinduced PGE₂ (A), TNF- α (B), 1L-1 β (C) and 1L-6 (D) production in RAW 264.7 macrophages, after the incubation of cells with LPS for 24 h in the presence or absence of NAMO. Values are expressed as mean ± SD in triplicate experiments. Values with different letters are significantly different at p < 0.05 as analyzed by DMRT.

Inhibitory effect of NAMO on LPS-induced proinflammatory cytokine (TNF- α , IL-1 β , and IL-6) Production

The inhibition of cytokine production or regulation is a key mechanism in the mediation of inflammation. Inflammatory stimulators such as LPS induce cytokines in the process of macrophage activation and mediate tissue response in different phases of inflammation. In this assay, the inhibitory effect of NAMO on the production of pro-inflammatory cytokines such as TNF- α , IL-1 β , and IL-6 in LPS-stimulated RAW 264.7 macrophages was studied. The NAMO did not show significant inhibitory activities on the production of cytokine TNF- α and IL-6 at the tested concentrations (Figure 3B, D). However, the strong suppression of the production of cytokine IL-1 β was observed after the pretreatment of NAMO with RAW macrophages (Figure 3C). Therefore, the inhibition pathway might be mediated by the association of IL-1 β formation.



Figure 4: Inhibitory effect of the isolated NAMO compound from cultured marine diatom *Phaeodactylum tricornutum* on LPS-induced iNOS and COX-2 protein expression in RAW 264.7 macrophages. The cells were incubated with LPS for 24 h in the presence or absence of the NAMO. Then, cell lysates were subjected to electrophoresis and the expression levels of iNOS and COX-2 were detected with specific antibodies.

Inhibitory effect of NAMO on LPS-induced iNOS and COX-2 protein expression

Western blot analysis was carried out to determine the protein expressions of iNOS and COX-2 proteins which mediate the suppression effect of NAMO on NO and PGE₂ production (Figure 4).

According to the results, iNOS and COX-2 protein expressions were markedly induced when the macrophages were treated with LPS compared to the control. However, upon incubation with NAMO followed by LPS treatment, both iNOS, and COX-2 protein expression was downregulated, and at the dose of 25 μ g mL⁻¹, the COX-2 protein expression was completely suppressed. Overall, it was suggested that the strong inhibitory effect on PGE2 production by NAMO was correlated with the down-regulation pattern of COX-2 protein expression. In addition, NAMO inhibited the production of NO levels by suppressing iNOS proteins.

P. tricornutum is a well-known marine diatom that accumulates fatty acids with varying degrees of unsaturation. In particular, the compositions of the sterols, PUFA, and fatty acids which are associated with *P. tricornutum* biomasses as lipids have different functional values that are of significant importance for the advancement of marine biotechnology (Radakovits *et al.*, 2011). The cultivation of marine microalgae in an aquaculture system has been practised for a few decades. Open or closed (photobioreactor) microalgae culture systems are applicable based on the energy source and type of culture. Each one of the culturing systems has either desirable or undesirable effects, since

the photosynthetic organisms and their mass production may directly correlate with the source of energy (Hong *et al.*, 2012). The controlled optimum conditions with operational inputs such as salt, dissolved CO_2 , water, nutrients, pH, and O_2 provides a great opportunity for a steady environment without being contaminated by photo-bioreactors. Researchers and technologists have been attracted to cultured marine microalgae to disclose the biochemical constituents of the crude extracts and to find out which components have pharmacological effects (Plaza *et al.*, 2009). Based on the major ingredients available in cultured marine microalgae, it is a promising source for the functional food industry (Garcia *et al.*, 2006; Samarakoon & Jeon, 2012;).

The results of our previous work, culturing and harvesting the biomass of the marine diatom, P. tricornutum, were discussed (Samarakoon et al., 2013). In addition, biological activity-guided fractionation led to the isolation of five different compounds from the hexane fraction of cultured P. tricornutum. The isolated novel fatty alcohol ester; NAMO was evaluated for inhibiting the growth of human leukemia cancer cells significantly through the p53 and caspase-3 mediated cell apoptotic pathway (Samarakoon et al., 2014). In the present study, the same isolated compound from P. tricornutum was evaluated for anti-inflammatory effects in vitro. Nitric oxide (NO) acts as an intercellular and intracellular signal molecule and plays crucial roles in many cellular functions in the nervous, cardiac, vascular, and immune systems (Wichard & Pohnert, 2006). However, the low levels of NO production are important in protecting organs such as the liver from ischaemic damage but the chronic expression of NO is associated with various inflammatory conditions (Kassim et al., 2010). Over the pathological conditions, NO production is increased by iNOS (Kim et al., 1999; Sahu et al., 2013). In terms of the potential therapeutic value when related to inflammation, inhibition of NO production by suppression of iNOS expression can be considered. A few recent publications have confirmed that the organic solvent extractions and compounds isolated from plant-based materials attenuated the inflammations caused by LPS-induced RAW macrophages via key signaling pathways such as the NF-kB (Ahujaa et al., 2019; Hao et al., 2019), and the iNOS pathway (Bharadwaj et al., 2020). The ethanol extract of a new microalgal strain, Aurantiochytrium limacine showed the significant suppression of genes for pro-inflammatory cytokines mediated by the NFkB pathway (Takahashi et al., 2019). With the profound inhibitory effect on NO production evidenced by NAMO, it was further evaluated for its ability to inhibit the LPSinduced PGE2 production in RAW264.7 macrophages.

The present findings indicated that NAMO has antiinflammatory properties, and the effects might be caused by inhibiting iNOS, COX-2, TNF- α , IL-1 β , and IL-6 expression via affecting the NF- κ B pathway. The accessible nature of the cultured marine microalgae is useful not only for the development of pharmacological prospects, but also to meet an increasing demand for nutraceuticals and food supplements. Exploring the sustainable use of cultured marine diatoms is considered important in future industrial and viable applications, and for bioprospecting for the novel bioactive metabolites. Developing new and safer treatment strategies using biomaterials extracted from marine microalgae to prevent and treat inflammatory diseases could be possible (Morais et al., 2015). Therefore, we have evidence that the isolated NAMO from the cultured marine diatom P. tricornutum is a promising source for both in vitro anticancer and anti-inflammatory effects. The findings of the study strongly emphasize the clinical and biotechnological potential of the isolated novel fatty alcohol ester from the cultured marine diatom P. tricornutum for further developments.

CONCLUSION

NAMO, isolated from the cultured marine diatom P. tricornutum, was identified as an anti-inflammatory active compound and showed promising inhibitory effects against cytokine production or regulation, which is a key mechanism in the mediation of inflammation against 264.7 RAW macrophages. NAMO demonstrated an inhibitory effect against NO with the 50% inhibitory concentration (IC₅₀) value of 8.50 μ g mL⁻¹ (24.8 μ M). In addition, NAMO attenuated PGE, and 1L-1ß production significantly in LPS-induced RAW cells. Moreover, the protein expression of iNOS and COX-2 confirmed the down-regulatory effect in a dose-dependent manner against RAW macrophages. Hence, the isolated NAMO can be a novel alternative with anti-inflammatory efficacy for future lead compounds in pharmacological applications.

Data availability

The data used to support the findings of this study are available from the corresponding author upon request.

Conflicts of interest

The author(s) declare(s) that there are no conflicts of interest regarding the publication of this paper.

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RESEARCH ARTICLE

Mechanical Engineering

Implementation of a fuzzy logic approach for a smart production system

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Abstract: This paper considers the monitoring of the entire process of an installation of a smart autonomous production system, and continuous analysis of its operation. The smart production system consists of an industrial robot and a computer numerical control (CNC) machine. Research is focused on the industrial robot subsystem; an implementation of a fuzzy logic method for end effectors is applied, and results regarding relevant clamping force are presented. The grippers and magnetic switches are described in more detail, and devices for measuring the clamping force are shown and analyzed. After installation and levelling of the machine, it is necessary to perform calibration - the correction of movement along the axis of the CNC machine. In theory, if the CNC machine is programmed to follow an ideal circular path, when the positioning accuracy of the machine and its tooling is perfect, then the actual curve that the machine produces, would correspond exactly to the ideally programmed circle. In practice, many factors affect CNC machine performance causing deviations from the ideal circuit. Factors influencing the accuracy are various, stemming from machine geometry, through the control system, to machine wear over many years of the operation. A possible upgrade of the existing system, in order to increase the autonomy, and reduce the need for a human operator, would include the installation of a 3D camerasensor. The robot would take the correct pieces in accordance with production or assembly process, from the box next to it. The box is being filled with preparations from a conveyor belt.

Keywords: Artificial intelligence, automation, fuzzy logic, industrial computer systems, Industry 4.0.

INTRODUCTION

Complex working conditions and the size of machines are the key challenges facing modern industrial plants today (Mikić *et al.*, 2022). Nowadays industrial production companies strive to make the transition from automation to digitization, and reach the requirements and exploit opportunities of Industry 4.0. Due to the necessity for modernization of the companies, reduction of production costs, increase of productivity, quality of service, improvement of working conditions, timely supply of material and parts, as well as finished products, it is crucial to develop methods for the achievement of these goals, and apply these appropriately in the real work environment (Günter *et al.*, 2006; Novaković *et al.*, 2021).

The aim of this paper is to examine and evaluate the possibility of successful installation of a smart production system of the latest generation, based on fuzzy logic implemented in the decision making, and to compare the results with the production systems of the previous generations. The research includes all the necessary conditions for the successful implementation of a flexible production cell in an automated production line in a modern factory. All conditions for the implementation of the Industry 4.0 are fulfilled, with the use of robots and

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intelligent control, as well as trends in their development and application.

A concrete example of the application of an intelligent autonomous system in the production of machine elements is explained. The HAAS CNC machine is chosen for an experiment (HAAS, USA) and the intelligent production is described in detail, with the flexible cell, robot and smart control.

Choosing the appropriate grippers, with respect to the grasping force, is presented and it is implemented by using an artificial intelligence technique – the SWARA method (Step-wise weight assessment ratio analysis).

