

Environmental Science and Engineering

Aravind Jeyaseelan
Kamaraj Murugesan
Karthikeyan Sivashanmugam *Editors*

Sustainable and Cleaner Technologies for Environmental Remediation

Avenues in Nano and Biotechnology

 Springer

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Editors

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Chapter 1

Catalytic Degradation of 4-Nitrophenol and Organic Dyes Using Silver and Platinum Nanoparticles Synthesized by Plant Extracts



M. Karthikeyan, V. Blessy, and A. Thirumurugan

Abstract For a decade, nanotechnology research has been one of the most attractive field of wastewater treatment. In the present study, green route synthesized noble metallic nanoparticles of silver and platinum catalytic activity were assessed for degradation of 4-Nitrophenol as a model by UV–visible spectroscopy. Then, catalytic activity of typical organic dyes of Alizarin red S and Brilliant blue G were also performed and monitored by UV–visible spectroscopy. The efficiency of silver and platinum nanoparticles in reducing the 4-Nitrophenol and dyes were compared. The catalytic activities of both nanoparticles were also quantified by analyzing the sodium Borohydride reduction of 4-Nitrophenol. As a result, it has been observed that the silver nanoparticles show more reducing ability than platinum nanoparticles.

Keywords Silver NP's · Platinum NP's · Green synthesis · Catalytic activity · 4-Nitrophenol · Organic dyes

1.1 Introduction

The need for the production of nano materials has increased due to their various applications in the field of electronics, catalysis, medicine etc., Since the materials are at nano scale have different and unique properties when compared with bulk materials, they gain more attention in various applications (Akilandaeaswari and Muthu 2021). Mostly the nano materials are prepared from noble metals such as gold, silver, platinum etc., (Khan et al. 2022). Silver is one of the most used metals which are widely used as a disinfecting agent (Lee and Park 2020). Silver nanoparticles have effective antibacterial activity when compared with other nanoparticles and hence it is widely used in the field of dentistry, catheters, and making burn wounds to control bacterial growth (Wan et al. 2021). The platinum nanoparticles also have

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wide application in the field of catalysis, biomedicine, sensing, cancer therapy etc., but the production of platinum nanoparticles was not much explored in the field of nanotechnology (Joseph and Vijayanandan 2021).

Organic dyes are the major pollutants of water source which are widely used in textile industries, research activities etc., and hence the removal of such dyes are major concern (Haverkamp and Marshall 2009). Alizarin red S is the most used dyes in textile industry. It is 1, 2-dihydroxy anthraquinone sulfonic acid sodium salt which is synthesized by sulfonation of alizarin. Alizarin Red S also gains applications in various fields such as staining in microscopy, acid base indicator, determination of fluorine etc., (Kavitha et al. 2013). Coomassie brilliant blue which is a non-azo dye find its wide application in research laboratories especially in protein estimations. It is also widely used for staining in electrophoresis, Bradford assay etc., (Akhtar et al. 2013). Since these dyes heavily contaminate the water, they should be treated before discharged into the environment (Banerjee et al. 2014) various physical and chemical methods such as adsorption, coagulation, flocculation, photo catalysis etc., are employed to treat such dye containing water (Sharma et al. 2014). Nanoparticles synthesized from various noble metals have ability to catalyze various chemical interactions like hydrocarbons oxidation, hydrogen bonding, as well as redox reactions. As they have more surface area compared with their bulk materials, they have more and new catalytic properties (Dash et al. 2015). Because of these advantages, nanoparticles play a major role in catalyzing the reduction of organic dyes. The sodium borohydride reduction of 4-Nitrophenol to aminophenol group is the well-known chemical reaction. Hence it is widely used to quantify the catalytic activity of nanoparticles (Abou-Gamra 2014).

Therefore, in the present study, we have made an attempt on previously synthesized silver and platinum nanoparticles by plant extracts was used to determine the catalytic reduction of 4-Nitrophenol and assessment of catalytic reduction of organic dyes.

1.2 Materials and Methods

1.2.1 *Catalytic Activity in the Sodium Borohydride Reduction of 4-Nitrophenol*

An aliquot of 4-Nitrophenol (0.2 and 1 mM) was treated with freshly prepared sodium borohydride solution (3.6 and 16.5 mM) and 0.2 mL freshly prepared colloidal Ag NPs and Pt NPs in a 1 cm quartz cuvette. The reaction mixture was shaken thoroughly, and the UV-visible spectrum was recorded at room temperature (23–27 °C). The progress of the reaction was monitored by recording the absorption intensity of 4-nitrophenolate ion and the intensity of 4-aminophenol with certain time intervals, and the spectrum was recorded.

1.2.2 Assessment of Catalytic Activity in Reduction of Typical Dyes

The organic dyes such as Alizarin Red 1 ml of 7×10^{-4} M and Brilliant Blue-G 1 ml of 3×10^{-4} M was prepared. Similarly, 0.001 M NaBH_4 and 1 M H_2O_2 was prepared. Six test tubes were collected, washed thoroughly and labeled as D + NB, D + NB + SNP's, D + HP, D + HP + SNP, D + NB + Pt NP's, D + HP + Pt NP. (D–Dye, NB–Sodium borohydride, SNP's–Silver nanoparticles suspension, Pt NP's –Platinum nanoparticles suspension, HP – Hydrogen peroxide). In each test tube was added 1 ml of labeled material and made the volume up to 4 ml in all test tubes. The degradation of Alizarin Red and Brilliant Blue–G dyes by NaBH_4 and H_2O_2 was monitored in the presence or absence of platinum and silver nanoparticles spectrophotometrically.

1.3 Results and Discussion

1.3.1 Catalytic Reduction of 4-Nitrophenol

In this study, green route synthesized silver and platinum nanoparticles by plant extracts was used to determine the catalytic reduction of 4-Nitrophenol as model and assessment of catalytic reduction of typical organic dyes. The Fig. 1.1 shows the UV–visible spectrum of 4-nitrophenol mixed with deionized water. As compared with Fig. 1.1a the UV–visible spectrum at Fig. 1.1b indicates the formation of nitrophenolate ion while sodium borohydride was mixed with nitro phenol. When the sodium borohydride was added with aqueous solution of 4-nitrophenol, the absorption maxima of the solution was shifted from 319–401.5 nm due to the formation of more phenolate ions. Without the addition of nanoparticles there will be no reduction of nitro group to amino group (Rauf et al. 2005). Once the nanoparticles were added the intensity of peak at 401 nm was decreased and there was increase in intensity at 319 nm which indicates the formation of aminophenol group (Fig. 1.2a and b).

1.3.2 Assessment of Catalytic Activity in Reduction of Typical Organic Dyes

The rate of degradation of alizarin red-S dye by sodium borohydride and hydrogen peroxide were monitored in the presence and absence of nanoparticles spectrophotometrically. The Fig. 1.3a and b shows the catalytic degradation of organic dye Alizarin red-S by sodium borohydride and hydrogen peroxide respectively in the presence and absence of nanoparticles. In both the reduction profiles, the addition of nanoparticles

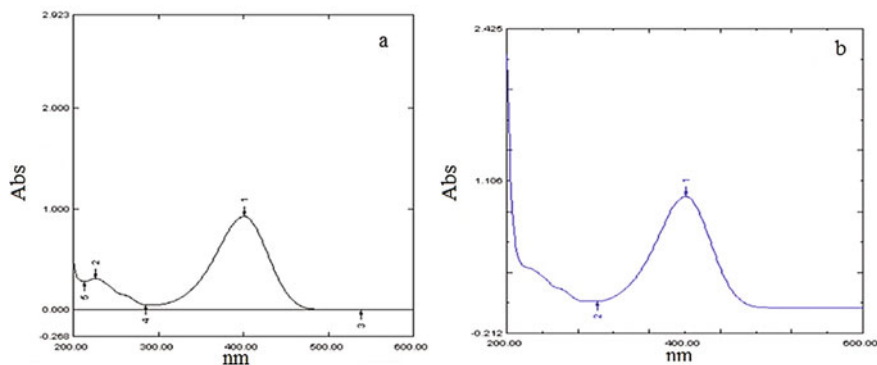


Fig. 1.1 UV–visible spectrum showing the peak for **a** 4-Nitrophenol **b** 4-Nitrophenol mixed with sodium borohydride (nitrophenolate ion)

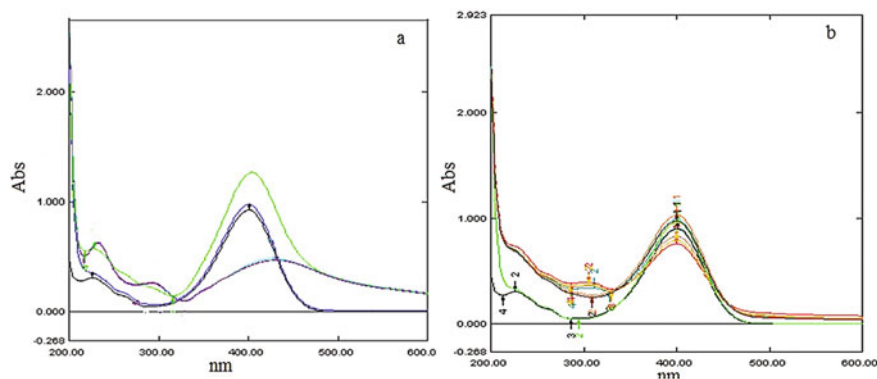


Fig. 1.2 Overlay of UV–visible spectrum at various time intervals using **a** SNP's **b** Pt NP's during catalytic reduction

enhanced the degradation of dye and have desirable difference between the initial and final dye concentration. From the reduction profile, the platinum nanoparticles have more catalytic activity in degradation of alizarin red – S compared to the silver nanoparticles.

Similarly, the rate of degradation of brilliant blue–G dye by sodium borohydride and hydrogen peroxide were monitored in the presence and absence of nanoparticles spectrophotometrically. The Fig. 1.4a and b shows the catalytic degradation of organic dye Brilliant blue–G by sodium borohydride and hydrogen peroxide respectively in the presence and absence of nanoparticles. In both the reduction profiles, the addition of nanoparticles enhanced the degradation of dye and have desirable difference between the initial and final dye concentration. When compared to alizarin red degradation, the silver nanoparticles show enhanced catalytic activity for the reduction of brilliant blue–G in combination with both reducing agents.

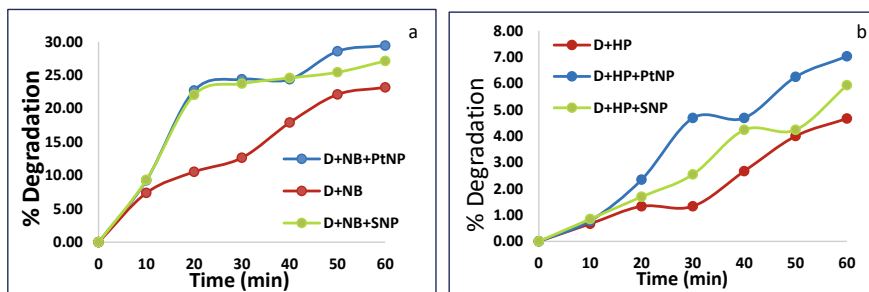


Fig. 1.3 **a** Sodium borohydride (NB) and **b** Hydrogen peroxide (HP) reduction profile of alizarin red-S in presence and absence of nanoparticles (Pt NP's, SNP's)

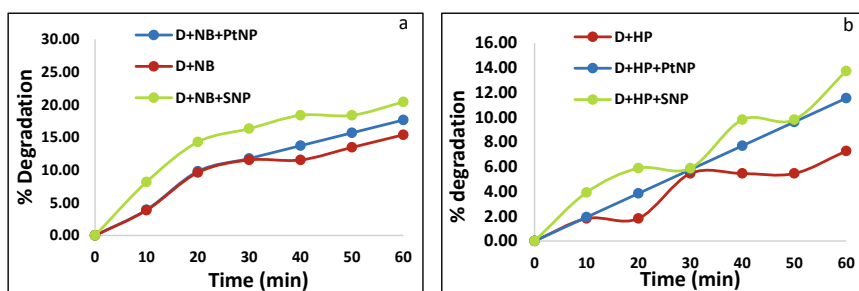


Fig. 1.4 **a** Sodium borohydride (NB) and **b** Hydrogen peroxide (HP) reduction profile of Brilliant blue-G in presence and absence of nanoparticles (Pt NP's, SNP's)

1.4 Conclusion

In this study, previously synthesized silver and platinum nanoparticles were successfully used for reduction of 4-Nitrophenol and assessment of catalytic activity for alizarin red-S and brilliant blue-G. The obtained results show that the nanoparticles have significant ability to catalyze the reduction reactions. Hence, these nanomaterials can be used as catalyst for various industrial dye degradation.

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Chapter 2

Biological and Eco-Friendly Cost-Effective Measure for Remediation



Anil Kumar Moola , **Selvam Sathish, S Mari Selvam, Balasubramanian Paramasivan, Sujatha Peela, Harish Kumar Seenivasan, and Dhandapani Gurusamy**

Abstract The natural environment is being spoiled with harmful chemicals from tannery effluent, industrial effluent, and domestic wastewater by human uses. Therefore, the avoidance of pollution and the degradation of harmful chemical compounds are essential for the future mankind. Several microorganisms are involved in the biodegradation of harmful contaminants in the soil including bacteria, fungi, and algal species. Phytoremediation is the best choice for many developing and developed nations to as it is an eco-friendly process for the decontamination of several environmental pollutants. Green plants absorb the wastes materials in the soil through their roots. Nowadays, *in-vitro* studies on the phytoremediation of emerging pollutants are giving better results. This chapter focuses on the recent studies on phytoremediation through various roots, including hairy roots, adventitious roots, and normal roots. It also reports the mechanism of biodegradation and enzymatic reactions that

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trigger the phytoremediation process. This information is essential for assessing the feasibility of a plant for the phytoremediation process before its application in the target site.

Keywords Biodegradation · Harmful chemicals · Microorganisms · Plant · Phytoremediation · Soil

2.1 Introduction

Nowadays, the global nations were having a challenging problem with the need for groundwater. But, the groundwater is rapidly polluted by urbanization, use of pesticides, heavy metals (HMs), tannery effluents, phenols, organic matter, explosives, azo dyes, and industrial wastes (Carolin et al. 2017). From the industrial wastewater, large quantities of heavy metals are released, such as Hg (II), Pb (II), Cd (II), Ag (I), Ni (II), and Cr (VI). These are highly toxic, non-degradable, and have infinite lifetimes (Mokarram et al. 2020). The harmful chemical compounds are synthesized in two different ways, including natural elements and industrial wastewater. Natural compounds can easily be degraded by external factors than industrial wastewater. In this sense, many biological methods can remove these harmful substances from industrial wastewater and contaminated wastewater (Crini and Lichtfouse 2019). In removing toxic compounds from soil and water, various methods are followed like adsorption on activated carbon, chemical oxidation, incineration, and solvent extraction. These methods have several disadvantages, including low efficiency, high cost, or by-product generation. Therefore, phytoremediation is the most valuable method to remove harmful chemical compounds from soil and wastewater (González et al. 2006; Al-Baldawi et al. 2018). In this technology, the root system of the green plants absorbs the harmful chemicals, while emerging pollutants are also removed from the wastewater simultaneously. Phytoremediation is classified based on the mechanism of action employed by the plants as follows: phytoextraction (accumulation of toxic chemicals and elimination of toxicity), rhizofiltration (root system for removing toxic chemicals from polluted water), phytostabilization (removal of bioavailability of toxic chemicals in the soil), phytovolatilization (evaporation of several toxic chemical compounds) (Antoniadis et al. 2017). Various strategies and processes are involved in phytoremediation for improving the uptake of heavy metals, such as genetic strategies (genetic modification of plants) and chelate-assisted strategies (Sharma 2021). Hairy root culture is one of the most valuable phytoremediation processes to uptake the toxic compounds from soil and water. It can grow faster in a toxic environment than the regular root system (Gujarathi et al. 2005; Perotti et al. 2020). Microorganisms also help in the process of phytoremediation as bacterial and fungal pathogens in hyperaccumulator plants. Based on the 'literary reports, the review on comprehensive discussion on synergistic mechanism of plants, microbes and other additives has been rarely reported.

Hence in this chapter, features and the enzymatic reactions that take place in plants and microbes during the process of phytoremediation has been described. The key concepts of plant, microbe and biochar assisted remediation has been annotated. This chapter helps the researchers to understand the benefits of phytoremediation and its significance on synergistic relation with bacteria and other additives.

2.2 Bio-based Technologies

In recent days, the world faces various problems owing to pollution of oil, air, water, and land. These problems are temporarily solved with several bio-based technologies like bioaugmentation, biostimulation, composting, bioventing, and biosparging. Bioremediation is the process of removing contaminants from polluted soil or water with the help of several processes including bioaugmentation and biostimulation (Tyagi et al. 2011; Lellis et al. 2019). The types of bioremediation techniques has been illustrated in Fig. 2.1 (Sharma 2020). Biostimulation is one of the bio-based technologies to promote the degrading rate of hazardous chemical compounds from contaminated soil. In this technology, low-energy lasers are used to stimulate the rate of wound healing and degradation of contaminants (Wasewar 2022). Bioaugmentation is another most important method in bio-based technology for bioremediation of polluted environments and it is one of the best methods to degrade pollutants than other methods. Two different processes are involved in this method including the addition of a pre-adapted pure bacterial strain and the addition of a transformed gene vector into indigenous microorganisms (Li et al. 2018; Huang et al. 2020). Bioaugmentation is successful in agricultural land, forestry, and wastewater treatment with the help of indigenous microorganisms. Bioaugmentation has been reported significant achievement in anaerobic digestion of lignocellulosic residues (Tsapekos et al. 2017), waste activated sludge (Cayetano et al. 2021), food waste (Jiang et al. 2020) to enhance the process with the help of microbial enrichment culture. In bioaugmentation, microbial communities play a major role in remediation. Microbial strain selection and their subsequent survival and activity determine the success rate of bioaugmentation. Hence the basic or initial and most important step of bioaugmentation is strain selection due to the involvement of microorganisms in fulfilling the whole process. For instance, Yang et al. (2019) reported that both hydrogenotrophic and acetoclastic methanogenic pathway has to be accounted for bioaugmentation where the former and latter showed 71.1 and 59.7% higher efficiency than control. In addition to microbial strains, the efficiency of the plant in bioaugmentation is determined based on several factors including the chemical nature and concentration of pollutants, reactor setup and physicochemical characteristics of the environment (El Fantroussi and Agathos 2005; Yang et al. 2019). Several new approaches were developed in bioaugmentation including bioaugmentation with nanomaterials comprehensively defined as nanobiohybrids (Guo et al. 2020), cells encapsulated/immobilized performing more efficiently compared to free system, gene modified bioaugmentation (Nzila et al. 2016), rhizosphere and plant assisted augmentation

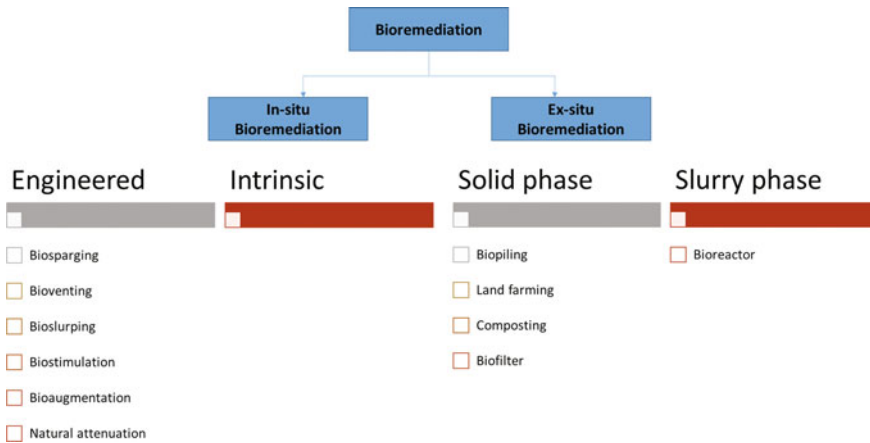


Fig. 2.1 Types of bioremediation techniques

(Itusha et al. 2019; Viji et al. 2022). Phytoremediation is one of the best techniques in bioaugmentation to remove pollutants from green plants with help of microorganisms. Microbial-enhanced sugarcane (*Saccharum officinarum*) was successfully eliminated the lindane from doped soil (Salam et al. 2017). Phytoremediation has various applications including degrading the pollutants, stabilizing the soil condition, increasing the biomass, decreasing the carbon dioxide level, and increasing the oxygen level. In the phytoremediation process, the plant roots thoroughly uptake the pollutants with the help of microorganisms in the rhizosphere. The pollutants then translocate into the upper regions of the plant for nutrient utilization and the remaining pollutants evaporate into the atmosphere through transpiration (Weyens et al. 2009; Ekta and Modi 2018).

2.3 Phytoremediation—An Overview

The term phytoremediation (Latin word) indicates the process of removal of the toxic contaminants by plant roots from the environment (Etim 2012). Generally, phytoremediation is defined as cleaning the polluted environment with natural plants. In phytoremediation, the green plants reduce the number of toxic compounds from the environment with root-associated soil microbes (Ali et al. 2013). Phytoremediation checks out the pollution of the contaminated environment through a series of processes namely phytofiltration, phytostabilization, phytoextraction, phyto-volatilization, and phytotransformation, rhizosecretion. The detailed mechanism has been represented schematically in Fig. 2.2. Its mechanism includes various processes such as heavy metal uptake, metal translocation to shoots, heavy metal tolerance, exclusion of toxic compounds, vacuolar compartmentalization, production of phytochelatins, and metallothioneins (Sharma 2021). In phytoremediation,

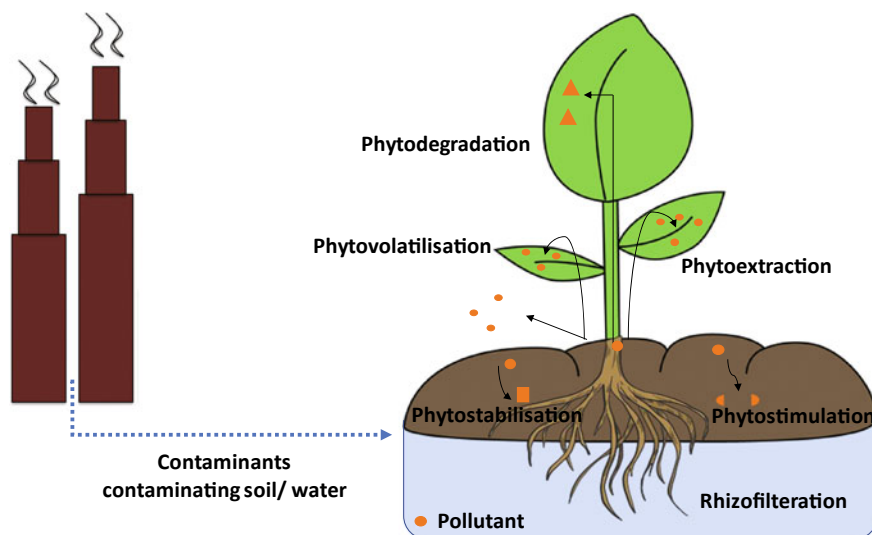


Fig. 2.2 General mechanism involved in phytoremediation by using plant roots

bioabsorption has several significant steps, including chemisorption, complexation, adsorption on surface and pore, ion exchange, micro precipitation, hydroxide condensation onto the surface (Shinomol & Bhanu 2021). Several factors including redox potential, pH, soil organic matter, clay content, nutrient balance, cation exchange capacity (CEC), concentrations of other trace elements in soil, soil moisture and temperature influences the uptake of metal from the contaminated soil (Neilson and Rajakaruna 2015). In translocating pollutants, the plant system has different tolerance processes including sequestration/ compartmentalization, binding/ chelation, excretion from aerial plant components, enzymatic and non-enzymatic antioxidants, and protection, stress recovery, and damaged protein repair (Antoniadis et al. 2017). Particularly, persistent organic pollutants have gradually become one of the most critical compound pollutants in the world due to their long-term bioavailability. Specific attention is paid to polycyclic aromatic hydrocarbons, polybrominated diphenyl ethers, organochlorine pesticides, and halo hydrocarbons.

The phytostimulation is the process of promoting microbial biodegradation in the rhizosphere. Phytotransformation is one of the most important metal absorption processes in the degradation of harmful pollutants comprising the degradation, decolorization and transformation to less toxic compounds. The phytotransformation of methylene blue from water using *Azolla pinnata* has been found to show decolorization efficiency of 85% removal validating the fact of ability of plants to transform organic compounds (Al-Baldawi et al. 2018). The taller plants have two different strategies for adsorbing heavy metals including exclusion (uptake of pollutants from soil to shoot through root) and accumulation of toxic compounds (detoxification of toxic compounds) (Luo et al. 2016). The accumulators can accumulate the harmful chemical compounds in the aerial parts of plants and exclude

the accumulation of toxic compounds in plants' biomass. Metal uptake and translocation of toxic compounds in the plant cell membrane also involves proton pumps (ATPases that consume energy and generate electrochemical gradients), co- and anti-transporters (proteins that use the electrochemical gradients generated by ATPases to drive the active uptake of ions), and channels (proteins that facilitate the active uptake of ions). Some organic acids also enhance the uptake of metal ions from contaminated soil. Heavy metal contaminated soil treated with citric acid assisted *Brassica napus* in hydroponic method reduced significant amount of cadmium, hydrogen peroxide (H_2O_2), electrolyte leakage, malondialdehyde (MDA) accumulated in plant roots (Ehsan et al. 2014).

Nowadays, air pollution is a primary challenge for all developed nations due to urbanization, vehicles, and industrialization. This makes the ozone in the surrounding environment depleted. Various methods were discovered to remove the toxic gases (including carbon monoxide, carbon dioxide, and greenhouse gases) from the natural air. Phytoremediation is one of the best methods to remove toxic gases from a contaminated environment. *Sida hermaphrodita* is one example of the phytoremediation of toxic gas through absorption from fossil-fuel combustion (Uchman et al. 2017).

2.3.1 Plant Roots in Phytoremediation

The harmful contaminants in the environment include heavy metals and complex chemicals that the plants directly uptake by their roots and few specific plants limit the uptake amount of heavy metals such as arsenic since its toxicity limit is 40 and 200 mg /kg in sandy and clay soils. The plants involved in phytoremediation adopt different mechanisms including passive uptake through the apoplast, direct transcellular transport from the environment to the plant vascular system, and active uptake through the symplast by plants during heavy metal uptake. The uptake of trace metal elements is controlled by root factors, including soil acidification by root exudates, translocator activity and selectivity, root membrane activity, avoidance strategy mechanisms, the release of redactors or oxidants, chelator root excretion, phytosiderophores, acids, and hydrogen ions (Vithanage et al. 2012).

Rhizosecretion is one of the essential roles of plant roots involved in phytoremediation. In this process, the valuable natural products, including enzymes, alcoholic compounds, and other substances, were secreted by plant roots (Takkar et al. 2022). The plant roots exudate compounds with specific biological activity in low quantity. Plant exudates are in multiple forms of sugar, alcohol, and acids. The exudates are annually produced ranging from 10–20% by photosynthetic plants. The exudates level is identified by the chemical or chromatographic method due to the low amount of secreted compound. The secretion of exudates is affected by various factors, including plant species, age of the plant, temperature, light, plant nutrition, soil microorganism, medium composition, soil moisture, and root damage. The primary source of exudates is the tip of the root (Antoniadis et al. 2017). Some of the essential

enzymes including laccases, dehalogenase, nitroreductases, nitrilases, and peroxidases were secreted by the plant for degrading the organic chemical compounds. The xenobiotic compounds are also degraded by the exudates of plants (Hussain et al. 2018). The rhizospheric microorganisms help to improve the uptake of metal by their enzymatic action on these metals (Kumar et al. 2017). With root exudates, microbial biodegradation is stimulated to minimize the toxicity of the chemicals. The microbial interaction of plants was highly influenced in phytoremediation to remove the pollution from contaminated soil and water. The plant growth-promoting rhizobacteria promote nitrogen fixation, phytohormone production, specific enzymatic activity, and plant protection (Moola et al. 2021). The energy is more important for uptake of the base metals (Ca^{2+} , Mg^{2+} , K^+ , and Na^+) through the active transport mechanism. However, organic chemicals can easily be diffused into the plant root cells by a passive transport mechanism. If the substrate concentration is low, the degradation can be explained by Michaelis–Menten kinetics. The first-order reaction is considered, while the higher substrate concentrations are limited. According to Monod kinetics, bacteria can be used as a substrate for development, and the growth of substrates is better at higher availability. However, it was noted that bacteria have limited degradation capability if the substrate's concentration is minimal, which can prohibit the biodegradation of contaminated sites. Combining with the phytoremediation potential of plants, bacteria can therefore be used to overcome problems with environmental pollutants. (Hussain et al. 2018). Examples of some plant species and their root system utilized for enhanced remediation of inorganic and organic pollutants are depicted in Table 2.1.

2.3.2 Adventitious Roots in Phytoremediation

The adventitious roots develop from various tissues, cells, and neighboring cells of vascular tissues. Besides, they develop in the postembryonic stage of plant growth (Geiss et al. 2018). Plants generally adapt to their native natural environment, including water availability and micro- and macronutrient availability, concentration, and localization. Hence, the plants change their organ and tissues depending upon the environmental conditions. The adventitious roots are also a modification of the shoot by the plant's adaptation mechanism (Bellini et al. 2014). The growth of adventitious roots is inhibited when an excess amount of heavy metal is accumulated in the surrounding soil environment. The inhibition is high when the solute potential of the external medium is high that may deny the plant from nutrient uptake. Direct inhibition of enzymes that are of physiological importance, and inhibition of mitotic division of the meristematic cells may also occur (Odjegba and Fasidi 2004). *ThMT3* gene expression in the transgenic *Salix matsudana* increases the tolerance of copper stress and nitric oxide (NO) production (Yang et al. 2015). A high concentration of copper (5–500 mM) causes leaf toxicity, strongly affects adventitious root formation and produces ethylene at higher levels in *Populus alba*'s shoot culture (Franchin et al. 2007). The aquatic adventitious roots have the photosynthetic ability

Table 2.1 Examples of the plant species and their root system utilized for enhanced remediation of inorganic and organic pollutants

S.no	Plant species	Type of root system	Uptake elements	References
1.	<i>Armoracia rusticana</i>	Hairy root system	Uranium	Soudek et al. (2011)
2.	<i>Pteris vittata</i>	Hairy root system	Arsenic and chromium	Kalve et al. (2011)
3.	<i>Eleocharis acicularis</i>	Rhizome	Cu, Zn, As, and Cd	Sakakibara et al. (2011)
4.	<i>Brassica juncea</i> L	Hairy root system	Zinc and nickel	Ismail et al. (2012)
5.	<i>Nicotiana tabacum</i>	Hairy root system	Arsenic	Talano et al. (2014)
6.	<i>Brassica napus</i> HR and <i>Pantoea</i> sp. FC 1	Hairy root system	Chromium	Ontanon et al. (2014)
7.	<i>Sorghum bicolor</i> × <i>sudanense</i>	Adventitious roots	As, Cu, Pb and Zn	Babu et al. (2014)
8.	<i>Ipomoea aquatica</i>	Adventitious roots	Copper	Chanu and Gupta (2014)
9.	<i>Nicotiana tabacum</i>	Hairy root system	2,4-Dichlorophenol	Angelini et al. (2014)
10.	<i>Nicotiana tabacum</i>	Hairy root system	Copper	Perez-Palacios et al. (2015)
11.	<i>Solanum nigrum</i> L	Tap-root	Cadmium	Ji et al. (2015)
12.	<i>Alyssum bertolonii</i>	Hairy root system	Nickel	Ibañez et al. (2016)
13.	<i>Hyptis capitata</i>	Hairy root system	Copper	Malik et al. (2016)
14.	<i>Adenophora lobophylla</i> <i>A. potaninii</i>	Hairy root system	Cadmium	Malik et al. (2016)
15.	<i>Moringa oleifera</i> <i>Typha latifolia</i> <i>Cymbopogon proximus</i>	Hairy root system	Cadmium	Ghada et al. (2017)
16.	<i>V. zizanioides</i>	Hairy root system	Arsenic	Moogouei (2018)
17.	<i>Vetiveria zizanioides</i>	Hairy root system	Nitrate	Moogouei (2018)
18.	<i>Amaranthus chlorostachys</i>	Hairy root system	Metformin and cesium	Moogouei (2018)
19.	<i>Elodea canadensis</i>	Adventitious roots	Cobalt	Mosoarca et al. (2018)
21.	<i>Lemna minuta</i> Kunth	Hairy root system	Cr(VI)	Paisio et al. (2018)
22.	<i>Brassica napus</i>	Hairy root system	Chromium	Perotti et al. (2020)

through the uptake of oxygen microelectrode and CO₂. The adventitious root system of *Meionectes brownie* has the maximum rate of photosynthesis (0.38 μmol O₂ m⁻²s⁻¹) (Rich et al. 2011). The function of the adventitious root system is based on the location of adventitious roots in plants. If the adventitious root generates from the stem, they help in tolerating the biotic stresses (Steffens and Rasmussen 2016). When there is flooding, it affects the plant that produces the aquatic adventitious roots due to the absorption of air (Ayi et al. 2016).