In order to perform fusion of the computer numerical control (CNC) machine and industrial robot, it is necessary to:

- Install an experimental setup, consisting of a CNC machine, with work-piece plate - tilting table with two axes, the industrial robot as an active part of the clamping system and work-piece;
- Connect the devices into a network so that all can communicate with each other;
- Create a numeric control program for the HAAS CNC machine in computer aided manufacturing (CAM) software;
 Program a robot and synchronize it with the CNC

machine program;

 Test whether pieces produced by the CNC machine are within the given tolerances.

The basic challenge is to connect these subsystems and ensure their mutual M2M (machine to machine) communication in order to determine whether the CNC cell system works well as a whole. There are many disturbances the robot is subjected to while being in operation – the weight it lifts can vary, as well as the dimensions of the work-piece, the tools are changing too, and the travel path is different from one task to another (Owen *et al.*, 2006; Chen & Dong, 2013; Cen & Melkote, 2017; Gualtieri *et al.*, 2021).

After the service department of the company had completed the installation of the components on the customer's premises, it is necessary to perform leveling and calibration of the machine and the robot. The leveling of the CNC machine and the robot is performed with the help of a special digital measuring device. This measuring device has the ability to communicate to mobile devices and computers over the Bluetooth connection. Information on the position of the work-piece, plate, or some other reference part of the machine is sent in real time to the mobile phone of the service technician who performs the adjustment (Dumas *et al.*, 2011; Tyapin *et al.*, 2014). That way, the service technician does not need to go to the device after checking each foot rate. To add to that, there is no need for another service technician who would constantly read the data during levelling (Lehmann *et al.*, 2012).

CRITERIA

The end effectors are the crucial part of the robot subsystem, which is the focus of this project, and the appropriate choice is obtained by using a fuzzy logic approach (Pérez Armayor & Díaz Batista, 2010; Dihovični & Škrbić, 2020).

In robotics, the end effectors are devices at the end of a robotic arm designed to work with the environment. The exact nature of this device is determined by the application of the robot (Wan *et al.*, 2015; Wu *et al.*, 2015).

By strict definition, which comes from serial robotic manipulators, the end effectors represent the last connection (or end) of the robot. In a broader sense, the end effectors can be viewed as part of a robot that communicates with the work environment (Owen *et al.*, 2006).

End effectors can consist of a gripper, or a tool. In general, there are four categories of robotic gripping (Wan *et al.*, 2015):

- Impactive jaws, or claws that directly physically grip an object;
- Ingressive needles, or buttons that physically penetrate the surface of an object (used to handle textiles, carbon fibers, and glass fibers);
- Astrictive attractive forces applied to the surface of an object (vacuum, magnetic, or electro adhesion);
- Contiguous requires direct contact (such as gluing, surface tension, or freezing).

Most often, a mechanical gripper can have two, three, or even five grippers (fingers). In general, the gripping mechanism is operated by grippers, or mechanical fingers. Two-finger clamping is commonly used for industrial robots that perform specific tasks in less complex applications (Martinov *et al.*, 2012).

The mechanism used in gripping with two-finger grips is characterized by:

- 1) the shape of the surface to be gripped;
- 2) the force required to grip an object.

The shape of the grips can be selected according to the shape of the object being manipulated. For example, if the robot is designed to lift a round object, the shape of the gripper may have a concave contour.

Although there are numerous forces acting on a body raised by a robotic arm, the main force is the force of friction. The surface of the grip can be made of soft material with a high coefficient of friction, so that the surface of the object is not damaged. The robotic gripper must withstand not only the weight of the object, but also the acceleration caused by the frequent displacement of the object.

To determine the force required to grasp an object, the following formula is used:

$$F = \frac{m(a+g)}{\mu n},\tag{1}$$

where: *F* is a force required to grasp an object; *m* is a mass of the grasped object; *a* is an acceleration of the object;

 μ is a coefficient of friction;

n is a number of grippers;

g is an acceleration of gravity.

The clamping force, in addition to acceleration, is also affected by the length of the grippers.

The diagram in Figure 1 displays how the clamping force is changing with respect to the length of the grippers.



Figure 1: The diagram of the gripping force (F) with respect to the length of the grippers (L) (source: SCHUNK GmbH & Co. KG)

The grippers are pneumatically actuated. The opening and/or closing of the grippers is controlled with programmable magnetic switches SCHUNK MMS 22-PI1. A pneumatic nozzle between the grippers is installed, which is used for cleaning work-pieces and clamping accessories from sawdust and coolant.

DISCUSSION

Decision making in many areas can be very risky and difficult, affected by competing constraints and therefore cannot be individual. Multi Criteria Decision Making (MCDM) is a renowned methodology in decision making and evaluation. One of the famous MCDM tools is SWARA - Step-wise Weight Assessment Ratio Analysis method (Zolfani & Saparauskas, 2013).

Choosing end effectors can be enhanced by implementing the SWARA method in decision making. The objective of this study is to calculate relative weights of the average criteria by implementing the SWARA method (Merigó & Engemann, 2010; Zolfani & Saparauskas, 2013).

Based on their expected significances, the arithmetic mean values for the average clamping force in N are shown in the Table 1, sorted in descending order.

 Table 1:
 The presentation of arithmetic mean of clamping force in [N] for different gripper types

Criteria – Gripper type	Mean clamping force [N]
PGN-plus 64-1	633
PGN-plus 64-2	450
PGN-plus 64-1-AS	322
PGN-plus 64-2-AS	225

Table 2 displays the relative importance of the mean criteria for average clamping force. The relative importance of the mean criteria j, denoted as \tilde{S}_j to previous criteria (j-1), is given in the Table 2.

Solving the weight q of the j-th criteria, denoted by \tilde{q}_j , is given by Equation 2 (Keropyan & Gil-Lafuente, 2010):

$$\widetilde{q}_{j} = \begin{cases} 1, j = 1\\ \widetilde{x}_{j-1}\\ \widetilde{k}_{j}, j > 1 \end{cases}$$

$$(2)$$

Criteria – Gripper type	\widetilde{s}_{j}
PGN-plus 64-1	
PGN-plus 64-2	0.710
PGN-plus 64-1-AS	0.705
PGN-plus 64-2-AS	0.698

 Table 2:
 The relative importance of the mean criteria for average clamping force

The relative weight of the evaluation criteria j is determined in Equation 3 (Nicolas, 2004; Pérez Armayor & Díaz Batista, 2010), and is given in the rightmost column of the Table 3:

$$\widetilde{w}_{j} = \frac{\widetilde{q}_{j}}{\sum_{k=1}^{n} \widetilde{q}_{k}}$$
(3)

Table 3: The relative weight of the evaluation criteria

Criteria – Gripper type	\widetilde{S}_{j}	\widetilde{k}_{j}	\widetilde{q}_{j}	\widetilde{W}_{j}
PGN-plus 64-1		1	1	0.470
PGN-plus 64-2	0.710	1.710	0.584	0.274
PGN-plus 64-1-AS	0.705	1.705	0.342	0.160
PGN-plus 64-2-AS	0.698	1.698	0.201	0.096
			2.127	1.000

The SWARA method is often applied for two reasons. Firstly, SWARA's perspective is different from other similar methods – this method gives the chance to decision makers to select their priority based on the current situation of environment and economy. Secondly, the role of the experts is very important in the SWARA method. Experts have a key role in process of decision making on very important projects. Finally, SWARA has the advantage of more logical calculation of weights and relative importance of criteria (Zolfani & Saparauskas, 2013).

The summary of resulting weight of the mean criteria for the clumping force for all of the four selected types of grippers is shown in Figure 2.



Figure 2: Graphical representation of resulting weight of the mean criteria for average clumping force

Two universal parallel grippers such as SCHUNK PGN-plus 64-1 grippers are chosen as the end effectors. The maximum force of external clamping of the workpiece is 135 N, and the step is 4 mm per grip.

The grippers are pneumatically actuated, where the control of opening and closing of the grip is done via programmable SCHUNK MMS 22-PI1 magnetic switches. A pneumatic nozzle is installed between the grippers, which is used to clean the work-pieces and clamping accessories from sawdust and coolant (Ma *et al.*, 2007).

Two universal parallel grippers, with two grippers SCHUNK PGN-plus 64-1, in Figure 3 are presented. The grippers are made of aluminum bars having a profile that corresponds to the shape of the work piece, and the profiles are oriented towards the gripping position. The grippers are manufactured by the custom design and machined on a CNC machine on the customer's premises, in the customer's plant.

The solution with two grippers placed perpendicularly on the fork, at an angle of 90° to each other, is chosen (as shown in Figure 3) in order to save time. While the robot brings the preparation in one gripper, with the other it takes and lifts-up the finished work-piece. After that the robot pulls the work-piece out of the clamping accessories and rotates the last joint by 180° to swap the places of both grippers. This way, the double stroke of the robot arm from the clamping accessories to the pallet is eliminated (Ezugwu *et al.*, 2009).



Figure 3: Parallel grippers with two grippers SCHUNK PGN-plus 64-1

The programmable magnetic switches SCHUNK MMS 22-PI1 has the function of sending the information to the robot in order to close the feedback.

The sensor is mounted on the body of the catcher, and connected with the channels provided. The sensors can also be programmed using special programming tools (plug-in or magnetic). The sensors have LED lights that send signals to operator by varying the number and frequency of flashes. The sensors also send the information to the robot via cable.

In case of any physical problem while catching the work piece (jamming, incorrectly placed work piece, etc.), the robot receives a signal to stop further execution, and signals the stop by turning on the red light on its signal traffic light, as well as by sending a message with a description of the error to the control panel. If the light signal on the robot's traffic light fails, the CNC machine recognizes that the waiting time is longer than the usual cycle (the allowed value of time delay expressed in seconds previously entered in the programme).