2.3.3 Hairy Roots in Phytoremediation

The hairy root (HR) system of the plants plays a major role in phytoremediation during the metal uptake. Generally, HRs have fast growth and development than normal root systems in natural plants. It have potential energy in the phytoremediation process (Majumder and Jha 2012). The root system has several processes to remove the pollutants, including adsorption, transport, translocation, hyperaccumulation or transformation, and mineralization. The carrot's hairy root culture was eliminating more than 90% of exogenous phenolic compounds from contaminated culture medium within 120 h (Zhou et al. 2013). The hairy root culture of *Brassica napus* removed 97–98% of 2, 4-dichlorophenol within 1hr treatment (Agostini et al. 2003). The root system of *Vetiveria zizanioides* has reached a high amount of Pb (2280 mg kg⁻¹ DW) in 5000 mg Pb kg⁻¹ of contaminated soil (Chen et al. 2004). *Agrobacterium rhizogenes* strain ATCC 15,834 infected *Helianthus annuus* has fast-growing hairy root culture and that catalyzes the disappearances of antibiotics including tetracycline (TC) and oxytetracycline (OTC) from the aqueous medium (Gujarathi et al. 2005). The hairy root culture induced by *A. rhizogenes* (NCIM 5140) in explants of *Sesuvium portulacastrum*, removed the harmful chemicals (Reactive green 19A HE4BD) from textile effluents within 5 days incubation periods (Lokhande et al. 2015). The normal root system of *Canna indica* Linn removed 41–55% of triazophos (O, O-diethyl-O-(1-phenyl-1, 2, 4-triazole-3-base) sulfur phosphate, TAP) in a hydroponic system (Cheng et al. 2007). *Vetiveria zizanioides* removed the phenolic contents through hydroponics technology (Singh et al. 2008).

2.3.4 Remediation with Microorganisms

Few microorganisms could act as biofertilizers in polluted soil and water. Naturally, plants interact with microorganisms for improved nitrogen fixation. Recent developments in bioremediation approaches have been made during the last 20 years with the intention of restoring damaged areas rapidly and inexpensively (Balloi et al. 2010; Leong and Chang 2020). Examples of some microbes involved in heavy metal bioremediation has been summarized in Table 2.2. Remediation through bioaugmentation with *Bacillus* sp., *Lysinibacillus* sp., or *Rhodococcus* sp. diminished

Table 2.2 Example of microbes are mostly involved in degrading pollutants

S.no	Microbes	Pollutant	References
1.	<i>Kocuria flava</i>	Copper	Achal et al. (2012)
2.	<i>Stenotrophomonas sp.</i>	Copper	Zaki and Farag (2010)
3.	<i>Enterobacter sp.</i>	Copper	Bestawy et al. (2013)
4.	<i>Desulfovibrio desulfuricans</i>	Nickel	Ockjoo et al. (2015)
5.	<i>Enterobacter cloacae</i>	Nickel	Banerjee et al. (2015)
6.	<i>Acinetobacter sp.</i>	Zinc	Tabaraki et al. (2013)
7.	<i>Desulfovibrio desulfuricans</i>	Cadmium	Ockjoo et al. (2015)
8.	<i>Enterobacter cloacae</i>	Lead	Banerjee et al. (2015)
9.	<i>Pseudomonas aeruginosa</i>	Lead	Ahmady-Asbchin et al. (2015)
10.	<i>Termitomyces clypeatus</i>	Chromium	Fathima et al. (2015)
11.	<i>Trametes versicolor</i>	Zinc	Sahan et al. (2015)
12.	<i>Pleurotus mutilus</i>	Uranium	Mezaguer et al. (2013)
13.	<i>Pleurotus ostreatus</i>	Lead	Zhang et al. (2016)
14.	<i>Aspergillus lentulus</i>	Lead	Mishra and Malik (2012)
15.	<i>Candida parapsilosis</i>	Mercury	Muneer et al. (2013)
16.	<i>Bacillus cereus</i>	Cadmium	Jan et al. (2015)
17.	<i>Cupriavidus metallidurans CH34</i>	Chromium	Alviz-Gazitua et al. (2019)
18.	<i>Pseudomonas aeruginosa and Bacillus cereus</i>	Cadmium	Nath et al. (2018)
19.	<i>Pseudomonas aeruginosa</i>	Cadmium	Chellaiah (2018)
20.	<i>Bacillus megaterium</i>	Boron, lead, and cadmium	Esringu et al. (2014)
21.	<i>Bacillus licheniformis</i>	Mercury	Muneer et al. (2013)
22.	<i>Enterobacter cloacae</i>	Mercury	Al-Garni et al. (2010)
23.	<i>Enterobacter, Klebsiella, Leifsonia, Bacillus</i>	Zinc	Jain et al. (2020)
24.	<i>Rhizophagus clarus</i>	Mercury	Gontia-Mishra et al. (2016)
25.	<i>Trichoderma virens</i>	Cadmium	Khanna et al. (2019)

the metal contents of the HM-induced contaminated soils, according to Fauziah et al. (2017). The soil microbes have the efficiency of remediation to degrade the harmful chemical compounds. *Burkholderia fungorum* DBT1 is a soil microorganism that transforms dibenzothiophene, phenanthrene, naphthalene, and fluorine (Andreolli et al. 2013). *Lolium multiflorum* plant system degrades the petroleum hydrocarbons (Arabian medium crude oil) with the help of petroleum-degrading microorganisms including *Glomus intraradices* (AMF), *Sphingomonas paucimobilis* (bacterium),

and *Cunninghamella echinulata* (filamentous fungi). Phytoremediation of metals in maize plants is successful with the addition of *Bacillus licheniformis* and *Rhodotorula dairenensis* (García et al. 2013). The plant's endophytic microorganisms produce economically valuable substances and plant growth-promoting substances such as phytohormones, plant growth regulators, and host protecting molecules. Various endophytic genera were found in plants, including *Serratia*, *Enterobacter*, *Acinetobacter*, *Agrobacterium*, *Bacillus*, *Herbaspirillum*, and *Klebsiella* strains. The endophytic microorganisms can improve plant growth by enhancing nutrient acquisition, water uptake, and decreased oxidative stress enzyme activities in host plants grown in contaminated soils (Feng et al. 2017). The bacteria's resistance to heavy metals like zinc and copper will be evaluated based on biochemical investigations of the bacteria. Some biodegradative bacteria were degrading the toxic chemical substances like herbicides, pesticides, refrigerants, solvents, and other organic compounds to beneficial plant growth substances (Zhang et al. 2020). The soil microbes participate in various remediation processes, including soil formation, energy transfer, and nutrient cycling. Through these processes, the microorganisms can promote revegetation and increase the stability of polluted ecosystems. But a high concentration of pollutants reduces biomass, population number, and diversity (Li et al. 2012). Soil microorganisms and plant root-associated bacteria were enhancing the production of plant root exudates, including organic molecules. Heavy metal phytotoxicity can be decreased by altering the bioavailability of metals in soil and preventing metal translocation through phytoremediation (Ma et al. 2016). The plant-associated microorganisms have the capacity of absorbing pollutants and promote plant growth through several direct and indirect mechanisms. The plant-associated microorganisms can produce plant growth regulators, including auxins, cytokinins, and gibberellins (Weyens et al. 2015). Bacterial and fungal strains degrade petroleum hydrocarbons from contaminated soil. From this study, the maximum phytoremediation rate was reported as 0.51 mg of TPH g⁻¹ of dry plant d⁻¹ at 60 days of culture and 90% of hydrocarbon was removed from the contaminated soil (Escalante-Espinosa et al. 2005). *Lolium perenne* plant was inoculated with microbial strains of *Bacillus subtilis*, *Sphingobacterium multivolume*, *Acinetobacter radioresistance*, *Rhodococcus erythropolis*, and *Pseudomonas fluorescens* for remediation of petroleum-contaminated soil. From this study, the maximum degradation rate was achieved up to 58% after 162 days from inoculation (Tang et al. 2010).

2.4 Biochar Mediated Remediation

Biochar is gaining remarkable attention in bioremediation techniques over years owing to its significant physicochemical characteristics. The porous surface, functional groups, surface area and charge properties of biochar enhances its use as a stand-alone bioadsorbent for various pollutants viz organic, petroleum, dyes, heavy metals, etc. (Cheng et al. 2021). Being produced from biomass like agricultural residues, biosolids, municipal solid wastes, it is carbon rich product possessing the

heterogeneous characteristics mainly depending on feedstock and reaction conditions (Selvam and Paramasivan 2022). As concerned to the scope of this chapter, literary reports on biochar for bioremediation validate it to be cost-effective and sustainable bioadsorbent compared to commercial adsorbents. The effectiveness of its adsorption has been proved for antibiotics, hydrocarbons, pesticides and metal contaminants. The proposed mechanism for the biosorption is primarily upsurging the microbial community to degrade the contaminants in amended soils. In addition to that, biochar efficiently adsorbs contaminants like heavy metals and decreases its bioavailability and toxicity (Karppinen et al. 2017). Song et al. (2021) reported that bacteria immobilized on biochar in polycyclic aromatic hydrocarbon contaminated soil increased the removal rate of the contaminant compared with the individual applications. Bioremediation efficiency is improved significantly by the biochemical material especially the biochar being immobilized with plant growth promoting bacteria drives soil micro-ecology and enhances the remediation pathway with this accumulator system (Wu et al. 2019). The studies on utilizing biochar with phytoremediation report that amendment of biochar reduces heavy metal and metalloid bioavailability. However, few plants need elevated doses to accumulate the bioavailable metals and metalloids (Narayanan and Ma 2022). Summarizing these entries, advantages of biochar for bioremediation involve improved physicochemical properties of contaminated soil, enhanced microbial activities and population, and improved agricultural productivity.

2.5 Conclusion and Future Perspective

Using phytoremediation technology, heavy metal uptake by roots seems to be a prosperous way to remediate the polluted environment. Microbial remediation and phytoremediation are reliable, cost-effective, efficient, eco-feasible alternatives. When compared with other conventional methods, these methods are better biological tools to remediate polluted conditions. However, several factors must be considered to accomplish better remediation results. The future perspectives to enhance the research includes the following.

- The technologies providing low-cost solution and eco-friendly to remediate contaminates has to be explored further.
- Also, focus on developing agricultural techniques to enhance remediation efficiency to reduce the time and cost utilized in the removal of heavy metals from soil to a considerable level for the well-being of mankind and nature has to be enhanced.
- Exploring the plants that has higher biomass, environmental affordability and high heavy metal accumulation potential for overcoming the associated limitations of phytoremediation.
- Eco-friendly additives with phytoremediation for enhancing the process efficiency and simultaneous enhancement of soil structure has to be explored more.

- Most of studies so far reported is based on laboratory scale. Translative research on field demonstration and its execution has to be implemented.

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Chapter 3

Performance of Lead (Pb) Resistant *Staphylococcus aureus* from Tannery Waste Discharge Site



P. Muthukumaran and S. Karthika

Abstract Heavy metal resistant bacteria were identified from tannery effluent wastewater in and around Ezar Tanneries in Vaniyambadi, Vellore district, South India. In the first step, eleven bacterial isolates were collected from tannery effluent wastewater and tested on Luria Bertani (LB) agar plates enriched with 5 mg/l concentrations of five distinct heavy metal ions, including Cr, Cu, Ni, Pb, and Zn. Heavy metal resistance was a primary factor in choosing these two isolates. *Staphylococcus aureus* was determined to be the causative agent based on morphological and biochemical characteristics of the isolates. All of the detected isolates showed resistance to both copper and lead. Maximum growth of *Staphylococcus aureus*, a sewage isolate, occurred at 30 °C and pH 6.5. Bioremediation of heavy metal-tainted sewage and wastewater may benefit from the use of the discovered *Staphylococcus aureus* strain that is resistant to the metals in question.

Keywords Tannery effluent · Lead (Pd) · *Staphylococcus aureus* · Bioremediation

3.1 Introduction

Raw tannery effluent waste water contain huge amount of heavy metals that are not degraded by the conventional process of wastewater treatment and removal of these heavy metals from waste water needs advance chemical technology and it's expensive too. Presence of high concentration of toxic heavy metals in wastewater directly leads to both contamination of receiving water bodies and deleterious impact on aquatic life. Particularly, the mining industry generates residues with high levels of heavy metals (Ni, Cd, Pb, Mn, Cu, and Zn) that are deposited in open areas presenting an ecological and human health risk. Lead (Pb) a major pollutant that is found in soil, water and air is a hazardous waste and is highly toxic to human, animals, plants and

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microbes (Low et al. 2000; Elahi et al. 2019). The microorganisms respond to these heavy metals by several processes; including transport across the cell membrane, biosorption to the cell walls and entrapment in extracellular capsules, precipitation, complexation and oxidation–reduction reactions (Rai et al. 1981; Macaskie and Dean 1989; Huang et al. 1990; Brady and Duncan 1994a; Brady et al. 1994b; Elahi et al. 2022). Naturally occurring bacteria that are capable of metal accumulation have been extensively studied since it is difficult to imagine that a single bacterium could be capable to remove all heavy metals from its polluted site (Clausen 2000). The objective of this study is to determine heavy metals resistance of bacteria, biochemical characteristics and growth studies were used to exploit these isolates for clean-up of tannery effluent waste water.

3.2 Materials and Methods

3.2.1 Isolation of Screening of Bacterial Isolates from Tannery Effluent

The sewage water samples were collected in and around Tanneries, Vaniyambadi, Vellore district, South India. The samples were collected in sterile plastic container and transport to laboratory for bacteriological analysis. The bacterial isolates were screened on Luria Bertani (LB) agar plates supplemented with 5 mg/l concentration of Cr, Cu, Ni, Pb and Zn metals by the standard pour plate method. Plates were incubated at 37 °C for 48 h and colonies differing in morphological characteristics were selected and used for further studies (Elahi et al. 2019, 2022).

3.2.2 Identification and Biochemical Characterization of Screened Bacterial Isolates

The shape and color of the colonies were examined under the microscope after Gram staining. Isolates were biochemically analyzed for the activities of catalase, starch hydrolysis and gelatin hydrolysis, mannitol test, motility, indole production and citrate utilization. The tests were used to identify the isolates according to Bergey's Manual of Systematic Bacteriology (Claus and Berkeley 1986).

3.2.3 Growth Studies of Screened Bacterial Isolates

Growth studies of sewage bacterial isolates was studied in 250 ml flasks containing 50 ml LB medium supplemented with 0.1 mM concentration lead. Flasks were inoculated with 0.5 ml of overnight culture and agitated on a rotary shaker at 37 °C. Growth was monitored as a function of biomass by measuring the absorbance at 620 nm using spectrophotometer and the optimal growth conditions with reference to pH and temperature were determined. The effect of pH on the growth of bacteria in the presence of lead was analyzed at five pH values 4.5, 5.5, 6.5, 7.5 and 8.5 and different temperatures 30, 37 and 45 °C was studied (Elahi et al. 2022).

3.2.4 Determination of MIC

Maximum resistance of the bacteria against increasing concentrations of lead on LB broth was evaluated until the strains unable to grow in the broth. The initial concentration used (100 ppm) was added from 100 mg/100 ml of stock solution. Concentration of metal up to 3000 mg/l was taken. The stock solution of Pb (NO₃)₂ (Ranbaxy) was prepared in distilled water and sterilized by autoclaving at 121 °C, 15 psi for 15 min (Kalaimurugan et al. 2020).

3.3 Results and Discussion

3.3.1 Isolation of Screening of Bacterial Isolates from Tannery Effluent

The microbial colonies were isolated by pour plate method. Eleven bacterial isolates were purified using streak plate method. The bacterial isolates named based on the source from which it was isolated. The isolated bacterial colonies were screened with the metal concentration of 100 mg/l and the metal resistant isolates were isolated based on their cell density values measured at 620 nm by using UV–visible spectrophotometer (ELICO, India) (Kalaimurugan et al. 2020) (Table 3.1).

3.3.2 Identification and Biochemical Characterization of Screened Bacterial Isolates

From the gram-staining procedure it is found that the microbial strain was a gram-positive spherical bacterium that occurred in microscopic clusters formed yellow

Table 3.1 Screening of metal resistant bacterial isolates

S.No	Metals (5 mg/lit)	Days	Microbial strains															
			VLB*-1	VLB-2	VLB-3	VLB-4	VLB-5	VLB-6	VLB-7	VLB-8	VLB-9	VLB-10	VLB-11					
1.	Cr	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	
		1	+	+	+	+	++	+	+	+	++	+	+	++	+	+	+	
		2	++	+	+	+	++	++	++	+	+	++	++	++	++	++	++	++
2.	Cu	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
		1	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+
		2	+	+	+	++	+	+	+	+	+	+	+	+	++	+	+	+
3.	Ni	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
		1	+	+	+	+	+	+	+	+	++	+	+	+	+	+	+	+
		2	+	+	+	+	+	++	+	+	++	+	+	++	++	++	++	++
4.	Pb	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
		1	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+
		2	++	++	+	+	++	++	++	++	++	++	++	++	++	++	++	++
5.	Zn	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
		1	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+	+
		2	+	++	+	+	++	++	+	+	++	++	++	++	++	++	++	++

Note No growth; + poor growth; ++ moderate; +++ growth

*VLB-Vaniyambadi Leather effluent Bacteria

Table 3.2 The morphological and biochemical characterization of bacterial isolates (VLB-9)

Sl. No	Characteristics	<i>Staphylococcus aureus</i> (VLB-9)
1.	Colony diameter	1–3 mm
2.	Colony color	Yellow
3.	Colony morphology	Slightly elevated, circular
4.	Gram staining	Positive
5.	Turbidity in broth	Uniform dispersed growth
6.	Motility	Non-motile
7.	Oxidase	Negative
8.	Catalase	Positive
9.	Mannitol sugar fermentation	Negative
10.	Coagulase	Positive
11.	Temperatures	
	30	+ + +
	37	+ +
	45	–
12.	pH	
	4.5	+
	5.5	+ +
	6.5	+ + +
	7.5	+ +
	8.5	–

Note No growth; + poor growth; + + moderate growth; + + + growth

colonies in nutrient medium. Further biochemical tests resulted positive for catalase, mannitol sugar fermentation and coagulase test confirmed that the isolated microbial strain VLB-9 was identified as *Staphylococcus aureus* (Elahi et al. 2022) (Table 3.2).

3.4 Growth Studies of *Saphylococcus Aureus*

3.4.1 Effect of Temperature

The measurements from the cultures incubated for 24 h were in good agreement according to bacterial resistance. Growth was monitored as a function of biomass by measuring the absorbance at 620 nm using spectrophotometer and the optimal growth conditions with reference to different temperature such as 30, 37 and 45 °C. Figure 3.1 shows maximum removal of the metal was found to be at 30 °C. Thus

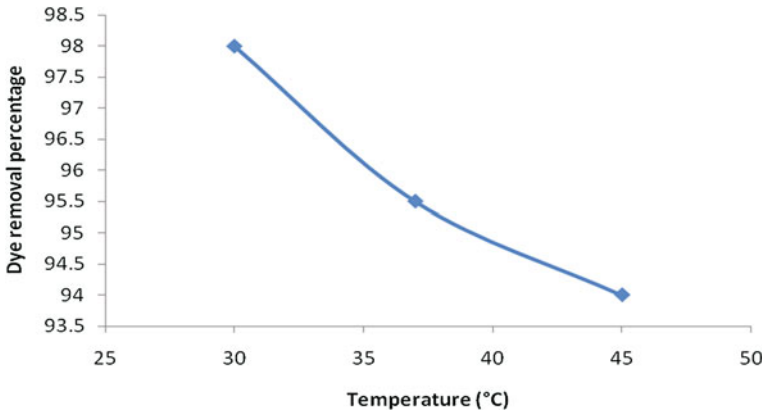


Fig. 3.1 Effect of temperature on biosorption of lead by *staphylococcus aureus*

the optimum temperature was found as 30 °C (Raja et al. 2006; Kalaimurugan et al. 2020).

3.4.2 Effect of pH

The effect of pH on the growth of bacteria in the presence of lead was analyzed at five pH values such as 4.5, 5.5, 6.5, 7.5 and 8.5. Figure 3.2 shows maximum removal of the metal was found to be at 6.5 pH. Thus the optimum pH was found as 6.5 (Ahmed et al. 2005; Moodley et al. 2019).

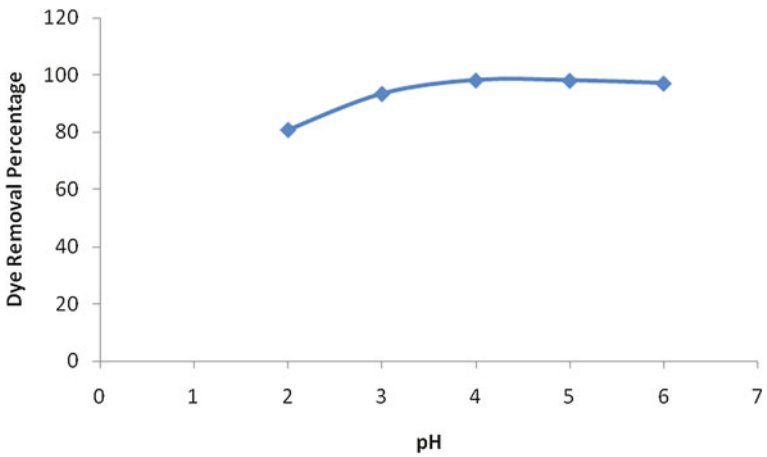


Fig. 3.2 Effect of pH on biosorption of lead by *staphylococcus aureus*

3.4.3 Determination of Minimum Inhibitory Concentration (MIC)

Staphylococcus aureus was highly resistant for various concentration of metal. The concentration such as 100, 500, 1000, 1500, 2000 and 2500 mg/l. Visible growth was observed in the broth containing up to 1000 mg/l concentration of lead as shown in Fig. 3.3 and Table 3.3. The growth of the isolated bacteria was inhibited from 1500 mg/l concentration of metal. Thus the minimum inhibitory concentration of the isolated bacteria was found as 1500 mg/l (Muthukumaran et al. 2014; Filali et al. 2004; Marzan et al. 2017).

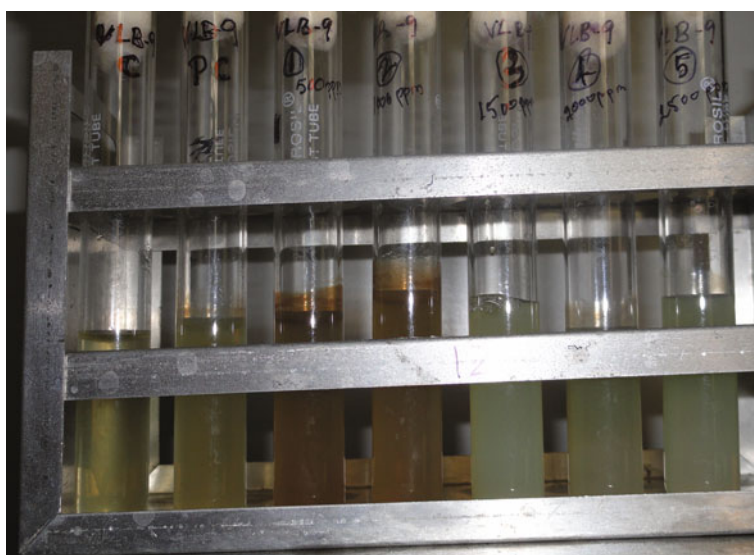


Fig. 3.3 Minimum inhibitory concentration of *staphylococcus aureus*

Table 3.3 Determination of minimum inhibitory concentration (MIC)

Sl. No	Metal ion concentration (ppm)	Growth of <i>Staphylococcus aureus</i>
1.	100	+++
2.	500	+++
3.	1000	++
4.	1500	—
5.	2000	—
6.	2500	—

Note No growth; + poor growth; ++ moderate growth; +++ growth

3.5 Conclusion

The heavy metal resistant organism could be a potential agent for bioremediation of heavy metals pollution. *Staphylococcus* resistant to heavy metals and antibiotics. In the present study high degree of heavy metals resistance associated with multiple antibiotic resistances was detected in *Staphylococcus aureus*. *Staphylococcus aureus* exposed to high levels of heavy metals in their environment have adapted to this stress by developing various resistance mechanism. These mechanisms could be utilized for detoxification and removal of heavy metals from polluted environment. According to these results, the present study evaluates that *Staphylococcus aureus* were used to remediate heavy metal contaminated tannery effluent waste.

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Chapter 4

Characterization of Chlorpyrifos Degrading *Pseudomonas* sp. Isolated from Contaminated Soil



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Abstract Chlorpyrifos is chemically known as O,O- diethyl-O-(3,5,6-trichloro-2-pyridyl) phosphorothioate belongs to organophosphorous pesticide. Usage of pesticide is still problematic and cause drastic changes in the ecosystem due to its moderate toxicity. The soil contaminated with chlorpyrifos was used to isolate CP degrading bacteria by standard pure culture technique. After isolation, the identified isolates were *P. aeruginosa* and *P. fluorescens* and their pure cultures stored for further degradation studies. In nutrient broth and MSM medium, the growth of bacteria was observed by turbidometric measurements with different concentration of pesticides. The cultures of chlorpyrifos degrading bacteria were centrifuged for 20 min at 5000 rpm and the supernatants were collected. The collected supernatants subjected to analysis of UV-Vis spectrophotometer at 300 nm to find out the maximum absorption spectra. The control was used as a blank. *P. aeruginosa* and *P. fluorescens* has the range of different values at different concentration of pesticides when absorbed at the maximum spectra of 300 nm. In *P. aeruginosa*, the chlorpyrifos residue was recorded in the 0.5c followed by 1.0c, 1.5c, 2.0c and 2.5c. The highest chlorpyrifos residue was obtained in 0.5c and 1.0c. In *P. fluorescens*, the highest chlorpyrifos was recorded in the 0.5c, followed by 1.03c, 1.5c, 2.0c and 2.5c. The highest chlorpyrifos residue was obtained in 0.5c and 1.0c. After the degradation studies by UV-Vis spectrophotometer analysis, comparatively *P. aeruginosa* have shown the higher degradation of chlorpyrifos. The compound present in the treated sample after degradation of chlorpyrifos was separated and identified by GC-MS analysis. The chlorpyrifos was detected at Retention Time (RT) of 11.27 and the compound was identified as Thienol-3,5 Pyridine. This compound confirmed the degradation of chlorpyrifos by *Pseudomonas* sp.

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Abbreviations

CP	Chlorpyrifos
GC-MS-Gas	Chromatography-Mass Spectroscopy
MSM	Mineral Salt Medium
UV-Vis	Ultraviolet—Visible
Nm	Nanometer
RT	Retention Time
OD	Optical Density
μl	Microliter
ml	Milliliter
hrs	Hours
CFU	Colony Forming Unit
rpm	Rotation Per Minute
IMViC	Indole, Methylred, Voges Proskauer and Citrate Test
TSI	Triple Sugar Iron

4.1 Introduction

India is a country that relies overwhelmingly on agriculture; about 60–70% of the population work in this sector. Industries are rapidly occupying a significant percentage of arable land that is presently being cultivated. A substantial rise in agricultural output has been produced by the employment of modern technologies and agricultural inputs including fertiliser, hybrid seeds, and pesticides. The widespread usage of pesticides over time has led to issues brought on by their interactions with the environment's biological system. Based on their chemical composition, pesticides are synthesized in a wide range of forms, including organochlorine pesticides, organophosphorus pesticides, carbamate, pyrethroids, and biopesticides. Organophosphates are a significant component of herbicides, insecticides, pesticides and nerve gas. The ester derivatives of phosphoric acid are known as OPs, which undergo degradation quite quickly and are thought to be safest for use in agriculture. OPs, which are the ester forms of phosphoric acid, are normally regarded as acceptable for use in agriculture due to their relatively quick rates of degradation (Renita et al. 2021). Because of the prohibition of organochlorine pesticides like lindane and dichlorodiphenyltrichloroethane (DDT), the dependency of organophosphorus insecticides in agriculture has increased. Pesticides are poisonous agents that are purposefully introduced into the ecosystem to eradicate living organisms like insects

(insecticides), weeds (herbicides), rodents (rodenticides), fungi (fungicides), and others. Insecticides are generic poisons that affect more than just the “pests” they are intended to kill. Nearly 90% of pesticides never actually affect target species; rather, contaminates soil and water with an impact on unintended organisms (Vidali 2001). Organophosphate pesticides have an adverse effect on non-target plants and non-target animal species that inhabiting an extensive range of aquatic and terrestrial habitats (Blann et al. 2009).

Organophosphorous pesticides are a major and common category of pesticides now in use throughout, contributing for greater than 36% of the global market (Kanekar et al. 2004). Pesticide exposure not only has a lot of negative health consequences and connected to a number of significant human illnesses and diseases, ranging from respiratory issues to cancer. The signs of this pesticide exposure include nausea, blurred vision, breathing issues, sweating, headaches, slow heartbeats, numbness, and cramps in the abdomen (Vidya Lakshmi et al. 2009). In 2001, the United States Environmental Protection Agency (EPA) controlled the use of chlorpyrifos to specific crops and outlawed its residential usage in the country due to its toxicity (Eaton et al. 2008).

Chlorpyrifos is an agricultural insecticide used globally against various pests and insects. Based on the soil nature and texture, CP has a half-life varying between 10 and 120 days in the soil that undergoes degradation slowly in soil, arising a major environment and public health threats (Silva 2020). This chemical has broad spectrum action on the insect nervous system and cause mortality by inhibiting the important acetylcholine esterase enzyme. In humans, severe CP poisoning cases shown symptoms such as headache, nausea, dizziness, muscle twitching, weakness, increased sweating, and salivation (Ambreen and Yasmin 2021). It also shows immunotoxic effects on humans and animals. When these pesticides build up in agricultural fields, they permeate into the nearby water bodies and interrupt the metabolic processes of aquatic plants and animals. The persistent use of CP has generated numerous issues in environmental and biological systems. As a result, it is critical to remove chlorpyrifos (CP) from contaminated surroundings. CP detoxification is accomplished by physical, chemical, and biological techniques (Huang et al. 2021). A number of physicochemical techniques for removing chemicals from the environment, such as photodecomposition, chemical treatment, volatilization, and combustion are followed. These methods are ineffective for totally destroying CP and its metabolic degradation products (Natarajan et al. 2021).

Chlorpyrifos must be safely degraded by microorganisms in order to avoid leaving hazardous residues. Different types of bacteria can break down various pesticides, herbicides and insecticides (Munazza 2005). The technique of bioremediation involves employing microorganisms to transform hazardous or poisonous complex organic substances into smaller, non-toxic inorganic molecules (Pang et al. 2020). The persistence of most pesticides in soil is impacted by microbial activities. Toxic chemical removal from the environment through microbial biodegradation is a very appealing strategy. A simple, affordable, and eco-friendly method for removing CPF from soil and water is bioremediation (Rayu et al. 2017).

Microbial remediation effectively removes pollutants from the environment using bacteria, fungi, or algae. Many bacteria from the genera *Pseudomonas*, *Alcaligenes*, *Bacillus*, *Xanthomonas*, *Enterobacter*, and *Ochrobactrum* have been found and characterised as being capable of degrading CPF (Huang et al. 2021). According to reports, algae and fungi may also effectively break down TCP and chlorpyrifos. The major result of CPF's degradation in the soil is 3,5,6-TriChloro-2-Pyridinol (TCP) (Chen et al. 2012). *Pseudomonas* is one of them, a bacterial genus that is widely distributed in soil and is essential for the mineralization of organic matter. They can breakdown the majority of pesticides due to their metabolic adaptability (Sarkar et al. 2009).

Pseudomonas species can be used for bioremediation and some of its members are capable of metabolising chemical contaminants in the environment.

Pseudomonas is a diverse genus that includes several catabolic processes and enzymes involved in the breakdown of pesticides. The aim of this work was to identify and characterise CP-degrading microorganisms in order to assess their potential for degradation. In the present study, chlorpyrifos degrading *P. aeruginosa* and *P. fluorescens* was isolated from chlorpyrifos contaminated soil. The growth of chlorpyrifos degrading *Pseudomonas* was assessed. The breakdown of chlorpyrifos by *Pseudomonas* was characterized by UV-Vis spectrophotometer. For further confirmation of chlorpyrifos degradation, the presence of breakdown product by GC-MS analysis was studied.

4.2 Materials and Methods

4.2.1 Collection of Soil Samples

The collection of soil samples was done randomly in the chlorpyrifos contaminated agricultural field, Erode District, Tamil Nadu, India. The selected agricultural field is regularly applied with agrochemicals for crop cultivation. The surface soil and debris layer were removed, at 10 cm depth soil samples were collected and transferred into a sterile container. The collected soil sample were transported to the laboratory and stored at 4 °C for further experiments (Singh et al. 2004).

4.2.2 Isolation of Chlorpyrifos Degrading Bacteria

From the collected soil sample, 1 g soil was weighed and serially diluted by the following standard microbiological plating methods. For the isolation of CP degrading bacteria, serially diluted samples were plated on the nutrient agar plate amended with 0.1% chlorpyrifos by spread plate technique. The inoculated plates were subjected to incubation at 37 °C for 24 to 48 h (Reetha and Kavi Karunya

2012). Then the samples were plated on the *Pseudomonas fluorescens* isolation agar plate by spread plate technique. After incubation duration for 24 h at 37°, bacterial colonies appeared. Further, the colonies were streaked into nutrient agar (NA) plates and sub-cultured sequentially to obtain pure colonies (Sharma et al. 2017).

4.2.3 Identification of CP Degrading Bacteria

The colonies were subjected to morphological and biochemical characterization by following Bergey's Manual of Determinative Bacteriology (Holt 1994). Colony characteristics used for identification include size, shape, color, margin, and elevation. Besides, the bacterial cells were stained and visualized under a compound microscope, supported by the Gram Staining Technique (Rayu et al. 2017). Subsequently, biochemical tests were performed, including Indole, Methyl-Red (MR), Voges Proskauer (VP), Citrate Utilization, Catalase, Oxidase, Urease, Lactose fermentation, and Hydrogen sulphide production.