After reporting an error on its control unit and turning on its traffic light, the CNC machine sends an SMS (Short Message Service) with a description of the problem to the mobile phone of the responsible person, and a message to the predefined e-mail addresses. All this is provided via the HaasConnect application. The ability to equip grippers of the robotic arm of the CNC machine with inductive sensor systems, magnetic switches, or work-piece-specific grips, improves their functionality, while increasing both the reliability and productivity of the process. The quality control of machining of the parts (with measuring and checking dimensions) is automated and performed using WIPS - Wireless Intuitive Probing System. A WIPS measuring probe manufactured by RENISHAW (RENISHAW mark - OMP40) is delivered with the CNC machine, as displayed in Figure 4. This probe enables fast and precise calculation (within the accuracy of the CNC machine) of the machined work piece on the CNC machine.



Figure 4: Measuring probe RENISHAW OMP40

The CNC machine manufacturer HAAS offers a complete system with WIPS equipment as a standard option. Considering the growing demands regarding precision and reduction of cycle time, as well as reduction of scrap, it is almost impossible to use a machining centre without this device.

Programming of a CNC machine for a cycle of inspection and quality control is done directly on the machine's control unit. The WIPS probe communicates

with the CNC machine over transceiver located in the workspace in the vicinity of the machine using an infrared (IR) radiation connection (Figure 6). The probe can be programmed to find the zero point of the workpiece, measure the dimensions of the work piece, touch the holes on the piece, tactilely scan the work piece, etc.

Communication between the measuring device and the CNC machine is shown in Figure 5.



Figure 5: Communication of the measuring device with the CNC machine



Figure 6: Measuring the tool dimension on the CNC machine

When the tool sizing cycle is started, the tool itself approaches the hard metal cylinder located on the top of the device and after touching the tip of the probe sends a signal to the machine control unit (Figure 6). At that moment, the machine compares the difference in the lengths of the axis of the movement and the previously calibrated (known) length at which the tip of the probe is located. Thus, the exact length of the tool is obtained, which is measured from the tip of the tool to the front of the spindle, and the machine automatically saves the measurement – it enters value in the tool table.

The same process is repeated along the second axis in case it is necessary to measure the diameter of the tool. Then the tip of the probe approaches the cylinder of the tool sideways, with the spindle rotating the tool in the opposite direction then while performing cutting or turning.

Measuring the tool dimension on the CNC machine is presented in Figure 6.

This measuring device is very useful in series production, because it has the ability to detect tool breakage. The tool check cycle and repetition interval are set on the machine control unit in the tool table. The principle of the operation is such that the machine has a predefined allowable path length that can be travelled when the tool approaches the measuring probe. If there is no contact between the tool tip and the measuring probe within this range, the machine signals a fault on the control unit. The cycle can be repeated periodically in predefined equal time intervals (after the tool has been in the grip for a certain period of time), or after each change of the tool.

The use of automated measuring devices for the machining quality control and inspection process reduces significantly the occurrence of scrap in automated production, where the human presence is not perpetual. The initial investment in systems of this type returns rather quickly through the decrease of the preparation time, which includes adjusting the position of the part or preparation and defining the dimensions of the cutting tools.

The classic calibration methods last approximately 11 minutes for the preparation, with 8 tools stacked to be ready to work. This system reduced that time to less than 2 minutes.

A possible plan for future work is to upgrade the existing system, described here, by installation of a 3D camera-sensor, which would provide an even higher level of autonomy and eliminate the need for a worker who serves the machine. Succesful implementations of artificial vision systems exist (Roberti et al., 2009). Instead of a human worker who places preparations on the pallet matrix and transports the finished pieces the robot would take the pieces from the box next to it, which would be filled with preparations by a conveyor belt. The robot would also place the finished pieces on another conveyor belt that would take these to the next operation (inspection and packaging). If this is the case, it would not be necessary to place the workpieces in the proper position on the pallet. This aproach would be a significant improvement towards Industry 4.0.

In recent years fuzzy logic has provided advances in the design and control of hybrid intelligent systems based on nature-inspired optimization. Applications of fuzzy and hybrid intelligent systems in areas such as intelligent control and robotics, pattern recognition, medical diagnosis, time series prediction, and optimization of complex problems have been explored (Melin *et al.*, 2018).

Dumitrescu *et al.*, (2021) introduced a concept of distributed intelligent control (DIC) and applied fuzzy logic to the control of an autonomous robot (vehicle). From the point of view of the DIC concept, three types of components can be defined and fuzzified (Dumitrescu *et al.*, 2021):

- Fuzzy sensors, which provide a representation of measurements as fuzzy subsets;
- Fuzzy actuators, which can act and operate in a real world; depending on the fuzzy subsets they receive;
- Fuzzy components of inference, also received by the fuzzy actuators, which can perform distributed fuzzy logic.

Fuzzy components can be integrated for different applications into a compact model of components, called the fuzzy cell. The configuration of the cells in the network is done through a specific programming language (Dumitrescu *et al.*, 2021).

Increasing the variety of products that market needs impose on manufacturers requires high adaptability of production systems that can be achieved through the introduction of flexible production systems composed of interoperable devices with variable architecture. Control and management of such a complex system requires fast and reliable real-time virtualization of real-world applications, as well as real-time feedback from realworld virtual (cyber) models (Wan *et al.*, 2015). The boundary line between the real production system and its cyber-representation is characterized by extremely high information throughput, combining these two systems into a single system, the Cyber-Physical Manufacturing System (CPMS). Recent advances in the fields of Cyber-Physical Systems (CPS) and the Internet of Things (IoT) have enabled the creation of CPMS.

CONCLUSION

In this study the SWARA method was applied on given data for the clamping force, obtained from the SCHUNK manufacturer. For the chosen criteria such as clamping force, the resulting weight was calculated. The degree of importance is presented in the appropriate tables, as well as in graphs. Taking into account the relative weight, the SCHUNK PGN-plus 64-1 grippers are chosen for the end effectors out of group of four types of the grippers.

In order to achieve adequate grip and capture of the object, the positioning and orientation of the grippers play a very important role. The industrial robot must achieve the precise relative positioning and orientation of the catcher with respect to the object, which in the case of hierarchical intelligent control can be placed in an arbitrary position.

Movement of an autonomous industrial robot in accordance with the recognized object, initiates many difficulties during completion of other defined phases of object manipulation. Thanks to visual feedback and learning based on artificial neural network systems, the robot catcher can determine and take a proper position and orientation with respect to the recognized object. This can be obtained in different ways, primarily by using systems with adaptive flexibility.

Smart behavior, which involves an autonomous industrial robot, is provided through adequate coordination between visual information about the object and surrounding space obtained from the sensor-camera and the control system. The smart control of the robot, realized by using sensor-cameras and artificial neural networks, has proven to be effective in overcoming the problems of modeling and the emergence of uncertainty in the real environment (*e.g.*, changing the position of objects).

The successor of the contemporary CNC machine could be called the ICNC machine (intelligent computer numerical control). In the future, there will be more flexible production cells controlled by artificial intelligence which will be able to 'learn', and the human presence will be decreased further and reduced only to assigning tasks.

Two decades ago, there were less than 700,000 industrial robots worldwide; however nowadays the number is over 3 million. In developed countries, there has been a simultaneous increase in production and a decrease in employment, and one, but not the only, cause is the development of technology. Today, 3D printing is used in the production of cars and airplanes, biotechnology is changing the way we grow crops and produce food, and the further development of nanotechnology and artificial intelligence will greatly reshape numerous economic activities, probably faster than we expect.

Currently the need for unskilled laborers whose work can be automated shows a downward trend and this is expected to continue, while the need for experts having skills to develop, launch and maintain highly automated systems is growing year by year.

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RESEARCH ARTICLE

Biomedical Engineering

Non invasive automated approach for eczema lesions segmentation using colour space normalization

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Abstract: Eczema is a common type of atopic dermatitis. Eczema skin lesions can be identified visually by observing the difference between the colour and texture of the lesions and the normal skin. Dermatologists assess eczema by direct visual assessment and record their observations in specialized forms. These methods are tedious as well as time consuming and introduce inter-rater and intra-rater variability in the results. To make the assessments objective and easier for the dermatologists, we have proposed a framework for the segmentation of eczema skin lesions. Red-Green-Blue (RGB), CIELab and their normalized colour spaces were considered. For segmentation, a two step K-means algorithm was proposed. In the 1st step, a conventional K-means algorithm segments the image into three regions, *i.e.*, skin, lesion, and mixed region. This is followed by a 2nd K-means segmentation step. The performance of this method was better than the conventional methods. The algorithm was evaluated using 85 eczema images of different severity and grades. To assess the performance of the algorithm, the gold standard segmentation for eczema lesions was manually drawn and verified by a dermatologist. The Green-channel of normalized (CSN-I) RGB colour space provided the best result for a semi-supervised approach giving the segmentation accuracy of 88.28% whereas for fully automated approach a segmentation accuracy of 84.43% was achieved using support vector machine (SVM).

Keywords: Atopic dermatitis, K-means segmentation, selection modes, skin lesions.

INTRODUCTION

Atopic eczema is a chronic, relapsing and inflammatory skin condition that is widespread in all age groups and may cause social and psychological problems (Carroll et al., 2005). Measuring the severity of a skin disease is the main objective in dermatology. It is essential to assess the course of the disease before, during and after the treatment. In skin diseases like atopic eczema, there are no serologic markers to identify the severity and extent of the disease. Hence evaluation of the disease is based on the visual signs and symptoms (Finlay, 1996). The general symptoms of atopic eczema are dry reddish skin with signs of inflammation. Due to scratching, the skin patches may be thickened, blistered, and sometimes infected. The different symptoms on eczema plaques 'erythema, edema/swelling, oozing/crusting, are: excoriation, and lichenification' (HOME, 2014). Subjective symptoms are itching and loss of sleep which also includes difficulty in falling asleep and/or waking up at night. Eczema is commonly found at the front side of elbows and wrists, around the neck, and back of the knees.

The prevalence of eczema appears to vary across the world as observed in key international studies (Lee &

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Detzel, 2015). The latest data based on the phase three of the 'International Study of Asthma and Allergies in Childhood' (ISAAC) (Mallol et al., 2013) showed that atopic dermatitis (AD) has reached a plateau in countries like the UK and New Zealand, but it continues to increase in prevalence, specifically in younger children (age 6-7 years) in low-income countries, in regions such as South East Asia and Latin America. This skin condition affects about 31.6 million people in the United States, and one in ten individuals will develop eczema during their lifetime (Silverberg & Hanifin, 2013). In the Asia Pacific, the 12-month prevalence of AD in children (age 13-14 years) was reported to be as high as 9% in Malaysia and Singapore, and as low as 0.9% in China, which has the lowest prevalence in the world. The reasons for the differences in prevalence are not fully understood, but industrialization and socio-economic factors have been implicated (Lee & Detzel, 2015).

The main objective of measuring the severity level of eczema is for assessing the progression of the disease in patients. The quantitative scales used for measuring eczema severity are difficult to interpret in clinical practice. Doctors use systems like 'Eczema Area and Severity Index' (EASI) (Barbier *et al.*, 2004), and 'Patient Oriented Eczema Measure' (POEM) (Charman *et al.*, 2002) to assess the severity of eczema. The process of manual form filling is time consuming, and involves subjective assessment that results in inter-rater and intrarater variability. Therefore, the objective of this research is to develop an automated eczema lesion segmentation system based on the EASI index, for assessment and grading of atopic eczema. The symptoms and severity levels for EASI scoring scheme are shown in Figure 1.

A comprehensive literature review of eczema has shown that very little research has been done for the development of a non-invasive, automated, objective method for the assessment and grading of the severity of eczema. A study conducted by Tremp et al. (2011) proposed the use of digital imaging techniques to assess the severity of atopic eczema using the EASI scoring system. They called their method EASIdig. In this method digital camera is used to take images of the patients; these images were evaluated by general physicians trained in EASI and SCORAD scoring systems. It took about thirty minutes to calculate the score using EASIdig. Another study reported by Alam et al. (2016) used 85 images to develop an automated approach for eczema detection and severity measurement. K-means clustering, and morphological image processing were used to segment eczema lesions using CIELab colour space. For classification ten colour and four texture features were used, reporting an accuracy of 90% using SVM classifier.



Figure 1: Symptoms and severity levels used for assessment in EASI scoring scheme (HOME, 2014)

Based on a literature review it is observed that other skin conditions such as Psoriasis, acne vulgaris, and melanoma have been studied extensively. The research is normally conducted for the classification of skin lesions originating from different diseases. For example, Hegde et al. (2018) used RGB colour and gray level cooccurrence matrix (GLCM) texture features to classify different types of skin lesions, i.e., chronic eczema, lichen planus, and plaque psoriasis. The machine learning methods SVM, naïve bayesian classifier (NBC), artificial neural networks (ANN), and linear discriminant analysis (LDA) were used. The classification accuracy using SVM classifier was 81.61%. Hameed et al. (2020) have proposed a framework for classification of healthy, benign, and malignant melanoma, and eczema images. A classification accuracy of 96.47% was achieved using colour and texture features.

Recently neural networks were investigated for melanoma lesion segmentation. Zafar *et al.* (2020) segmented dermoscopic images of melanoma using convolutional neural networks (CNN). They have combined UNet and ResNet architectures to form a novel architecture known as Res-Unet. A Jaccard Index of 77.2% was obtained. Abraham and Khan (2018) proposed U-Net and attention U-Net for segmentation of melanoma lesions. They achieved a dice score of 0.838 by U-Net and 0.856 by attention U-Net. The above literature review shows that tremendous research has been carried out for the classification of diseases where lesions are significantly different from each other in terms of surface properties, size, and shape.

In this paper our aim is to segment eczema skin lesions having the characteristic feature of no specific size and shape with a fuzzy boundary. The shape, size and colour of eczema lesions are not well defined; it can take any shape, spread to any size, and its colour varies a lot with the increase in the severity level. Some lesions are concentrated in a smaller area whereas others are scattered in a larger area. This makes the segmentation and classification of eczema lesions extremely difficult.

In the following sub-sections, we will give a brief overview of the pre-processing methods, colour spaces, segmentation, and classification techniques that we found useful for eczema lesion segmentation.

Image pre-processing

Image pre-processing removes noise and the effect of irregular illumination from an image. Ambient light condition is a very important criterion for image acquisition. The image segmentation may be affected by over-exposure or under-exposure of light. Two illumination correction techniques are found suitable for eczema lesion image illumination correction; these are adaptive light compensation (ALC) (Ch'ng *et al.*, 2014a) and Frankle-McCann retinex (retinex) (Funt *et al.*, 2004). In ALC the value of the luminance denoted as Y is extracted from the input image and average luminance (Y_{avg}) is computed. The maximum value of 200 and minimum value of 80 for good luminance exposure is obtained empirically from the images. A luminance factor is computed for RGB image correction as given in equation (1).

$$Factor = \begin{cases} 200 / Y_{avg}; & \text{if } Y_{avg} > 200 \\ 1 & ; & \text{if } 80 < Y_{avg} < 200 \\ 80 / Y_{avg}; & \text{if } Y_{avg} < 80 \\ & \dots(1) \end{cases}$$

The retinex method is designed to correct the illumination of under-exposed (dark) areas of an image while maintaining the visual discrimination of lighter areas (Funt *et al.*, 2004).

Colour spaces

Colour is an important property in visual perception. Different colour spaces are useful for different applications. A preliminary study with four colour spaces (HSI, YCbCr, CMY, CIELab) for eczema lesion segmentation using k-means showed that hue (H) channel of HSI has the highest segmentation accuracy of 76.63% (Nisar *et al.*, 2013).

Yang *et al.* (2010) proposed a colour space normalization (CSN) technique to improve the face recognition rate. His analyses of different colour spaces showed that those colour spaces perform better when the sum of elements in the 2^{nd} and 3^{rd} row of transformation matrix are zero. The RGB colour space does not possess this property; however, the YUV colour space has this property as shown in equations (2) and (3).

$$\begin{bmatrix} R \\ G \\ B \end{bmatrix} = \begin{bmatrix} 1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \end{bmatrix} \begin{bmatrix} R \\ G \\ B \end{bmatrix} \qquad \dots (2)$$

$$\begin{bmatrix} Y \\ U \\ V \end{bmatrix} = \begin{bmatrix} 0.2990 & 0.5870 & 0.1140 \\ -0.1471 & -0.2888 & 0.4359 \\ 0.6148 & -0.5148 & -0.1000 \end{bmatrix} \begin{bmatrix} R \\ G \\ B \end{bmatrix} \qquad \dots (3)$$

The CSN technique can normalize any colour space and the normalized colour space has a transformation matrix in which the sum of elements in the 2^{nd} and 3^{rd} rows are zero. Colour space normalization-I (CSN-I) is also known as 'within-colour-component normalization.' It is obtained by removing the means of 2nd and 3rd row vectors respectively. As a result, the sum of the row vectors is equal to zero in the transformation matrix. Equations (4) and (5) (Yang et al., 2010) show the transformation matrix for the CSN-I RGB, \tilde{A}_{RGB-} and CSN-I CIE-XYZ, \tilde{A}_{XYZ-I} colour spaces, respectively. Colour space normalization-II (CSN-II) is known as 'across-colourcomponent normalization.' In this technique the zeromean vector of the 2nd and 3rd rows is obtained via linear combination. Equations (6) and (7) (Yang et al., 2010) are the transformation matrices obtained for CSN-II RGB, \widetilde{A}_{RGB-I} and CSN-II CIE-XYZ, \widetilde{A}_{XYZ-I} colour spaces, respectively. Figure 2 gives an overview of different colour components.

$$\widetilde{A}_{RGB-I} = \begin{bmatrix} 1 & 0 & 0 \\ -1/3 & 2/3 & -1/3 \\ -1/3 & -1/3 & 2/3 \end{bmatrix} \dots (4)$$

$$\widetilde{A}_{XYZ-I} = \begin{bmatrix} 0.6070 & 0.1740 & 0.2 \\ -0.0343 & 0.2537 & -0.2193 \\ -0.3940 & -0.3280 & 0.7220 \end{bmatrix} \dots (5)$$

$$\widetilde{A}_{RGB-I} = \begin{bmatrix} 1 & 0 & 0 \\ -0.5774 & 0.7887 & -0.2113 \\ -0.5774 & -0.2113 & 2/3 \end{bmatrix} \dots (6)$$

$$\widetilde{A}_{XYZ-I} = \begin{bmatrix} 0.6070 & 0.1740 & 0.2 \\ -0.0901 & 0.3631 & -0.2730 \\ -0.4600 & -0.1986 & 0.6586 \end{bmatrix} \dots (7)$$



Figure 2: Illustration of different colour components for RGB, CIE Lab and their normalized CSN-I and CSN-II colour spaces

Segmentation and classification methods

In image processing, clustering methods are commonly used for segmentation of the desired regions of interest (ROI). The classification methods are divided into two categories: unsupervised learning and supervised learning classification methods. Unsupervised learning classifiers do not require any reference data for carrying out the classification. The classifier groups the data to a class itself. Clustering requires the user to define the hypothesized number of clusters. Commonly used methods of clustering are K-means clustering (Hartigan

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& Wong, 1979) and Fuzzy C-mean (FCM) (Bezdek *et al.*, 1984). However, supervised learning classifiers require a set of training data to work as a reference for classification. Supervised learning classification groups similar data based on the training data groups. Commonly used approaches are region-based, energy-based, and region and contour-based approaches. There are also some other classifiers such as SVM (Hegde *et al.*, 2018).

MATERIALS AND METHODS

The first step in any image processing-based application is image acquisition. It is followed by image preprocessing for image enhancement and noise removal. This is followed by image segmentation to obtain the ROI. In this paper we have proposed an algorithm to segment eczema skin lesions. We will discuss all the steps in the algorithm in the following sub-sections.