4.2.4 Enrichment Technique and Isolation

Enrichment culture technique (Barathidasan et al. 2014) was followed to isolate potential CP degrading bacteria using Minimum Salt Medium (MSM). MSM composition as follows (in grams per liter of distilled water): Ammonium Nitrate ($(\text{NH}_4)_2\text{NO}_3$)—1 g; Hydrated Calcium Nitrate ($\text{Ca}(\text{NO}_3)_2 \cdot 2\text{H}_2\text{O}$)—0.04; Hydrated Magnesium Sulphate ($\text{MgSO}_4 \cdot 7\text{H}_2\text{O}$)—0.1 g; Potassium Chloride (KCl)—0.2 g; Hydrated Iron (ii) sulfate ($\text{FeSO}_4 \cdot 7\text{H}_2\text{O}$)—0.001, Dipotassium phosphate ($\text{K}_2\text{HPO}_4 \cdot 12\text{H}_2\text{O}$)—1.5; and Potassium dihydrogen phosphate (KH_2PO_4)—4.8. The pH of the above medium was adjusted to 7 using a Benchtop pH meter. The degradation efficiency of bacteria was compared in two different media such as MSM and Nutrient broth amended with CPD. In MSM, chlorpyrifos is the only carbon source for utilization of bacteria (Kumar 2011). The prepared Mineral Salt Medium (MSM) was sterilized in an autoclave at 121 °C for 15 min at 15 lbs (Sharma et al. 2016). The medium was allowed to cool. The different concentrations of chlorpyrifos were added to the tubes aseptically from 0.5 to 2.5%. About 0.1 ml bacterial culture was inoculated into the tubes. Then, the incubation of inoculated tubes was done at 37 °C for 2 days.

The prepared Nutrient Broth (NB) was sterilized in an autoclave at 121 °C for 15 min at 15 lbs. The medium was allowed to cool. The different concentrations of Chlorpyrifos pesticides were added to the tubes from 0.5 to 2.5%. About, 0.1 ml of bacterial culture was inoculated aseptically into the tubes. The incubation of tubes was done at 37 °C for 24–48 h.

4.2.5 UV Spectrophotometric Analysis

When the bacterial cultures were inoculated in the MSM medium lacking carbon source, the bacteria start utilizing alternative carbon source (CP) supplemented in MSM (Kumar 2011). CP degradation in the media was analysed by taking the Optical Density (OD) after centrifugation of the minimal media and nutrient media amended with chlorpyrifos at the absorbance maximum of 300 nm (Kumar et al. 2015).

4.2.6 GC-MS for Determination of Chlorpyrifos Intermediates

The identification of chlorpyrifos by efficient bacterial isolate was analyzed by gas chromatography-mass spectrometry (GC-MS) to identify degradation metabolites. GC-MS was set up with a split-splitless injector and attached to a triple quadruple mass spectrophotometer (Yin et al. 2021). The capillary column was used for the separation (Li et al. 2010). The carrier gas used at a constant flow rate of 1 ml/min was Helium gas (99.999%) and 3 μ l injection volume was employed. The injector temperature was kept at 250 °C for 5 min in splitless mode. The oven temperature was set as follows:

- Initial temperature at 100 °C (hold for 2 min)
- 20 °C /min to 180 °C
- 10 °C/ min to 250 °C (hold for 2 min).

The total running time for GC-MS analysis was 15–20 min. Electron ionization and spectra were recorded at 70 eV. The GC-MS interface was placed at 250 °C and the ion source temperature was placed at 200 °C. The total ion current (TIC) chromatograms were recorded ranging from 40 to 500 m/z. The individual constituents of GC report were identified by comparing the Mass Spectra (MS) with the standard library of National Institute for Standard Technology mass spectra (NIST-MS).

4.3 Results and Discussion

The chlorpyrifos degrading and non-degrading bacterial colonies were appeared on the nutrient agar plates. The good numbers of chlorpyrifos degrading bacteria were obtained after 48 h of incubation. The bacterial colonies were obtained from the *Pseudomonas fluorescens* isolation agar plates. At 10^{-4} dilution, 153 CFU/ml was isolated. Isolation and exploration of indigenous microbial strains for *in-situ* detoxification is an environmentally friendly approach (Ortiz-Hern et al. 2010). Indigenous species are preferable as they do not destructively affect the population and diversity of soil micro-flora (Farhan et al. 2021).

Table 4.1 Morphological characterization of CDB₁ and CDB₂ isolates

Isolates	Colony morphology	Pigmentation	Gram staining	Motility
CDB ₁	Mucoid, elevated	Bluish green	Gram negative rods	Motile
CDB ₂	Smooth convex glistening	Yellowish green	Gram negative rods	Motile

The isolates tolerated to grow on chlorpyrifos containing media were selected for screening. For pure culture, the colonies were picked up from the plates and sub-cultured by quadrant streak plate method. After overnight incubation, the culture was stored at 4 °C.

The isolated bacteria were subjected to morphological characterization by following standard staining procedures. The dominating colonies isolated were mucoid and elevated with appearing bluish green pigmentation. Other colonies were showing smooth, convex, glistening with yellowish pigmentation. The staining procedures confirmed the bacterial cells as Gram negative rod shaped. The motility test by hanging drop technique showed the isolates as motile bacteria. The results are represented in the Table 4.1.

The biochemical tests were performed for the identification of the organisms. The Indole test, Methyl Red (MR), Voges-Proskauer (VP) and citrate test showed the negative result for both CDB₁ and CDB₂ isolates. The catalase and oxidase test showed the positive result for both isolates. The TSI test showed negative result for CDB₁ and CDB₂.

Both the isolates were subjected to partial identification based on various biochemical tests with reference to the Bergey's Manual of Determinative Bacteriology. The isolates were identified as a member of *Pseudomonas* sp. and it was identified that CDB₁ as *Pseudomonas aeruginosa* and CDB₂ as *Pseudomonas fluorescens*. However, the results are like those which were obtained by concerning the identification of *Pseudomonas* sp. The results are represented in the Table 4.2.

After the incubation period, the nutrient and MSM medium at different concentration of pesticides, the turbidity was observed. The cultures were centrifuged at 5000 rpm for 20 min and the supernatant were collected. Spectral analysis was made using UV-Vis Spectrophotometer and the cell free extract was scanned in the range of 300 nm to find out the maximum absorption of spectra (Table 4.3). The control was used as a blank (Reetha and Kavi Karunya 2012; Kumar et al. 2015).

P. aeruginosa and *P. fluorescens* had the range of different values at different concentration of pesticides when absorbed at the maximum spectra of 300 nm. The CP degrading *P. aeruginosa* had shown the maximum degradation when incorporated

Table 4.2 Biochemical characterization of CDB₁ and CDB₂ isolates

Isolates	Indole	Methyl red	Voges Proskauer	Citrate	Catalase	Oxidase	Gelatin	TSI		
								G	L	S
CDB ₁	–	–	–	+	+	+	+	–	–	–
CDB ₂	–	–	–	+	+	–	+	–	–	–

Table 4.3 UV-Vis spectrophotometer analysis of chlorpyrifos degrading *Pseudomonas*

Chlorpyrifos concentration	<i>P. aeruginosa</i>		<i>P. fluorescens</i>	
	Nutrient broth (OD at 300 nm)	MSM medium (OD at 300 nm)	Nutrient broth (OD at 300 nm)	MSM medium (OD at 300 nm)
Control	0.00	0.00	0.00	0.00
0.5	4.00	1.48	4.00	1.50
1.0	4.00	1.28	3.69	1.02
1.5	3.77	0.44	0.50	0.55
2.0	1.27	0.38	0.43	0.22
2.5	0.60	0.37	0.35	0.21

with 0.5c of chlorpyrifos after 2 days of incubation period. The results were showed in the Table 4.3. In case of *P. fluorescens*, the degradation of the chlorpyrifos pesticide was low comparing to *P. aeruginosa* under similar culture condition. For degradation of chlorpyrifos, MSM medium found suitable by both isolates when compared to nutrient broth. The change in the biodegradation rate depends on the concentration of the pesticides, substrate availability, incubation period and the ability of bacteria. The chlorpyrifos degradation by *P. aeruginosa* was further subjected to GC-MS analysis (Fig 4.1).

In the present study, GC-MS was used to identify degradation products of chlorpyrifos. The GC-MS is the most sensitive method to identify the degradation products of organochloropesticides (Farhan et al. 2021). After the degradation studies by UV-Vis Spectrophotometer analysis, comparatively the isolate I have given the higher degradation spectrum of chlorpyrifos. The compound present in the treated sample after degradation of chlorpyrifos was separated and identified by GC-MS analysis and it was showed in the Table 4.4 (Barathidasan et al. 2014).

The degradation of chlorpyrifos was confirmed by GC MS analysis. The chlorpyrifos was detected at Retention Time (RT) of 11.27 min. The reproducibility was confirmed using an injector (Kumar 2011). The cell free extracts of samples from the culture medium were subjected to GC MS analysis. This analysis was done to confirm the presence of degradative products of chlorpyrifos by microbial activity. The results of the present study represent that *P. aeruginosa* was known to degrade even high concentration of chlorpyrifos (2.5c). The active degradation of chlorpyrifos was obtained in 0.5c and 1.0c by *P. aeruginosa*. In *P. flourescens*, the highest chlorpyrifos degradation was recorded in the 0.5c, followed by 1.0c, 1.5c, 2.0c and 2.5c. The highest chlorpyrifos residue was obtained in 0.5c and 1.0c. The main potential CP degraders belonging to *Pseudomonas* sp. It has the potential to make use of CP when provided as sole carbon source (Amani et al. 2019).

By comparing both isolates, *Pseudomonas aeruginosa* have the higher degradation of chlorpyrifos (Manigandan and Nelson 2015). *Pseudomonas* sp. degrades many pesticides like Alachlor, Chlorprophan, DDT etc. and the residues found as Thienol 3,5 Pyridine, 3,4 Dihydroxy 1,6 bis (3-methoxy-phenyl)-hexa-2,4diene,

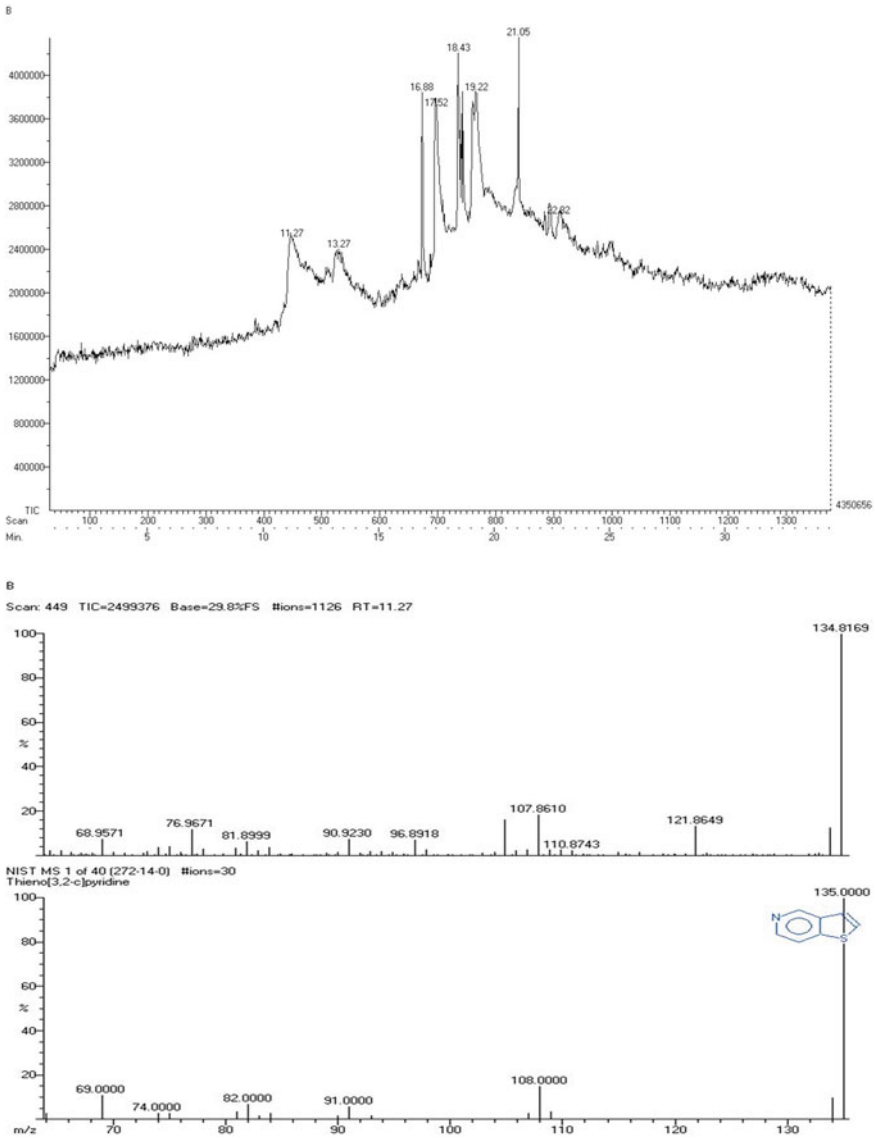
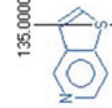
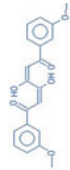







Fig. 4.1 Chromatogram showing Chlorpyrifos degradation by GC-MS Analysis

Oleic acid, n-hexadecanoic acid and Iso propyl stearate. The chlorpyrifos degradation was detected at Retention Time (RT) of 11.27 and the compound was identified as Thienol-3,5 Pyridine.

Although just a few species have been demonstrated to breakdown the compounds due to their complexity, bacteria and fungi are the primary transformers. Using microorganisms to immobilise or decompose contaminants is a common method

Table 4.4 GC-MS analysis of the treated sample

Compound name	Retention time	TIC	Structure
Thienol 3,5 pyridine	11.27	2,499,376	
3,4 dihydroxy-1,6-bis-[3-methoxy-phenyl]hexa-2,4-diene-1,6 diene	22.82	2,715,088	
n-hexadecanoic acid	17.52	3,702,304	
Isopropyl stearate	21.05	4,350,656	
Ascardiol epoxide	13.47	2,401,136	
Dodecanoic acid	16.88	3,841,936	
Oleic acid	18.43	4,204,384	

of removing them, and this process is known as bioremediation. The potential for these compounds to enter animals and subsequently human consumers of animal products makes biodegradation of OPs in the soil and water crucial (Chapman et al. 1986). Yadav et al. (2016) reported that bacteria contain catabolic genes that allow them to thrive in a variety of ecological niches at varying pH, temperature, and oxygen concentrations, which enables them to digest contaminants effectively.

From the study, it is concluded that CP undergoes degradation by many soil microorganisms. Bacterial degradations of CP are dependent on culture conditions, isolates and species of bacteria involved (Singh and Walker 2006). *Pseudomonas* sp. strains have been found to effectively breakdown the pesticide and its primary metabolite both in the absence and presence of external nutritional supplements, according to study on bacteria with the ability to degrade (Bose et al. 2021). Future research is proposed to investigate and develop the strains obtained in the current study for enhanced bioremediation of CP in soils. A bacterial consortium that uses symbiotic relationships among the strains to degrade CP must be developed. To improve our knowledge of the biodegradation process, additional research should focus on numerous biochemical factors, genes, and enzymes.

Declaration

Conflicts of Interest: All authors have no conflict of interest.

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Chapter 5

Studies on the Effects of Metribuzine on *Phormidium* and *Chroococcus* Species



S. Balakumar and N. Mahesh

Abstract The efficacy of metribuzine, a selective herbicide recommended for comprehensive weed control in rice, was investigated on *Phormidium* and *Chroococcus* species, common species of cyanobacteria obtained from the National Marine Cyanobacterial Facility (NFMC). A comparative study of the effects of metribuzine on both species of cyanobacteria measures their chlorophyll-a content and biochemical levels such as carbohydrate, protein, amino acid and lipid levels over 18 days of time- and dose-dependent exposure. was evaluated under laboratory conditions by Results showed that the herbicide exerted pleiotropic effects on both cyanobacterial species over the concentration range tested (2–10 μg). Chlorophyll a was more adversely affected by metribuzine at the highest concentration (10 μg) on day 18. On day 12, more than 50% inhibition of cyanobacterial growth was observed when 10 μg of metribuzine was applied. Carbohydrate, protein, amino acid, and lipid levels were reduced with herbicide, but by day 18, effects were noticeable at the highest concentrations for both cyanobacteria. Metribuzine concentrations in the 2 μg range also affected chlorophyll a and biochemical content in both cyanobacteria. *Phormidium* species were more affected by metribuzine than *Chroococcus* species. Since, *Phormidium* and *Chroococcus* species are abundant in paddy fields and can be used as inoculum for rice biofertilizer programs, protecting them from the potential residual effects of herbicides is critical to correctly managing local soil fertility.

Keywords *Phormidium* species · *Chroococcus* species · Metribuzin · Inhibitory action · Paddy fields

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5.1 Introduction

Although cyanobacteria are essentially microscopic, large colonies and mats are highly visible. They are a large, morphologically diverse group of photosynthetic prokaryotes that are involved in oxygenic photosynthesis and include unicellular to multicellular microorganisms found in nearly every habitat on earth (Carr and Whitton 1982; Vermass 2001). Cyanobacteria have a similar appearance to algae and have similar light, nutrient and carbon dioxide requirements. Even without light, some cyanobacteria can survive and grow using chemicals from the environment. They store carbon in different forms such as starches, oils and various sugars. Their main photosynthetic pigment is chlorophyll a (chl_a), and other photosynthetic pigments in cyanobacteria are various phycobiliproteins (phycoerythrin, phycocyanin, allophycocyanin) and carotenoids (beta-carotene, echinenone, myxoxanthophyll). Because of these later pigments and mucus, the color of cyanobacteria in nature can range from dirty yellow to various shades of blue-green, brown or black (Whitton and Potts 2000). Some species of the genus have small gas sacs inside their cells that float to the surface or sink to the bottom, depending on changing light and nutrient availability.