Data acquisition

The images were acquired using a digital single lens reflex (DSLR) camera. A dataset of 85 images in JPEG format is available for this study, the images were voluntarily provided. The gold standard for the skin lesions was manually drawn using the GNU Image Manipulation Program (GIMP) (The GIMP Team, 2014); and verified by a dermatologist. For performance assessment of segmentation, the gold standard images were used.



Figure 3: Comparison of pre-processing algorithms (a) input image; (b) adaptive light compensation (ALC) and (c) Frankle-McCann retinex (retinex)

Image pre-processing

Colour invariance due to uneven illumination makes the image segmentation more challenging. Hence, an ALC method (Ch'ng *et al.*, 2014a) was used to pre-process the images. We have also used another pre-processing method, Frankle-McCann retinex (Funt *et al.*, 2004) to compare the performance of pre-processing as illustrated in Figure 3.

Proposed segmentation approach

The K-means segmentation method is commonly used in unsupervised segmentation. We have proposed a bi-level K-means segmentation that divides the segmentation process into two steps (Ch'ng *et al.*, 2014b). The proposed technique is shown in in Figure 4. In the first step, it divides the image into three regions: the skin region, the mixed region, and the lesion region. Then the intensity feature [mean (μ) ± standard deviation (SD)], of skin or lesion region can be used as a reference value for the calculation of Mahalanobis distance to segment the mixed region into three sub-regions. We have used skin region as the reference as it gives better results.

For user supervised selection, the segmented cluster can be obtained by using one of the four different combinations as shown in the flowchart in Figure 4. The user will choose the most appropriate segmentation combination based on visual observation as shown in Figure 5. However, for the unsupervised method the SVM classifier with ranked features as training data will be used.



Figure 4: The proposed bi-level segmentation algorithm



Figure 5: Window for the operator to select a suitable cluster combination for manual selection in level 2

Level 1 clustering

After the pre-processing, the image was segmented into three regions using K-means clustering. The regions were labelled as skin region (cluster 1), mixed region (cluster 2), and lesion region (cluster 3). K-means clustering approach proposed by Yao *et al.* (2013) is used. Equation (8) shows the Euclidean distance (Brindha *et al.*, 2013) used in the 1st level of K-means clustering.

Euclidean Dist,
$$d(c, x) = \sqrt{\sum_{u=1}^{v} (x_u - c_u)^2}$$
 ...(8)

where *d* represents the distance, *x* and *c* represent the data points and centroid respectively, and *u* and *v* are the data dimensions used (here only 1 dimension of data is used).

Level 2 clustering

In Level 2 the mixed region is divided into three subregions (skin, mixed and lesion) using K-means clustering. The lesion is now composed of 5 clusters. Here the Mahalanobis distance (McLachlan, 1999) was used for the mixed region segmentation as it gives better results than the Euclidean distance (empirically observed). The Mahalanobis distance is given in equation (9).

Mahalanobis Dist,
$$d(x, y) = \sqrt{(y - \mu_x)/\operatorname{cov}(x) \times (y - \mu_x)^T}$$
...(9)

where y represents the data point and x represents the vector (mean \pm standard deviation) that is used as the reference in equation (9).

The segmentation of the final (mixed) cluster was the most difficult. It was done, using semi-supervised (operator input) or un-supervised (SVM) method.

Cluster combination for manual selection

For semi-supervised selection, operator input is required. The operator subjectively selects the most suitable combination of clusters that represents the segmented lesion. From Level 2 segmentation one cluster was combined with the lesion region at a time, and the option for four types of segmentation selection is present, as shown in the selection window in Figure 5. In the selection window the input image and pre-processed image are shown in the left column. The middle and right columns show the cluster combinations that were selected by the operator. In the right column, the boundary of the lesion was drawn using a black line. For the middle column

the region that does not belong to lesion region in the segmented image was excluded. The operator selects one of the four cluster combinations that best represents the segmented lesion, based on his judgement.

SVM classifier for automated selection

SVM was trained using selected features that are ranked by feature ranking methods. In this study, the T-statistic scoring method was used for feature ranking. The T-statistic scoring method calculates a score for the capability, C_T , of each feature to separate them between classes. Based on the value of C_T , the scoring of each feature was ranked in descending order. The two classes are denoted as 'a' and 'b'. For each feature, the mean, μ_a and μ_b ; standard deviation, σ_a and σ_b ; and the number of images used, n_a and n_b were calculated (Liu *et al.*, 2002). The mathematical expression for calculating C_T is given in equation (10).

$$C_{T} = \frac{\mu_{a} - \mu_{b}}{\sqrt{(\sigma_{a}^{2}/n_{a}) + (\sigma_{b}^{2}/n_{b})}} \qquad ...(10)$$

The score value obtained in equation (10) was used to sort the features in descending order. The combination of features was done by increasing one feature at a time. Finally, validation was done using 53 gold standard images; the validation error rate was obtained using equation (11).

$$\text{Error rate} = \frac{\text{Falsely classified data}}{\text{Overall data}} \qquad \dots (11)$$

Feature extraction and selection

For SVM training features were extracted from the skin and lesion regions of 20 gold standard images. These images were classified as mild images in context of the redness symptom. Five features extracted from the colour layers of four-colour spaces (RGB, CIELab, CSN-I RGB, CSN-I CIELab) are mean, standard deviation (SD), mode, skewness, and kurtosis, whereas the four features extracted from GLCM are contrast, correlation, energy, and homogeneity (Nisar et al., 2020). Colour layers of colour spaces were used for creating the GLCM; the intensity of the colour layer was normalized to '0' to '1'. A symmetrical GLCM was used with the horizontal direction only. The number of levels used for GLCM is 256. All extracted features were ranked using the T-statistical score ranking method and the combination of ranked features with lowest error was selected for SVM training.



Figure 6: Feature combination error rate graph

Figure 6 shows the feature combination error rate obtained during the training and classification of gold standard data. In this study, 108 features are used, of which 60 (5 features \times 4 colour spaces \times 3 layers) are colour features and 48 (4 features \times 4 colour spaces \times 3 layers) are texture features. It is observed from Figure 6 that the lowest error rate (less than 5%) is achieved for 10 features. Hence, we have used these 10 features for segmentation and classification, which are SD and kurtosis of the R channel, the energy of the G channel, correlation and homogeneity from CSN-I R, homogeneity from CSN-I G, Mean, skewness and homogeneity from CSN-I b*.

RESULTS AND DISCUSSION

In this paper we have proposed an algorithm for segmentation of eczema skin lesions. For pre-processing, the ALC method is used, and the results are compared with the retinex algorithm. The performance of the proposed segmentation algorithm is compared with K-means and FCM algorithms with three and five clusters. Two core colour spaces, RGB and CIELab, and two normalization techniques for colour spaces, CSN-I and CSN-II, are studied. The green channel of RGB colour space is selected for segmentation as it gives better segmentation performance when compared with red and blue channels; we denote it as RGB(G) (Ch'ng et al., 2014a). Similarly, the green channel is also chosen for CSN-I RGB and CSN-II RGB, and we denote it as CSN-I RGB(G) and CSN-II RGB(G), respectively. For CIELab colour space, chrominance channel a* is selected, and we denote it as CIELab(a*); for CSN-I CIELab and CSN-II CIELab illumination channel L* is selected and we denote it as CSN-I CIELab(L*) and CSN-II CIELab(L*), respectively.

For performance assessment, different metrics are used. True positive, true negative, false positive and false negative are denoted as TP, TN, FP, and FN respectively. These are used to calculate the precision, recall/ sensitivity (Sens), specificity (Spec), and accuracy (Acc), as given in equations (12)–(15). All the three parameters should have a high value for good classification results. The DICE coefficient is calculated using equation (16) to compare the similarity between the gold standard and segmented images.

$$Precision = TP/(TP + FP) \qquad \dots (12)$$

$$Recall/Sensitivity (Sens) = TP/(TP + FN)$$
 ...(13)

Specificity (Spec) =
$$TN/(TN + FP)$$
 ...(14)

Accuracy
$$(Acc) = (TP + TN)/(TP + TN + FP + FN)$$

...(15)

$$DICE \ Coefficient = (2 \times TP)/(2 \times TP + FP + FN)$$
...(16)

Table 1 shows the performance comparison of segmentation algorithms using the semi-supervised method: ALC as pre-processing technique and CSN-I RGB(G) colour channel. It is observed that all methods can achieve above 98% accuracy for the best-case segmentation, image 1; however, for image 2, the proposed algorithm has the highest accuracy whereas K-means and FCM with 5 clusters have better accuracy than with 3 clusters.

Tables 2 to 4 show the performance assessment of different segmentation algorithms. Out of the six

colour spaces evaluated, results for only top three best colour spaces are shown. Table 2 shows the overall segmentation performance for the proposed technique using semi-supervised and un-supervised methods. Among the three-colour spaces, CSN-I RGB(G) has the highest accuracy. Between pre-processing methods, ALC has a higher accuracy as compared to retinex.

Tables 3 shows the segmentation assessment for K-means using five and three clusters respectively. It is observed that the ALC pre-processing technique gives better segmentation accuracy. In K-means, for both pre-processing methods CSN-I RGB(G), CSN-II RGB(G) and CIELab(a*) are the three top ranked colour spaces that give the highest segmentation accuracy. Finally,

K-means with k = 5 gives higher segmentation accuracy as compared to the k = 3 case.

Table 4 shows the segmentation performance for FCM using five and three clusters respectively. For both pre-processing techniques, ALC has a better segmentation accuracy for FCM compared to retinex. For FCM the top three ranked colour spaces are the same as with K-means, as in Table 3. For both FCM (k = 5) and FCM (k = 3), CSN-I RGB(G) has highest segmentation accuracy. However, for the semi-supervised method using the retinex pre-processing technique there is an exception, as the CIELab(a*) colour space has a slightly higher segmentation accuracy compared to other colour spaces.

Table 1: Performance comparison of segmentation algorithms (semi-supervised method) using CSN-I RGB (G channel).