Cyanobacteria also live in a variety of terrestrial environments. Soil is one of the most potential habitats for algae, especially in moist or damp conditions. Cyanobacteria are one of the major constituents of nitrogen-fixing biomass in paddy fields. The agricultural importance of cyanobacteria in rice cultivation is related to their ability to fix atmospheric nitrogen on plants and soil (Whitton 2000; Silva and Silva 2007). Some genera of heterocysts are diazotrophic and can use atmospheric nitrogen as the sole nitrogen source, thus contributing to light-dependent nitrogen fixation in rice paddies worldwide, maintaining and maintaining soil fertility. It plays an important role in building. In addition to heterocyst nitrogen fixation, some non-heterocysts fix atmospheric nitrogen under microaerobic conditions. Some non-heterocyst forms often crusts in soil. Crust formation impedes soil erosion, increases rainwater retention, and reduces evaporative water loss (Thajuddin and Subramanian 2005).

The use of herbicide may have harmful effects on aquatic life. Agricultural runoff during rainfall often concentrates in lakes and ponds (Richards and Baker 1993). Cyanobacteria are one of the potential organisms that could benefit humanity in many ways. Cyanobacteria represent a vast potential resource in a variety of applications, including aquaculture, food, feed, fuel, fertilizer, pharmaceuticals, industry and environmental protection (Patterson 1996).

The aim of this study to investigate the adverse effect of herbicide metribuzine on *Phormidium* sp. and *Chroococcus* sp., found in the paddy field and the conservation of local soil fertility. Therefore, this study may help to improve the implementation of effective integrated rice pest management and likely to succeed for future use of cyanobacteria in rice fields.

5.2 Materials and Methods

5.2.1 Cultivation of Cyanobacteria

Filamentous and rounded diazotrophic cultures of the cyanobacteria, *Phormidium* sp. and *Chroococcus* sp., were obtained from NFMC. Purchased cultures were propagated in artificial synthetic nutrient III (ASN III) broth and placed in a rotary shaker at room temperature. The inoculated plates were regularly examined for cyanobacterial growth. Grown cultures were sub-cultured periodically and stored at 4 °C for future use.

5.2.2 Herbicide Stock Solution

Herbicide stock solution was prepared by dissolving 1 mg of herbicide in 100 ml of sterile ASN III medium to obtain a stock solution. Solutions of various concentrations were prepared from this herbicide stock solution as 2, 4, 6, 8, 10 µg/ml respectively.

5.2.3 Herbicide Efficacy

To test herbicide efficacy against the cyanobacterial species *Phormidium* sp and *Chroococcus* sp., were selected and treated with different herbicide concentrations. The required amount of herbicide was added to the liquid culture medium after sterilization. A volume of 0.5 ml of exponentially growing culture was then added equally to each different set of replicates (100 ml). Morphological changes supplemented with different herbicide concentrations were regularly observed. Chlorophyll a content and biochemical content were measured at individual day (6, 9, 12, 15, and 18) intervals. Experiments were performed in duplicate for 18 days under controlled conditions.

5.2.4 Determination of Chlorophyll a

The isolates were centrifuged at 5000 rpm for 10 min and the pellet was suspended in 4 mL of 80% methanol. The tubes were sealed to prevent solvent evaporation and incubated at 60 °C in the dark for 1 h in shaker. The contents were cooled and centrifuged at 5000 rpm for 5 min. The supernatant was collected and repeated again for more pigment extraction. The supernatant was made up to a known volume with 80% methanol. Absorbance was measured at 663 nm.

5.2.5 Carbohydrate Estimation

The cultures were centrifuged at 5000 rpm for 10 min and 100 mg of the pellet was placed in a test tube and hydrolyzed with 2 ml of concentrated H₂SO₄ at 100 °C for 30 min. 1 mL of 5% phenol and 5 mL of concentrated H₂SO₄ were added to 0.5 mL of hydrolyzate and mixed well. Color development was measured at 490 nm using a spectrophotometer. Glucose was used as standard.

5.2.6 Protein Estimation

The 100 mg of the pellet collected after centrifugation was suspended with 10% TCA and centrifuged at 10,000 rpm for 10 min. The pellet was resuspended in 1N NaOH, boiled for 30 min, then re-centrifuged to remove light scattering material. Supernatant was made up to a known volume. To this 0.9 ml of distilled water and 5 ml of alkaline copper reagent were added to 0.1 ml of supernatant, left for 10 min, and finally 0.5 ml of Folin-Ciocalteu reagent was added. After 30 min, absorbance was read at 750 nm and protein levels were calculated using a standard chart made using bovine serum albumin.

5.2.7 Amino Acid Deduction

The 100 mg of the pellet after centrifugation was homogenized in 80% ethanol. The clear supernatant was adjusted to a known volume after centrifuged at 5000 rpm. 1 ml of this was pipetted into a test tube and diluted to 4 ml with distilled water. To this 1 ml of ninhydrin reagent was added and kept in a boiling water bath for 15 min. The tube was then cooled and 1 mL of 50% ethanol was added. The induced violet color was measured with a spectrophotometer at 540 nm.

A standard plot was made using a mixture of alanine, aspartic acid, tryptophan, proline, and lysine.

5.2.8 Lipid Estimation

The 100 mg of the pellet was homogenized and extracted by solvent (chloroform:methanol 2:1 (v/v)) and filtered by Whatman paper. The filtrate was vortexed with sodium sulfate to remove moisture. It was then collected in pre-weighed bottles and left overnight at room temperature in the dark. Total lipids were estimated by weighing the dried extract.

5.3 Results and Discussion

Successful rice farming is progressively reliant on the application of herbicides to remove unwanted weeds. However, heavy use of herbicides raises the level of chemical residues to levels that can inhibit the growth of native microbes such as cyanobacteria. Anything that affects these primary producers can affect the ecological balance of the entire paddy agroecosystem. Share the characteristics of This susceptibility of cyanobacteria to herbicides varies, but mainly depends on the type and type of herbicide. Evaluation of the toxic effect of metribuzine on cyanobacterial abundance and distribution shows progressive inhibition of cyanobacterial growth recorded compared to controls. These inhibitory effects could be attributed to the lethal effect of metribuzine on cyanobacterial growth and biochemical content. These results are consistent with those of Singh et al. (1983) reported that the thiocarbamate his benchocarb inhibited the growth of his *Nostoc linckia*. Similar results were also reported by Metting (2008), who found that the growth and dry weight of *Anabaena oryzae* and *Auroscilla fertilisima* were significantly reduced by the application of parathion. The herbicide metribuzine binds to soil components immediately after application, so cyanobacteria are constantly exposed to potential hazards from these herbicides (Table 5.1) (Gustavson et al. 2003).

Herbicides interfere with photosynthetic pigment synthesis by inhibiting perphyrin formation and biosynthetic steps in carotenoid synthesis, ultimately leading to chlorosis and precursor accumulation (Mohapatra et al. 1996). Such accumulation causes the necessary blockade of cell junctions that further impairs metabolic efficiency.

Cyanobacterial growth was measured in terms of chlorophyll 'a'. Chlorophyll a was recorded at 37.10 mg/mL, 32.57 mg/mL and 52.08 mg/mL, 44.26 mg/mL in the *Phormidium* and *Chroococcus* sp. controls on days 6 and 18, respectively. Concurrently, a loss of 5.45 and 34.60 mg/ml chlorophyll a content was observed in his *Phormidium* species treated with the lowest concentration (2 µg/ml) of metribuzine on days 6 and 18. At the highest concentration (10 µg/ml), a loss of chlorophyll a content of 24.03 and 51.62 mg/ml was observed in metribudine-treated formidium species on days 6 and 18 (Table 5.2; Fig. 5.1). In *Chroococcus* species treated with metribuzine, a loss of 5.61 and 30.

Chlorophyll a levels of 64 mg/ml were recorded at the lowest concentration (2 µg/ml) on days 6 and 18, respectively. At the highest concentration (10 µg/ml), a loss of chlorophyll a content of 25.12 and 43.52 mg/ml was recorded on days 6 and 18 (Table 5.2; Fig. 5.1). Similar observations were reported in.

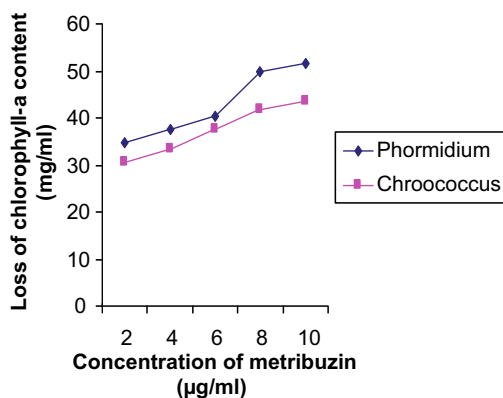
Anabaena cylindrica, with the herbicides bentazone and molinat, which decreased chlorophyll a levels with increasing herbicide concentrations (Galhano et al. 2009). Yan et al. (1997) confirmed a decrease in growth rate of *Anabaena sphaelica* with increasing molinate concentration. Eladel et al. (1999) and Battah et al. (2001) also reported that thiobencarb induced a decrease in the specific growth rate of *Anabaena he variabilis* and significantly reduced its biomass yield. Kobbia et al. (2001) all tested concentrations (0.2–0.8 mg/l) photosynthetic inhibitors of simazine, triazine,

Table 5.1 Physico-chemical properties of metribuzin (Maria Diva et al. 1998)

Herbicide	Chemical family and chemical abstract service number	Chemical name	Chemical formula	Molecular weight (g.mol ⁻¹)	Melting point (°C)
Metribuzin	S-triazine CAS no: 21097-64-9	(4-amino-6-tert-butyl-4,5-dihydro-3-methylthio-1,2,4-triazine-5-one)	C ₈ H ₁₄ N ₄ O ₂ S	214.28	125.5–126.5

Table 5.2 The effect of metribuzin on chlorophyll-a in cyanobacteria

S.No	Concentration of Metribuzin ($\mu\text{g/ml}$)	Loss of chlorophyll-a in <i>Phormidium</i> sp., (mg/ml)					Loss of chlorophyll-a in <i>Chroococcus</i> sp., (mg/ml)				
		(Days)					(Days)				
		6	9	12	15	18	6	9	12	15	18
1	2	5.45	14.27	20.89	27.61	34.60	5.61	12.70	17.47	24.75	30.64
2	4	8.48	16.01	23.83	29.17	37.54	8.83	15.37	21.9	28.71	33.49
3	6	12.07	18.13	26.41	31.38	40.39	13.89	19.60	25.85	32.39	37.54
4	8	16.58	23.10	29.63	35.24	49.87	15.73	22.73	28.15	36.07	41.68
5	10	24.03	30.65	36.16	42.60	51.62	25.12	29.82	34.04	40.3	43.52

Fig. 5.1 The effect of metribuzin on chlorophyll-a content on 18th day

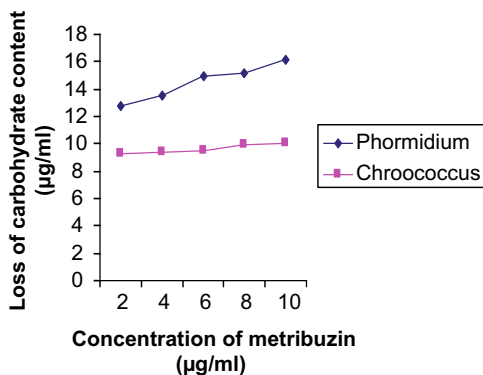
and reducing side PS II. Our results also contradict those of Gonzalez-Tome (1996), who studied *Nostoc* species. If double bentazone was used. Xia (2005) showed that thiobencarb had a negligible effect on chlorophyll-a synthesis in her *Nostoc sphaeroides* colonies.

Biochemical studies such as carbohydrate, protein, amino acid and lipid contents were analyzed in both control and herbicide-treated cyanobacterial species. At the highest concentration of herbicide (10 $\mu\text{g/ml}$), a significant decrease in the content of all biochemical components was observed on day 18 in both cyano-his bacteria. *Phormidium* species treated with the lowest concentration of metribuzine (2 $\mu\text{g/ml}$) on days 6 and 18 showed a loss of carbohydrate content at 0.8 and 12.8 $\mu\text{g/ml}$. At the highest concentration (10 $\mu\text{g/ml}$), a carbohydrate loss of 9.2 and 16.1 $\mu\text{g/ml}$ was observed in metribudine-treated formidium species on days 6 and 18, respectively (Table 5.3; Fig. 5.2). *Chroococcus* sp. treated with the lowest concentration (2 $\mu\text{g/mL}$) of metribuzine had carbohydrate losses of 1.7 and 9.3 $\mu\text{g/mL}$ on days 6 and 18, respectively.

At the highest concentration (10 $\mu\text{g/ml}$), a loss of carbohydrate content of 6.7 and 10.0 $\mu\text{g/ml}$ was recorded on days 6 and 18, respectively (Table 5.3, Fig. 5.2).

Table 5.3 The effect of metribuzin on carbohydrate in cyanobacteria

S.No	Concentration of Metribuzin ($\mu\text{g/ml}$)	Loss of carbohydrate in <i>Phormidium</i> sp., ($\mu\text{g/ml}$)					Loss of carbohydrate in <i>Chroococcus</i> sp., ($\mu\text{g/ml}$)				
		(days)					(days)				
		6	9	12	15	18	6	9	12	15	18
1	2	0.8	4.9	7.0	9.8	12.8	1.7	2.6	6.1	8.0	9.3
2	4	4.9	5.8	7.6	9.9	13.5	2.3	3.6	7.1	8.1	9.4
3	6	6.2	7.0	9.1	12.3	15.0	2.6	5.1	7.2	8.6	9.5
4	8	6.5	7.8	9.9	12.4	15.2	5.7	6.4	7.7	8.9	9.9
5	10	9.2	9.6	12.2	14.0	16.1	6.7	7.3	8.6	9.1	10.0

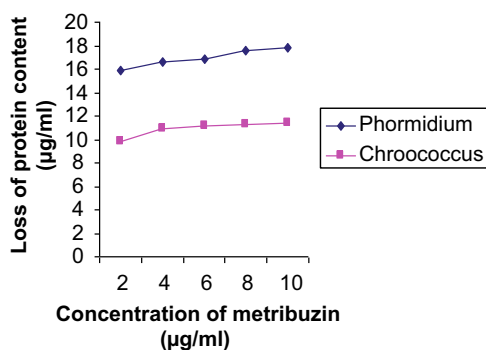
Fig. 5.2 The effect of metribuzin on carbohydrate content on 18th day

Concurrently, carbohydrate content losses of 10.6 $\mu\text{g/mL}$, 16.2 $\mu\text{g/mL}$ and 7.6 $\mu\text{g/mL}$, 10.2 $\mu\text{g/mL}$ were recorded in the *Phormidium* and *Chroococcus* sp. controls on days 6 and 7, respectively. Cobia et al. (2001) reported that simazine has the ability to suppress the carbohydrate content of *Anabaena cylindrica*. Garjano et al. (2009) reported that treatment with bentazone resulted in significant inhibition of carbohydrate synthesis. The loss of protein content was recorded as 9.4 $\mu\text{g/ml}$, 17.8 $\mu\text{g/ml}$, and 9.0 $\mu\text{g/mL}$ and 11.4 $\mu\text{g/mL}$ for *Phormidium* and *Chroococcus* sp. controls on days 6 and 18, respectively. were recorded at protein content losses of 1.7 and 15.9 $\mu\text{g/ml}$. At the highest concentration (10 $\mu\text{g/ml}$), a loss of protein content of 8.1 and 17.8 $\mu\text{g/ml}$ was observed in metribudine-treated formidium spp. on days 6 and 18, respectively (Table 5.4; Fig. 5.3). *Chroococcus* species treated with the lowest concentration (2 $\mu\text{g/mL}$) of metribuzine showed a protein loss of 1.6 and 9.8 $\mu\text{g/mL}$ on days 6 and 18, respectively. At the highest concentration (10 $\mu\text{g/ml}$), a protein loss of 7.1 and 11.4 $\mu\text{g/ml}$ was recorded on days 6 and 18, respectively (Table 5.4; Fig. 5.3).