Table 2: Segmentation performance of the proposed two-level K-means algorithm (%)

		ALC	Retinex					
	Precision	Recall/Sens	Spec	Acc	Precision	Recall/Sens	Spec	Acc
Semi-supervised se	election							
CSN-I RGB(G)	86.78	81.76	90.37	88.28	87.78	76.86	87.93	86.40
CSN-II RGB(G)	82.52	80.34	87.77	85.98	80.07	82.08	82.79	84.63
CIELab(a*)	79.93	82.45	84.91	85.27	76.31	85.09	78.12	83.08
Un-supervised selection								
CSN-I RGB(G)	80.05	84.98	84.28	84.43	77.91	81.80	81.96	81.99
CSN-II RGB(G)	73.52	86.80	77.87	81.04	73.73	84.51	77.91	80.28
CIELab(a*)	71.95	87.17	75.33	80.20	74.22	84.25	79.10	80.81

		ALC		Retinex				
	Precision	Recall/Sens	Spec	Acc	Precision	Recall/Sens	Spec	Acc
Semi-supervised selection	on (5 cluste	rs)						
CSN-I RGB(G)	87.51	79.23	90.70	87.51	83.38	78.44	87.01	84.87
CSN-II RGB(G)	84.24	77.52	88.41	85.77	80.15	80.94	84.24	84.25
CIELab(a*)	81.04	79.94	85.33	84.67	80.06	82.73	83.65	84.79
Un-supervised selection	(5 clusters))						
CSN-I RGB(G)	76.71	84.77	76.04	79.78	73.89	83.24	73.47	78.23
CSN-II RGB(G)	68.64	88.15	68.19	76.25	70.28	85.52	68.68	76.39
CIELab(a*)	68.12	88.33	67.28	75.61	71.02	85.46	71.03	77.08
Semi-supervised selection	on (3 cluster	rs)						
CSN-I RGB(G)	86.88	77.84	87.37	86.23	79.91	81.30	80.41	82.49
CSN-II RGB(G)	81.33	80.08	84.07	84.32	80.70	77.95	84.13	82.84
CIELab(a*)	75.98	84.91	78.34	82.91	80.21	81.29	83.25	83.67
Un-supervised selection (3 clusters)								
CSN-I RGB(G)	80.01	82.55	81.68	82.46	78.88	79.26	79.31	80.31
CSN-II RGB(G)	72.91	86.25	74.78	79.50	76.78	80.87	78.60	79.77
CIELab(a*)	71.91	87.72	73.61	79.66	73.51	82.74	76.70	78.33

Table 3: Segmentation performance of the K-means with 5 and 3 clusters (%)

Table 4: Segmentation performance of Fuzzy C mean with 5 and 3 clusters (%)

		ALC		Retinex				
	Precision	Recall/Sens	Spec	Acc	Precision	Recall/Sens	Spec	Acc
Semi-supervised select	ion (5 cluster	rs)						
CSN-I RGB(G)	87.52	78.60	91.37	87.36	84.25	78.59	88.65	85.65
CSN-II RGB(G)	85.02	77.15	89.18	86.08	81.10	81.16	84.27	84.44
CIELab(a*)	80.13	81.30	85.11	84.63	78.89	83.57	83.24	84.31
Un-supervised selection	n (5 clusters)							
CSN-I RGB(G)	77.79	84.16	76.71	79.90	74.48	82.57	73.76	78.10
CSN-II RGB(G)	68.17	87.80	67.96	75.94	72.21	84.55	71.62	77.02
CIELab(a*)	67.83	88.62	66.90	75.61	70.36	86.09	70.20	76.75
Semi-supervised select	ion (3 cluster	rs)						
CSN-I RGB(G)	87.78	76.86	87.93	86.40	83.34	78.02	83.05	83.43
CSN-II RGB(G)	80.07	82.08	82.79	84.63	80.59	79.08	83.66	83.11
CIELab (a*)	76.31	85.09	78.12	83.08	80.49	80.92	81.73	83.60
Un-supervised selection (3 clusters)								
CSN-I RGB(G)	79.80	81.74	81.43	81.70	80.00	79.74	79.87	80.90
CSN-II RGB(G)	72.57	86.38	73.97	78.90	76.39	81.66	77.62	79.38
CIELab(a*)	71.11	87.94	72.63	79.06	72.68	83.42	74.84	77.59

Table 5: Comparison of the segmentation algorithms in terms of DICE coefficient (%)

			Semi-suj	pervised	selection	Un-supervised selection					
		Proposed	K-means (k=5)	FCM (k=5)	K-means (k=3)	FCM (k=3)	Proposed	K-means (k=5)	FCM (k=5)	K-means (k=3)	FCM (k=3)
<i>T</i>)	CSN-I RGB(G)	83.47	81.72	81.74	79.96	79.73	80.17	76.74	77.01	77.41	76.44
ALC	CSN-II RGB(G) CIELab(a*)	80.56 80.62	79.70 79.21	79.69 79.52	78.51 78.48	79.06 78.72	77.28 76.47	73.94 73.30	73.47 73.32	75.87 76.27	75.46 75.67
Xa	CSN-I RGB(G)	80.41	79.34	80.29	77.75	77.88	76.99	74.09	73.65	74.97	76.02
etine	CSN-II RGB(G)	79.94	79.32	79.96	76.71	77.61	76.13	73.79	74.14%	75.65	75.60
R	CIELab(a*)	80.61	80.09	79.77	78.46	78.13	76.31	74.00	73.76	73.63	73.34

Table 5 shows the comparison in terms of DICE coefficient. When comparing between pre-processing methods, ALC performs better than retinex. CSN-I RGB(G) colour space outperforms all the other colour spaces with highest performance accuracy. For segmentation techniques, the proposed bi-level K-means segmentation gives a better accuracy for most of the cases. There are only six cases out of 24 where bi-level K-means does not give the best accuracy; these are RGB(G) in ALC for both the semi-supervised and unsupervised selection mode, CSN-II RGB(G) in ALC, and CIELab(a*) colour space with both normalized techniques in retinex for semi-supervised selection mode. Conventional K-means seems to have a slightly better performance than the proposed method for those cases. Finally, we can say that the proposed method gives an overall best accuracy for the semi-supervised method, which is 88.28%, whereas the un-supervised selection mode has an accuracy of 84.43%.

For the semi-supervised selection mode, segmentation with a higher cluster number (5) gives a better accuracy compared to fewer clusters (3) for both K-means and FCM, whereas the accuracy of the unsupervised selection mode tends to be lower for the higher number of clusters as compared to the lower number of clusters used. For the proposed segmentation algorithm, the input image will first be segmented into three regions in the first step and then the mixed region will further be segmented into another three sub-regions. For the unsupervised selection mode, the poor performance with a higher number of clusters for K-means and FCM may be due to the features selected for training by SVM, despite using the features with lowest validation error rate.

Hence from the results it is observed that the CSN-I RGB(G) colour space has better segmentation accuracy resulting in the highest similarity between the gold standard and segmented images. The colour space normalization technique proposed by Yang *et al.* (2010) shown in equation (4) amplifies the difference between normal skin and an eczema lesion in the green channel of CSN-I RGB colour space, that leads to a better segmentation for the proposed bi-level K-means segmentation algorithm.

CONCLUSION

In this paper we have investigated the performance of colour spaces, pre-processing methods, segmentation techniques and selection modes. Investigations show that G channel of CSN-I RGB outperforms all other colour spaces for the Eczema skin lesion segmentation.

The proposed segmentation method and pre-processing technique ALC performed better in most cases. The highest segmentation accuracy achieved is 88.28% for proposed bi-level segmentation method using CSN-I RGB (G channel) colour space for semi-supervised selection. While highest accuracy achieved for unsupervised selection is 84.43% with the proposed segmentation method using the CSN-I RGB (G channel) colour space. In future we plan to increase the image database for an effective evaluation. Secondly more features should be considered for feature selection; used for training the supervised learning classifier. Deep learning methods will also be considered.

Conflict of interest

The authors declare that they have no conflict of interest.

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RESEARCH COMMUNICATION

Natural Products

Bioactive cyclo-(S-Pro-*R*-Leu) from Aspergillus flavus, the marine endophytic fungus from brown alga, *Dictyota kunthi*

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Abstract: The marine environment is regarded as a rich source of bioactive secondary metabolites, and some of these compounds have been shown to be useful for novel drug discovery. In recent years, research on the chemistry of endophytic fungi from marine algae has shown them to be an important source of secondary metabolites with a wide range of biological activities. The objective of this study was to describe the isolation and identification of bioactive compounds obtained from Aspergillus flavus, hosted in the brown alga, Dictyota kunthi, together with β -glucuronidase bioassay of them. This fungal strain was cultivated on a large scale and extracted with ethyl acetate (EtOAc). Using various column chromatographic and spectroscopic techniques, we isolated and characterized two metabolites, cyclo-(S-Pro-R-Leu), 1 and genistein 2. Cyclo-(S-Pro-R-Leu) inhibited β -glucuronidase with an IC $_{_{50}}$ value of 83.9 \pm 0.1 μM when compared to glucosaccharo-1,4-lactone (IC₅₀ = $48.4 \pm 1.3 \mu$ M). This appears to be the first report of the isolation of cyclo-(S-Pro-R-Leu) from this entophyte. To the best of our knowledge, this is the first report of significant β -glucuronidase inhibitory activity of this compound. From these findings, we can confirm that the endophyte from the brown alga, Dictyota kunthi, produces a β -glucuronidase inhibitory metabolite.

Keywords: β -glucuronidase, *Aspergillus flavus*, cyclo-(*S*-Pro-*R*-Leu), *Dictyota kunthi*, marine endophytes.

INTRODUCTION

Due to the discovery of unique chemical structures and their bioactivity, the search for novel chemicals from the marine environment has rapidly increased in recent years (Teixeira et al., 2019). Endophytic fungi live in the inner tissues or even cells of their hosts (Zhang et al., 2009). Secondary metabolites of algal-derived endophytic fungi have proven to be a good source of novel bioactive secondary metabolites for the development of pharmaceuticals and agrochemicals, showing activities such as anticancer, antibiotic, antiviral, antioxidative, and kinase inhibitory or activating activities (Sarasan et al., 2017; Handayani et al., 2019). In continuation of our studies towards the discovery of biologically active compounds from the Sri Lankan seaweeds and their endophytes (Haroon et al., 2013; Haniffa et al., 2019) we isolated the fungal strain A. flavus, from the brown alga, D. kunthi, collected from the south coast of Sri Lanka. This fungal strain was cultivated on a large scale and extracted using EtOAc. We extracted and characterized two metabolites, 1 and 2, using various column chromatographic techniques. Compounds 1 and 2 were evaluated *in vitro* for β -glucuronidase inhibitory activity and compound 1 was found to have significant activity. This appears to be the first report of significant β -glucuronidase inhibitory activity of this compound. The activity of β -glucuronidase rises in various conditions, including cancer, some hepatic disorders, and AIDS (Awolade et al., 2020). Therefore, development of specific inhibitors of β -glucuronidase could be useful. The present study describes the isolation and identification of cyclo-(S-Pro-R-Leu), 1 and genistein 2 together with β -glucuronidase bioassay for screening of 1. To our knowledge, this is the first report of the isolation of cyclo-(S-Pro-R-Leu) from this endophytic fungus.

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MATERIALS AND METHODS

General procedure

A Büchi 535 melting point instrument was used to determine melting points. A Jasco DIP-360 digital polarimeter was used to quantify optical rotation in MeOH. Bruker AMX 300 and 500 MHz NMR spectrometers were used to record ¹H-NMR spectra, while the same instrument was used to conduct ¹³C-NMR studies at 75.45 and 100 MHz, respectively. Chemical shifts (δ) were measured in ppm relative to TMS, with coupling constants *J* measured in Hz. A Jeol JMS HX 600 mass spectrometer was used to detect EI MS.

Isolation and identification of the fungal strain

D. kunthi, a brown alga, was collected at Kirinda on the southern coast of Sri Lanka. The endophytic fungi were isolated from the blade of the seaweed using the method developed to isolate endophytes from higher plants by Petrini (Fokkema & Heuvel, 1986), with certain modifications (Haroon *et al.*, 2013). The endophytic strain (DK-11) was identified as *A. flavus*, by S.R. Premaratne (Institute of Fundamental Studies, Natural Products Programme, Kandy, Sri Lanka). A subculture has been deposited at the Natural Products Laboratory, Institute of Fundamental Studies, Kandy.

Cultivation of Aspergillus flavus

The growth medium (Czapex Dox) for *A. flavus* was prepared by mixing the following chemicals (g/L) in filtered (No. 3 Whatman filter paper) natural seawater (500 mL/L) and distilled water (500 mL/L): K_2HPO_4 (1.0 g), MgSO₄ 7H₂O (0.5 g), KCl (0.5 g), FeSO₄ 7H₂O (0.01 g), sugar (30 g), and peptone (30 g). Culture medium (6.0 L) was prepared and distributed (250 mL, each flask) among 24 Erlenmeyer flasks of 1000 mL capacity (each), and autoclaved at 121 °C for 30 min. Then, agar blocks inoculated with mycelia were added into a 1000 mL flask containing 250 mL of culture medium and incubated at 28 °C with a control (without fungal strain) for 14 ds.

Extraction and isolation of pure compounds

After 14 ds, 250 mL of EtOAc was added into each flask containing culture media and mycelium. They were mixed thoroughly using a sonicator and the EtOAc

layer was separated from each flask and evaporated using a rotary evaporator at 50 °C. The EtOAc extract was dried in a vacuum oven to afford a white residue (1.5 g). Flash column chromatography (CC) on silica gel (230 - 400 mesh size) produced two pure compounds from the EtOAc extract. This column was initially eluted with gradient polarities of n-hexane and EtOAc, which afforded five fractions (F -1 to F-5) and their weights were given as. F-1 (0.3 g), F-2 (0.1 g), F-3 (0.7 g), F-4 (10 mg), and F-5 (20 mg). Repeated flash silica gel (SiO₂) column chromatography of F-3 (20% EtOAc:hexane) yielded pure compound 1 (40 mg, 2.6%, 15% Acetone:hexane). Fraction F-4, when subjected to column chromatography, yielded compound 2 (2 mg, 0.13%, 20% Acetone : hexane).

β-glucuronidase inhibition assay

The β -glucuronidase inhibition assay was performed according to the method developed by Collins *et al.* (Collins *et al.*, 1997; Haroon *et al.*, 2013), with slight modification.

Characterization of compounds

Cyclo-(S-Pro-R-Leu) (1)

A white amorphous solid (40 mg, 2.6%) mp 170 °C; $[\alpha]^{27}_{D}$ -105.5° (c = 0.5, EtOH) UV (MeOH), λ_{max} at 242 nm; IR (KBr) ν_{max} 1432, 1635, 1670, 2955, 3261 cm⁻¹; ¹H NMR (400 MHz, CDCl₃); δ_{H} 0.92 (3H, d, J = 6.5 Hz, H₃-12), 0.96 (3H, d, J = 6.5 Hz, H₃-13), 1.51(1H, m, Ha-10), 1.74 (1H, m, H-11), 1.85 (2H, m, H₂-4), 2.06 (1H, m, Hb-10), 2.13 (1H, m, Ha-5), 2.33 (1H, m, Hb-5), 3.53-3.6 (2H, m, H₂-3), 3.91 (1H, dd, J = 9.5, 3.4 Hz, H-9), 4.15 (1H, t, J = 7.7 Hz, H-6), 6.16 (1H, s, NH); ¹³C NMR (75 MHz, CDCl₃); δ_{C} 21.2 (C-13), 22.7 (C-4), 23.2 (C-12), 24.6 (C-11), 28.0. (C-5), 38.6 (C-10), 45.4 (C-3), 53.4 (C-9), 59.4 (C-6), 166.2 (C-7), 170.3 (C-1); CI MS (positive mode) m/z 211 [M + H]⁺.

Genistein (2)

A pale yellow powder (2 mg, 0.13%) mp 286°C lit (Aida *et al.*, 1995) 285 °C; ¹H NMR (500 MHz, CD₃OD); $\delta_{\rm H}$ 6.21 (1H, d, J = 2.2 Hz, H-8), 6.31 (1H, d, J = 2.2 Hz, H-6), 6.84 (2H, d, J = 8.6 Hz, H-3'/5'), 7.35 (2H, d, J = 8.5 Hz, H-2'/6'), 8.04 (1H, d, H-2); EI MS 70 eV, *m/z* (rel.int, %): 270 (100) [M⁺], 153 (56), 137(10), 118 (21).

RESULTS AND DISCUSSION

Chemistry of isolated compounds

Repeated column chromatography on flash silica gel yielded two metabolites from the EtOAc extract of A. flavus. Compound 1 was found to be a white amorphous solid with a melting point of 170 °C. The compound's molecular ion peak was found at m/z 211 [M + H]⁺with the aid of CI-MS (positive mode). The amide NH group absorbed at 3261 cm⁻¹ in the infrared spectrum of 1, while the two amide carbonyl groups absorbed at 1635 and 1670 cm⁻¹. In the¹H-NMR spectrum, recorded in CDCl₂, the most downfield singlet, that of the amide NH, appeared at $\delta_{\rm H}$ 6.16. The other two downfield sets of protons, H-6 and H-9 appeared as triplet at $\delta_{\rm H}$ 4.15 (1H, t, $J_{\rm 6,\,5}$ = 7.7 Hz) and as double doublet at $\delta_{\rm H}\,3.91$ (1H, dd, $J_{9, 10a/10b} = 3.4$, 9.4 Hz,) respectively. The ¹³C-NMR broad-band decoupled, HMQC and DEPT spectra of 1, in CDCl, indicated 11 carbon signals, including 2 methyls, 4 methylenes, 3 methines, and 2 quarternary carbons. The most downfield signals, those of amide carbons, appeared at δ_c 170.3 and 166.2 while the second most downfield carbon sandwiched between NH and CO appeared at δ_c 58.9. The methine carbon attached to C-6 and the methylene carbon attached to C-3 of 1 appeared at δ_c 53.4 and 45.4 respectively. Compound 1 was identified as cyclo-(S-Pro-R-Leu) by direct comparison with reported ¹H- and ¹³C-NMR spectral data (Adamczeski et al., 1995; Pedras et al., 2005; Liu et al., 2012) (Figure 1). The absolute configuration was determined by the optical rotation, the $[\alpha]_{D}^{27}$ (c = 0.5, EtOH) value of 1 (-105.5°), well agreeing with reported data, $[\alpha]_{D} = -108$ (Adamczeski *et al.*, 1995; Li *et al.*, 2008). However, according to Yang et al. (2009), it showed the optical rotation of [a]D = -78.3.

Compound 2 was found to be a pale yellow powder with a melting point of 286 °C. The molecular ion and base peak of compound 2 was found at m/z 270, which corresponded to the chemical formula $C_{15}H_{10}O_5$, indicating eleven degrees of unsaturation. The most downfield singlet seen δ_H 8.04 in the ¹H-NMR spectra recorded in CD₃OD was ascribed to olefinic H. Moreover, H-2'/6' and H-3'/5' were ascribed to the paradisubstituted aromatic ring protons that resonated at δ_H 7.37 (2H, d, J = 8.6 Hz) and δ_H 6.84 (2H, d, J = 8.6 Hz), respectively.

The presence of tetra - substituted aromatic ring protons was suggested by meta coupling signals at $\delta_{\rm H}$ 6.32 (1H, J = 2.2 Hz, H-8) and 6.21 (1H, J = 2.2 Hz,

H-6) in the ¹H-NMR spectrum of 2. Compound 2 was confirmed as genistein based on a direct comparison of physical and spectral data (Aida *et al.*, 1995) (Figure 1). Isoflavones are known to occur commonly in higher plants, but they have also been isolated from bacteria, and from a culture of the fungus *Aspergillus niger*.