Yang et al. (1997) reported that molinate inhibited protein hydrolysis by inhibiting protease activity and resulted in total protein accumulation in cyanobacterial cells. Amino acid content losses of 3.0 and 11.1 $\mu\text{g/ml}$ were recorded in *Phormidium*

Table 5.4 The effect of metribuzin on protein in cyanobacteria

S.No	Concentration of Metribuzin ($\mu\text{g/ml}$)	Loss of protein in <i>Phormidium</i> sp., ($\mu\text{g/ml}$)					Loss of protein in <i>Chroococcus</i> sp., ($\mu\text{g/ml}$)				
		6	9	12	15	18	6	9	12	15	18
1	2	1.7	4.8	9.8	13.7	15.9	1.6	3.1	4.6	7.9	9.8
2	4	3.5	5.8	10.4	13.8	16.6	4.9	5.8	8.7	10.0	10.9
3	6	5.5	7.3	11.5	14.5	16.8	5.3	7.0	9.1	10.1	11.2
4	8	6.7	8.6	12.1	14.9	17.6	6.0	7.6	9.2	10.3	11.3
5	10	8.1	9.4	12.6	15.6	17.8	7.1	7.8	9.6	10.5	11.4

Fig. 5.3 The effect of metribuzin on protein content on 18th day

species treated on day 1. At the highest concentration (10 $\mu\text{g/ml}$), loss of amino acid content of 10.1 and 15.4 $\mu\text{g/ml}$ was seen in metribuzin-treated formidium species on days 6 and 18 (Table 5.5; Fig. 5.4). In his *Chroococcus* sp. treated with the lowest concentration (2 $\mu\text{g/ml}$) of metribuzine, a loss of amino acid content of 0.7 and 9.6 $\mu\text{g/ml}$ was observed on days 6 and 18, respectively. At the highest concentration (10 $\mu\text{g/ml}$), 5, 4 and 11 amino acid content is lost. 4 $\mu\text{g/ml}$ was recorded on days 6 and 18 (Table 5.5, Fig. 5.4). Concurrently, amino acid content losses of 10.8 $\mu\text{g/mL}$, 15.4 $\mu\text{g/mL}$ and 7.2 $\mu\text{g/mL}$, 11.6 $\mu\text{g/mL}$ were recorded in the *Phormidium* and *Chroococcus* sp. controls on days 6 and 18, respectively.

The *Phormidium* species treated on days 6 and 18 with the lowest concentration (2 $\mu\text{g/ml}$) of metribuzine showed a loss of lipid content at 0.11 and 1.07 mg/g . At the highest concentration (10 $\mu\text{g/ml}$), a lipid loss of 0.43 mg/g and 1.23 mg/g was seen in metribuzin-treated formidium species on days 6 and 18, respectively (Table 5.6; Fig. 5.5). No lipid loss in *Chroococcus* species treated with metribuzine 08 and 0.69 mg/g were recorded at the lowest concentration (2 $\mu\text{g/ml}$) on days 6 and 18, respectively. At the highest concentration (10 $\mu\text{g/ml}$), a loss of lipid content of 0.41 mg/g and 0.84 mg/g was recorded on days 6 and 18, respectively (Table 5.6, Fig. 5.5). At the same time, losses of amino acid content of 0.52 mg/g , 1.24 mg/g and 0.49 mg/g , 0.85 mg/g were recorded in *Phormidium* and *Chroococcus* sp. controls recorded on days 6 and 18, respectively rice field.

Table 5.5 The effect of metribuzin on amino acid in cyanobacteria

S.No	Concentration of Metribuzin ($\mu\text{g/ml}$)	Loss of amino acid in <i>Phormidium</i> sp., ($\mu\text{g/ml}$)					Loss of amino acid in <i>Chroococcus</i> sp., ($\mu\text{g/ml}$)				
		(days)					(days)				
		6	9	12	15	18	6	9	12	15	18
1	2	3.0	4.0	5.7	7.6	11.1	0.7	2.7	5.4	8.1	9.6
2	4	6.5	8.1	10.4	12.2	14.5	0.9	3.8	6.1	8.6	10.0
3	6	7.9	9.0	10.6	13.2	14.8	3.4	5.3	7.0	9.4	10.9
4	8	9.0	10.5	11.9	13.6	15.2	3.9	6.2	7.4	10.4	11.2
5	10	10.1	10.9	12.5	14.0	15.4	5.4	7.4	8.9	10.8	11.4

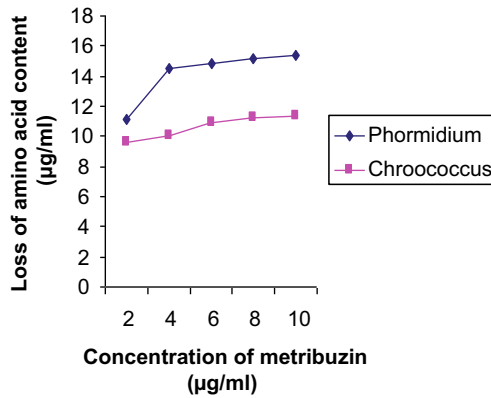
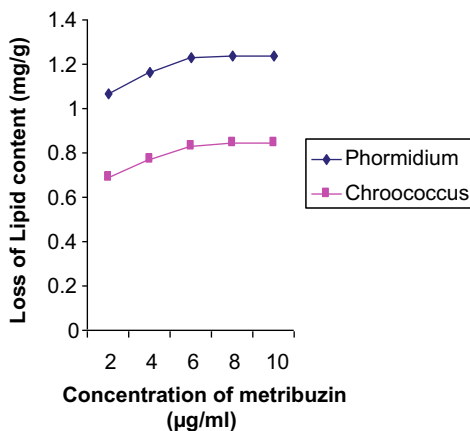


Fig. 5.4 The effect of metribuzin on amino acid content on 18th day

Table 5.6 The effect of metribuzin on lipid in cyanobacteria

S.No	Concentration of Metribuzin ($\mu\text{g/ml}$)	Loss of lipid in <i>Phormidium</i> sp., (mg/g)					Loss of lipid in <i>Chroococcus</i> sp., (mg/g)				
		(Days)					(Days)				
		6	9	12	15	18	6	9	12	15	18
1	2	0.11	0.32	0.41	0.68	1.07	0.08	0.23	0.35	0.53	0.69
2	4	0.22	0.44	0.50	0.75	1.16	0.19	0.33	0.45	0.59	0.77
3	6	0.30	0.55	0.63	0.89	1.23	0.27	0.39	0.54	0.69	0.83
4	8	0.39	0.62	0.68	0.921	1.235	0.35	0.48	0.59	0.74	0.842
5	10	0.43	0.67	0.70	0.925	1.239	0.41	0.53	0.64	0.76	0.848

Fig. 5.5 The effect of metribuzin on lipid content on 18th day



A gradual decrease in chlorophyll-a and biochemical content was observed in *Phormidium* and *Chroococcus* species treated with different concentrations of metribuzine. Their levels were found to increase in controls compared to all concentrations of cyanobacterial species treated with metribuzine. This result confirms that chlorophyll a and biochemical levels were significantly reduced in cyanobacterial species treated with metribuzine at his highest concentration (10 µg/ml) on day 18. The results also indicated that *Phormidium* was more sensitive to metribuzine than *Chroococcus*.

The data obtained in this study also demonstrated that metribuzine is capable of inhibiting the growth and biochemical components of *Phormidium* and *Chroococcus* species even at the lowest concentration (2 µg/ml). Similar observations were reported for haparasiphon with herbicide, where increasing herbicide concentrations decreased levels of chlorophyll, carbohydrates, proteins, amino acids, and lipids (Neveen et al. 2009). Similar results were also reported by his Shabana (1987), who found that the growth and dry weight of *Anabaena oryzae* and *Aulosira fertilissima* were significantly reduced by the application of atrazine. Our present results are in good agreement with those of Megharaj et al. match (1986) showed that herbicides exposed to the cyanobacteria *Nostoc linckia*, *Synechococcus elongatus* and *elongatus*, and his *Phormidium tenue* inhibited total carbohydrate levels even at the lowest concentrations. Other similar observations were made by many workers (Shabana 1985). Similarly, Machete-treated *Anabaena dryorum* showed the greatest reduction in carbohydrate, protein, and lipid content at all concentrations. Higher herbicide concentrations significantly decreased amino acid content (Kashyap and Pandey 1982). Gadkari (1988) reported that the photosynthetic herbicide metribuzine has the ability to inhibit the growth and nitrogenase activity of her *Nostoc muscorum*. *Phormidium* species have been shown to be sensitive to the herbicides 2,4-D, trifluralin, MCPA, and TCA (Cullimore and McCann 1977). Birmingham and Colman (1983) hypothesized that cyanobacteria may be more sensitive to herbicides than

eukaryotic algae. Also Kasai et al. (1993) concluded that, among the seven taxonomic groups, Volvocales and Cyanophyceae are the most sensitive to symmetry. Arvik et al. (1973) reported complete inhibition of cyanobacterial growth in liquid media in the presence of 1 ppm metribuzine.

A number of researchers have also reported the effects of herbicides on cyanobacteria tolerance, growth, and biochemical content (Venkataraman and Rajalakshmi 1971). Similar types of differential effects of different herbicides on the growth and biochemical content of different algae have been previously reported (DaSilva et al. 1975).

Herbicides are organic or inorganic chemicals used primarily to control pests. In high concentrations, these compounds are highly toxic to all plant systems. In addition to killing all target organisms when used in excess, they inhibit the growth of other non-target organisms (Cullimore and McCann 1977).

This study demonstrated the need to pay close attention to herbicide dosage and selection. Because a slightly higher dose of chemicals can kill beneficial microbes. From the foregoing discussion, it is clear that research on herbicide effects may affect the growth and biochemical composition of the cyanobacteria his *Phormidium* and his *Chroococcus* species.

5.4 Conclusion

Most studies of herbicide toxicity to cyanobacteria have been conducted with pure active ingredients. Currently we investigated the effect of marketable products used in the paddy field. The results obtained provided valuable information on the residual inhibitory effects of metribuzine herbicide formulations on the growth and biochemical content of cyanobacterial species. In terms of environmental protection measures, our results demonstrate the continuous and regular application of metribuzine in paddy fields. On the conflicting, we propose to ban the use of metribuzine in paddy fields because metribuzine has a strong inhibitory effect on soil microbiota, especially *Phormidium* and *Chroococcus* species. There is an urgent need to reduce herbicide contamination of the environment using safer, chemical-free pest control methods.

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Chapter 6

Emerging Methodologies for the Molecular Analysis of Soil Microbiota from Polluted Soil Sites



Ridhuvvarshini, Pavethra, Sophia Reena, and Sivaranjani

Abstract The soil microbiome performs a wide range of crucial functions; however, we have a limited understanding of its biodiversity. Extracting microbes from polluted sites could reveal potential microbes that could be used to mitigate pollution better than conventional microbes. Soil DNA may be extracted directly, amplified using polymerase chain reaction, and profiled to reveal more about the soil microbiome's taxonomy and function than ever before. Current procedures frequently combine DNA sequencing with other methods like denaturing gradient gel electrophoresis (DGGE), temperature gradient gel electrophoresis (TGGE), single-strand conformation polymorphism (SSCP), terminal restriction fragment length polymorphism (TRFLP), amplified rDNA restriction analysis (ARDRA), amplified ribosomal intergenic spacer analysis (ARISA), and cloning. The advantages and disadvantages of these methods are discussed, and new developments that have relevance as an appliance shedding light on the soil microbial ecology are also included. Soil diversity cannot be assessed using just one approach; therefore, picking the right one and using newly discovered information can significantly improve our understanding of soil microbes for their specific applications in mitigating.

Keywords Soil microbiome · Bacterial ecology · Polluted soil · Emerging methodologies · Microbial profiling · Direct DNA sequencing

6.1 Introduction

The diversity of the soil microbiome is little understood, even though the soil microbiome performs a wide variety of essential tasks. Direct DNA extraction, amplification by polymerase chain reaction, and profiling of soil microbes have made hitherto undisclosed information on the taxonomy and function of the soil microbiome accessible. It is common practise to combine DNA sequencing with other

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techniques, including DGGE, TGGE, SSCP, TRFLP, cloning, and amplification of rDNA restriction and ribosomal intergenic spacer regions. Innovations that have significance as an appliance shining a light on the soil microbial ecology are also mentioned, along with the pros and cons of these approaches. We can learn much more about soil bacteria if we use the correct method for assessing soil diversity and incorporate newly found information.

Additional advancements are provided by more recent techniques like microarrays and high-throughput sequencing. In this review, various molecular methods used to study soil microbiota are reviewed, along with the pros and cons of each method.

6.2 Bioinformatic Analysis

The initial computational method was based on the sequencing platform, with distinct software modules envisioned for early analysis of diverse systems. However, the programs need to be linked to establish automated “channels” to identify the possible genes and taxonomic groups illustrated by the sequencing data and their relative frequency of occurrence. When one genome is sequenced in short reads, it creates overlapping expansions that allow the “assemblage” of shorter reads into longer ones. When the genomic sequence is presented as a “scaffold,” the job becomes much less complicated. Extended sequences can be matched by searching the NCBI databases. Arranging the various sequences of the initiator in the best possible way is critical for further study. PHYLIP, the PHYLogeny Inference Package, can silently infer phylogenetic trees and sequence family relationships. Genome annotation is assigning biological meaning to DNA sequences and identifying genes. There is a need for software that can build de-novo genomes from short-reads (i.e., less than 50 b), with significant errors projected with extended-reads of 100 b; nevertheless, verifying restricted sequences is challenging. Therefore, it is highly improbable that microbial genomes from diverse and mixed populations can be reconstructed using high-throughput, short-read-length techniques. A mix of tactics catering to readers with varying attention spans (short, medium, and long) would be required. With the expanding breadth of metagenomic research and public databases comes the requirement for high-performance computing and automated applications. Independent and autonomous organizations are developing alternative public pathways to examine metagenomic data. Disparities in metabolic profiles among a few biomes have been uncovered using the mg-RAST server (Meyer et al. 2008; Dinsdale et al. 2008). The sequence was analyzed using BLASTX in addition to ribosomal (GREENGENES, RDP-II), chloroplast, and mitochondrial databases (via ACLAME). MEGAN is a one-of-a-kind method that rapidly assesses the biodiversity of metagenomic samples using visual outputs, allowing researchers to compare and contrast many data sets, such as functional evaluations and metadata (Huson et al. 2007) (Fig. 6.1).

Understanding the subject’s research hot spots and potential future research trajectories can be done using statistics and examining the literature’s subject keywords (Shi et al. 1999). This study employed VOS viewer version 1.6.16 (van Eck and

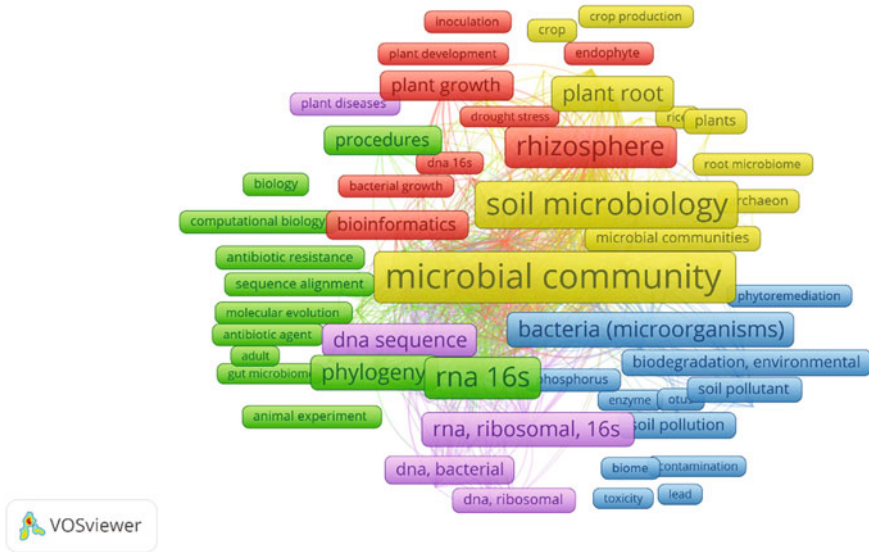


Fig. 6.1 VOSviewer mapping of keywords

Waltman 2010) to assess descriptive data and consider recent research and historical trends. The Scopus database output for the keyword “TITLE-ABS-KEY (Soil microbiota AND molecular analysis)” is also included for co-occurrence analysis in the VOS viewer. The nodes linked with the soil microbial community, soil bacteria, plant-microbe interactions, rDNA, and molecular analysis are deduced from the yellow, blue, red, violet, and green color clusters. The size of the nodes (frames) reveals how frequently they are used, and the separation between the frames and connecting lines reveals how interconnected and linked they are. All nodes are most connected to the cluster of adsorption keywords, the core content, and the most key phrase for all keywords. Figure 6.2 displays the global collaborations identified by the document search. The blue color of the map represents global research collaboration. The pink border that divides the states demonstrates the authors’ level of involvement. It is incredible how the countries that have published the most articles on the soil microbiome have worked together. A generic function summary of bibliometrix analysis done by Bibliometrix R-Tool (Aria and Cuccurullo 2017) for the Scopus database search of a given keyword is provided in Table 6.1.