Figure 1: Structures of isolated compounds 1-2.

Table 1: β -glucuronidase inhibition of pure compounds

Compound	IC ₅₀ (µM)
Compound	$Mean \pm S.E.M$
1	83.9 ± 0.1
2 glucosaccharo-(1,4)-lactone	- 48.4 ± 1.3

β-glucuronidase inhibitory activity

The compounds 1 and 2 were screened for β -glucuronidase inhibitory activity at 200 μ M. When compared with the standard glucosaccharo-(1,4)-lactone $(IC_{50} = 48.4 \pm 1.3 \mu M)$, compound 1 showed a significant β -glucuronidase inhibitory activity with an IC₅₀ value of $83.9 \pm 0.05 \,\mu\text{M}$ and 2 did not show any activity (Table 1). The substantial inhibitory activity of Compound 1 is most likely due to the enzyme's active site accepting a proton from the CONH. This compound was reported to inhibit the marine pathogenic bacteria, Vibrio anguillarum (Fdhila et al., 2003) and fungi Plasmopara viticola (Musetti et al., 2007). Further, it also exhibits antifouling activity against microfoulers (Li et al., 2006). To the best of our knowledge, cyclo-(S-Pro-R-Leu) from algae-associated fungi is uncommon compared to sponge-associated fungi, actinomycetes, and bacteria (Schmidtz et al., 1983; Stierle et al., 1988; Adamczeski et al., 1995; Fdhila et al., 2003; Li et al., 2006; Li et *al.*, 2008; Yang *et al.*, 2009). According to Liu *et al.* (2012), cyclo-(*S*-Pro-*R*-Leu), which has already been shown to be produced by the close species of *Aspergillus fumigates*, showed weak inhibitory activity against the release of β -glucuronidase from rat polymorphonuclear leukocytes (PMNLs). However, as per the results of this study, cyclo-(*S*-Pro-*R*-Leu) showed a significant β -glucuronidase inhibitory activity with an IC₅₀ value of 83.9 ± 0.1 µM. To the best of our knowledge this is the first report of this compound showing significant β -glucuronidase inhibitory activity.

CONCLUSION

The present study described the isolation and characterization of cyclo-(*S*-Pro-*R*-Leu) and genistein from EtOAc extract of *A. flavus* hosted in the brown alga, *Dictyota kunthi*. When compared to the standard glucosaccharo-(1,4)-lactone (IC₅₀ = 48.4 ± 1.3 μ M), the compound cyclo-(*S*-Pro-*R*-Leu) demonstrated significant β -glucuronidase inhibitory activity with an IC₅₀ value of 83.9 ± 0.1 μ M. Based on these findings, we can conclude that the endophyte from *Dictyota kunthi* produces a β -glucuronidase inhibitory metabolite.

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Conflict of interest statement: All authors should include a statement on conflict of interest disclosing any financial or other substantive conflicts of interest that may be construed to influence the results or interpretation of their research. All sources of financial support for the project should be disclosed.

Acknowledgement: Should be brief and made for specific scientific, financial and technical assistance only. If a significant part of the research was performed in an institution other than in those indicated by the authors' affiliations given in the title page, this fact should be acknowledged. All those who have made substantial contribution to the research but do not qualify to be authors should be acknowledged.

References :

All research work of other authors, when used or referred to or cited, should be correctly acknowledged in the text and in the References.

Citing references in the text:

- References to the literature must be indicated in the text and tables as per the Author-Year System, by the author's last name and year, in parenthesis (i.e. Able, 1997) or (Able & Thompson, 1998).
- Citation to work by more than two authors should be abbreviated with the use of et al. (i.e. Able et al., 1997).
- Multiple publications by the same first author in the same year should be coded by letters, (i.e. Thompson, 1991a; b).
- Multiple citations should be made in chronological order and separated by a semi-colon, (i.e. Zimmerman et al., 1986; Able et al., 1997).
- Reference to unpublished work, work in preparation or work under review should be cited in italics as (*unpublished data*) or, with the author's initials and surname given; such works should not be included in the Reference section.
- Personal communications may be mentioned in the text with the date of communication as (Personal communication, 2 June 2000).

List of references:

- The list of References should be arranged in alphabetical order based on the last name of the first author.
- Names of all the authors should be given except when there are more than 10 authors. When there are more than 10 authors, only the name of the first author can be given followed by *et al*.
- All the initials of the author must be given after the last name and the year of publication should follow in parentheses.
- This should be followed by the full title of the referred publication.
- When journal articles are listed, the journal name should be given in full and in italics and followed by the volume number in bold type, issue number in parentheses and then the inclusive pages.
- Where there are several publications by the same author(s) and published in the same year they should be differentiated by adding a lower-case letter after the year. When books are listed, the order should be: author(s), year, book title, volume number, edition, pagination/ inclusive pages, publisher and place of publication. The book title should be in italics. When sections of a book are listed, the order should be: author(s) of chapter, year, title of the section, title of the book, edition, inclusive pages, publisher and place of publication.
- Digital object identifiers (DOIs) should be included for all references where available.
- References should only be cited as 'in press' if the paper has been accepted for publication.

Examples of correct forms of references are given below.

Journal Articles

Boutin C. & Harper J.L. (1991). A comparative study of the population dynamics of five species of *Veronica* in natural habitats. *Journal of Ecology* **79**(01): 199 – 221.

DOI: https://doi.org/10.2307/2260793

Books

Burnham K.P. & Anderson D.R. (2002). *Model Selection and Multimodal Inference*, 2nd edition, pp. 488. Springer Science and Business Media, Inc., New York, USA.

Book Chapters

Hinrichsen R.A. & Holmes E.E. (2009). Using multivariate state-space models to study spatial structure and dynamics. In: Spatial Ecology (eds. R.S. Cantrell, C. Cosner & S. Ruan), pp. 145 – 166. CRC/ Chapman Hall, Florida, USA. DOI: <u>https://doi.org/10.1201/9781420059861.ch8</u>

Edited Books

Kimatl H., Amorim L., Rezende J.A.M., Bergamin Filho A. & Camargo L.E.A. (eds.) (2005). *Manual de Fitopatologia*, volume 2. Doenças das Plantas Cultivadas, 4th edition. Ceres, São Paulo, Brazil.

Conference Papers

Weaver D. (2002). Implementation of a learning management system using an integrated approach to professional development. In: Winds of change in the sea of learning. *Proceedings of the 19th Annual Conference of the Australasian Society for Computers in Learning and Tertiary Education (ASCILITE)* (eds. A. Williamson, C. Gunn, A. Young & T. Clear), volume 2, Auckland, New Zealand, 8-11 December. Unitec Institute of Technology, Auckland, New Zealand, pp. 711-720.

Agency Publications

U.S. Census Bureau (2009). World Population: 1950 - 2050. U.S. Census Bureau, Washington DC, USA.

Department of Health (2008). *Health Inequalities: Progress and Next Step* (pdf). Department of Health, London, UK. Available at *http:// PublicationsPolicyAndGuidance/DH_08 5307*, Accessed 9 June 2008.

Other

Robinson L.J. (2003) Spatial scale and depletion models of farmland birds in a fragmented landscape. *PhD thesis*, University of Reading, Reading, UK.

Efford M.G. (2008). Density 4.3: software for spatially explicit capture-recapture. Available at *http://www.otago.ac.nz/density* , Accessed 15 March 2009.

Abbreviations and Symbols: Unless common, these should be defined when first used, and not included in the abstract. The SI System of units should be used wherever possible. If measurements were made in units other than SI, the data should be reported in the same units followed by SI units in brackets, e.g. 5290 ft (1610 m).

Formulae and Equations: Equations should be typewritten and quadruple spaced. They should be started on the left margin and the number placed in parentheses to the right of the equation.

Nomenclature: Scientific names of plants and animals should be printed in italics. In the first citation, genus, species and authority must be given. e.g. *Borassus flabellifer* Linn. In latter citations, the generic name may be abbreviated, for example, *B. flabellifer* L.

Tables and figures: Tables and Figures should be clear and intelligible and kept to a minimum, and should not repeat data available elsewhere in the paper. Any reproduction of illustrations, tabulations, pictures etc. in the manuscript should be acknowledged.

Tables: Tables should be numbered consecutively with Arabic numerals and placed at the appropriate position in the manuscript. If a Table must be continued, a second sheet should be used and all the headings repeated. The number of columns or rows in each Table should be minimized. Each Table should have a title, which makes its general meaning clear, without reference to the text. All Table columns should have explanatory headings. Units of measurement, if any, should be indicated in parentheses in the heading of each column. Vertical lines should not be used and horizontal lines should be used only in the heading and at the bottom of the table. Footnotes to Tables should be placed directly below the Table and should be indicated by superscript lower case italic letters (a , b , c , etc.).

Figures: All illustrations are considered as figures, and each graph, drawing or photograph should be numbered consecutively with Arabic numerals and placed at the appropriate position in the manuscript. Any lettering to appear on the illustrations should be of a suitable size for reproduction and uniform lettering should be used in all the Figures of the manuscript. Scanned figures or photographs should be of high quality (**300 dpi**), to fit the proportions of the printed page (12×17 cm). Each figure should carry a legend so that the general meaning of the figure can be understood without reference to the text. Where magnifications are used, they should be stated.

Units of measurement

Length: km, m, mm, µm, nm Area: ha, km², m² Capacity: kL, L, mL, µL Volume: km³, m³, cm³ Mass: t, kg, g, mg, µg Time: year(s), month(s), wk(s), d(s), h, min, s Concentration: M, mM, N, %, g/L, mg/L, ppm Temperature: °C, K Gravity: x g Molecular weight: mol wt Others: Radio-isotopes: 32P Radiation dose: Bq Oxidation-reduction potential: rH Hydrogen ion concentration: pH