6.3 Community Profiling Techniques and Limitations

Community fingerprinting techniques have been widely employed in microbiota-ecology research, greatly enhancing our understanding of the variety of soil microbiota. For example, denaturing gradient gel electrophoresis (DGGE), temperature

Country Collaboration Map

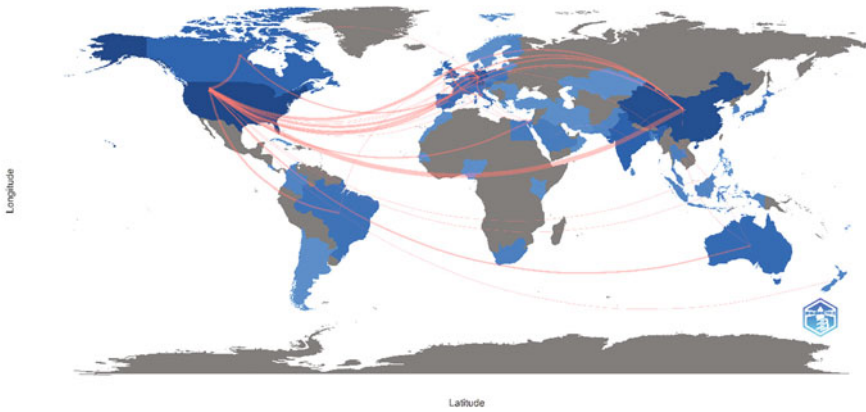


Fig. 6.2 Country collaboration map of research of the title

Table 6.1 Scopus database keyword search and paper hits

Description	Results
<i>Main information about the data</i>	
Timespan	2008:2022
Sources (Journals, Books, etc.)	139
Documents	277
Annual growth rate (%)	30.15
DOCUMENT AVERAGE AGE	3.6
Average citations per doc	52.39
References	20,472
<i>Document contents</i>	
Keywords plus (ID)	3319
Author's keywords (DE)	920
<i>Authors</i>	
Authors	1595
Authors of single-authored docs	6
Authors collaboration	
Single-authored docs	6
Co-authors per doc	6.49
International co-authorships (%)	35.02

(continued)

Table 6.1 (continued)

Description	Results
<i>Document types</i>	
Article	222
Book	2
Book chapter	8
Editorial	1
Erratum	1
Note	1
Review	41
Short survey	1

gradient gel electrophoresis (TGGE), single-strand conformation polymorphism (SSCP), terminal restriction fragment length polymorphism (T-RFLP), amplified rDNA restriction analysis (ARDRA). In addition, amplified tRNA-serine-leucine-nucleotide polymorphism (ART) is a method for identifying (van Elsas 2000).

While each community profiling approach is distinct, many share the same DNA features used in the separating process. Both denaturant gradient gel electrophoresis (DGGE) and thermal gradient gel electrophoresis (TGGE) use the melting behavior of DNA to separate fragments of the same size but different sequences; this is because DNA with different sequences (i.e., different G + C contents) melts at different locations in a polyacrylamide gel (TGGE). Because of the potential for many bands and an exaggerated estimation of diversity, the use of degenerate primers is limited in DGGE/TGGE. Since the advent of DGGE, these methods have become the most widely used community fingerprinting techniques in bacterial ecology, making their advantages and disadvantages clear to anybody interested in studying soil microbial communities. Soil fungal communities may be analyzed rapidly and effectively with DGGE/TGGE, making it a helpful tool for studying changes in community composition (Anderson et al. 2003a). The techniques' benefits include the ability to study and compare several samples on a single gel and the speed and simplicity of such comparisons. Specialized software programs have substantially improved the study of community fingerprints by permitting comparison of the position and relative intensity of distinct bands within gels. This paves the way for further statistical analysis of the data and an appreciation of the ecological conclusions drawn from it. Accurate comparison, however, is highly dependent on suitable internal standards and the prior establishment of gel resolution and quality. The need to compare multiple gels owing to a large number of samples emphasizes the significance of this. One of the fundamental limitations of DGGE is its lack of repeatability across gels, even if the gel-making process may be improved by using, for example, the same flow rate of denaturant solutions and the same equipment for each gel (Fromin et al. 2002).

The methods used to create such profiles of a whole community are not without their flaws. Compared to more oversized products, smaller DNA fragments (500 bp)

provide better resolution across bands in a profile, limiting the taxonomic information that can be gleaned by sequencing excised bands (Landeweert et al. 2003). More importantly, even the most sensitive staining techniques sometimes fail to identify all the variability that occurs in a sample, especially for the less prominent members of the group. It has also been shown that a single band on a gel may contain many sequence types (Schmalenberger and Tebbe 2003). T-RFLP uses automated DNA sequencing technology to provide significantly better throughput than gel-based community profiling approaches (Marsh 1999). T-RFLP varies from conventional RFLP in that it employs fluorescently tagged PCR primers (forward, reverse, or both primers) before restriction digestion and size measurement of fluorescently labeled terminal restriction fragments using a DNA sequencer. T-RFLP may simplify the community profile without compromising the variety found by identifying the last segment of each 18S rDNA or ITS sequence in a sample. T-RFLP is an improvement on the ARDRA method of community profiling. There are methods like ARISA, which are similar, but they assess whole amplicons instead of simply the terminal fragments that result after restriction digestion (Leckie et al. 2004). Using an automated DNA sequencer for ARISA and T-RFLP increases throughput and improves accuracy in sizing the generated fragments by adding an internal standard in each sample and maintaining stable operating parameters. Although T-RFLP has been used to analyze the diversity of soil fungi and to detect ECM fungi in soil samples (Lord et al. 2002; Dickie et al. 2002), a robust T-RFLP database is necessary for identifying individual fungi species. The lack of ease with which sequence information may be extracted from T-RFLP peaks makes identifying previously unknown species in a sample a formidable task. Occasionally, artificial identification has been made using virtual restriction digests based on public database sequences; however, this is risky because it presumes that only one species or operational taxonomic unit (OTU) can have a peak of that size in a sample and that the database sequence is correctly identified and of high quality.

Cloning PCR amplicons from environmental DNA has also been used to evaluate soil fungal diversity (Jumpponen 2003). However, clones may be screened using RFLP to categorize clones into OTUs prior to DNA sequencing, reducing the number of clones required for sequencing and making it difficult to determine how many clones must be analyzed to sample the diversity contained in a single sample effectively. The development of collectors or species abundance curves shows that the number of 18S rDNA clones that need to be analyzed in agricultural soil is far smaller than the number of clones that would have been required for a library of bacterial 16S rDNA to have covered the diversity microbiota (Anderson et al. 2003). These researchers found that more ITS clones than 18S rDNA clones from the same sample needed to be screened to ensure coverage of the fungal diversity in the sample. Without analyzing many clones, it is difficult to discriminate between familiar and unusual sequence types in a sample.

DNA sequencing and phylogenetic analysis are commonly used in tandem with community profiling methods for the taxonomic identification of species present in a sample; however, it is crucial to remember that environmental DNA samples may also contain chimera DNA sequences. The techniques mentioned above provide an

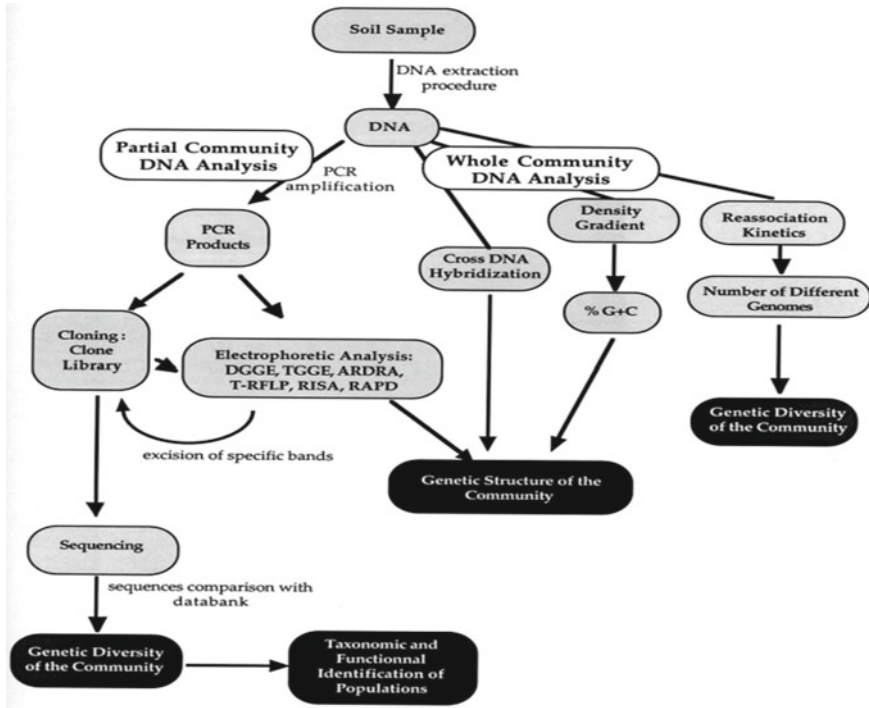


Fig. 6.3 Schematic representation of the different molecular approaches

array of adequate resources for assessing soil fungus populations. Although each has certain technical limitations that prevent it from being fully utilized, its widespread application in bacterial ecology over the past decade has led to steady development and refinement. Each community profiling method varies in its taxonomic resolution due to factors such as the PCR primers chosen for the first PCR amplification of the community DNA and the lack of extensive sequencing information in public databases. Fungus ecologists have been working on loosening these constraints for years, and their efforts will likely bear fruit soon (Figs. 6.3 and 6.4).

6.4 Current Developments in Molecular Ecology of Soil Fungi Profiling

The spatial and temporal dynamics of resident fungus, as well as the diversity of fungi in plant roots, the rhizosphere, and/or bulk soil, have been considerably improved by recent investigations of soil fungal ecology employing PCR amplification from total extracted DNA.

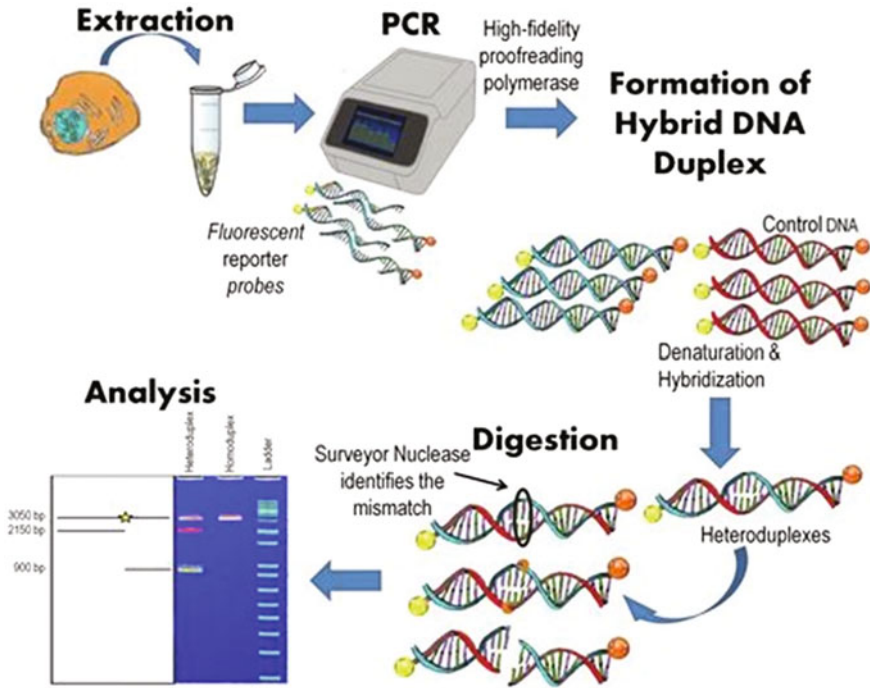


Fig. 6.4 Extraction and analysis of genetic information of soil microbes

Support for the long-held belief that growing fungi from the soil only allows for a limited perspective of variety was provided by the cloning and sequencing of 18S and ITS rDNA PCR products from total soil DNA (Hunt et al. 2004). Using 18S rDNA amplification in conjunction with TGGE and DGGE in both microcosms and field soil, we show that fungal diversity is lower in the rhizosphere compared to that in bulk soil, and that there is an influence of plant age. The bare soils around glacier ice edges and terminal moraines contain unique fungal communities, as shown by an analysis of 18S rDNA clones. Soil community composition has been shown to shift in response to various treatments, including petrochemical pollution, nitrogen addition, controlled vegetation burning, elevated atmospheric CO₂ concentration, and the addition of compost or manure, as determined by studies employing 18S or ITS rDNA amplification coupled with cloning, DGGE, SSCP, or T-RFLP analysis. The DGGE-detected communities lacked the taxonomic diversity of the T-RFLP-detected communities. However, because several DNA extraction methods and primers were employed, it is unclear to what extent these differences reflect variations in the relative resolution of the approaches.

However, the taxonomic richness of the DGGE profiles was similar to that of an isolated fungal assemblage from the same host, despite sequencing differences. While *Phoma* and *Microdochium* were predicted to be major players in the *A. arenaria* root fungal community based on isolated assemblages, they were not found using DGGE.

Additional research using 18S rDNA PCR and DGGE on *A. arenaria* endophytes demonstrated that the taxonomic structure of AM fungal assemblages in roots and that inferred from spore diversity are unrelated. It has been common practise to directly amplify AM fungus from whole root DNA for quite some time. Testing for AM infection in roots using competitive PCR has been shown to be doable. While it has been around for some time, it has just lately been put to use to resolve pressing ecological concerns. Helgason and coworkers (1998) used partial 18S rDNA PCR from total root DNA and cloning to compare AM fungal diversity in the roots of monoculture arable crop plant hosts and forest plant species. It was believed that the prevalence of AM fungus, which can infect a wide range of host plants, was not due to monoculture planting but rather an effect of agronomic management approaches on diversity. Different AM communities have been found in the roots of several plant species, however, according to a different study that used the same techniques in combination with T-RFLP (Vandenkoornhuysen et al. 2003). More evidence that floristic diversity may have a substantial effect on endophyte diversity was provided by Johnson and colleagues (2004), who compared the AM fungal assemblages in the roots of *Plantago lanceolata* bait plants in microcosms grown in monoculture and mixed plant communities. They were able to do so by employing a PCR and T-RFLP technique based on an 18S rDNA subset.

6.5 Popular Approaches

Community profiling techniques have recently made significant strides in soil mycology. These methods are already being used in labs all over the world, and it is expected that our knowledge of the composition and evolution of soil fungal communities will expand dramatically in the near future as a result. But researchers shouldn't just choose any old set of primers or community profiling technique to assess fungal diversity; they should weigh the pros and cons carefully in light of their goals, the fungi of interest, and the data they want to analyse. In addition, we have emphasised in this review the limits of the methodologies that must be taken into account in the context of individual studies and their respective hypotheses. Some of these restrictions apply across all fields of ecology, whereas others have been addressed in soil bacterial community investigations, leading to improved methods. Diversity analysis based on taxonomic richness and relative abundance, along with community dynamics, is far more difficult than analysing, say, the taxonomic richness of soil fungus. This is because fungi are mycelial in structure, and spores are likely to coexist in any given sample. The technique of stable isotope probing (SIP), which classifies organisms according to their nucleic acids' abilities to use certain stable isotope-marked substrates, has been successfully applied to the separation of soil bacteria into their respective functional groups (Morris et al. 2002). In particular, its benefits to soil fungus have just lately been recognised. These scientists present a convincing case for using this method to investigate microbial DNA in soil. Our findings show that SIP may be used to examine fungal RNA, which integrates ¹³C quicker

than DNA. This is in contrast to bacterial DNA, which incorporates carbon isotopes more slowly. In a practical setting, analysing fungal RNA obtained from soil enables the detection and identification of metabolically active community members rather than dormant individuals. Consequently, this is a major development in our understanding of fungus and the ecosystems in which they exist. Fungal DNA and RNA levels in soil have also been quantified using real-time PCR (Lueders et al. 2004). Opportunities for further development of metagenomics techniques and microarray technology to study the composition and activity of microbial communities seem promising. These methodologies, when used in tandem with community profiling techniques, allow us to get a more nuanced understanding of fungal communities and the roles they play in the ecological processes of soil.

6.6 Isotopic Labeling

This can be done with the use of circumstantial evidence, such as when scientists try to link a change in the abundance of a certain group or set of genes with the event in question. By using isotopes, substrates may be individually verified for membership in a certain category (Dumont and Murrell 2005). Micro-autoradiography (MAR) requires the introduction of radiolabeled substrates or nucleotides, such as ³H-labelled thymidine, into the soil. While FISH applied to soil may allow for the identification of live cells in their natural setting, it suffers from the same limitations as other direct imaging techniques (Wagner et al. 2006; Rogers et al. 2007). In order to better identify certain subpopulations, it is encouraged to label soil-extractable PLFAs or nucleic acids. When methane oxidizers in soil are exposed to ¹³CH₄, stable isotope probing (SIP) is most often observed in the interaction of ¹³C with PLFAs or DNA (Bull et al. 2000; Radajewski et al. 2000). Growing cells take up ¹³C from certain substrates in their DNA and rRNA. In order to separate the ¹³C-enriched fraction from the unlabeled fraction of isolated soil nucleic acids, density gradient centrifugation can be used. Methods for SIP are outlined by Kreuzer-Martin (2007); these techniques are useful for examining separate groups that make use of specific substrates like methane, and they may also give more broad information on the members of actively forming soil communities. Many soil communities, for instance, have been studied using various ¹³C substrates in conjunction with DNA and RNA-SIP (Manefield et al. 2002; Lueders et al. 2004; Whiteley et al. 2006).

6.7 BrdU Labeling

The thymidine nucleotide analogue 5-bromo-2-deoxyuridine (BrdU) can be used alone or in conjunction with an exogenous substrate to mark dividing soil cells. Using any of the current community DNA analysis techniques, DNA from the less active minority may be compared to DNA from cells containing BrdU. In soils from several

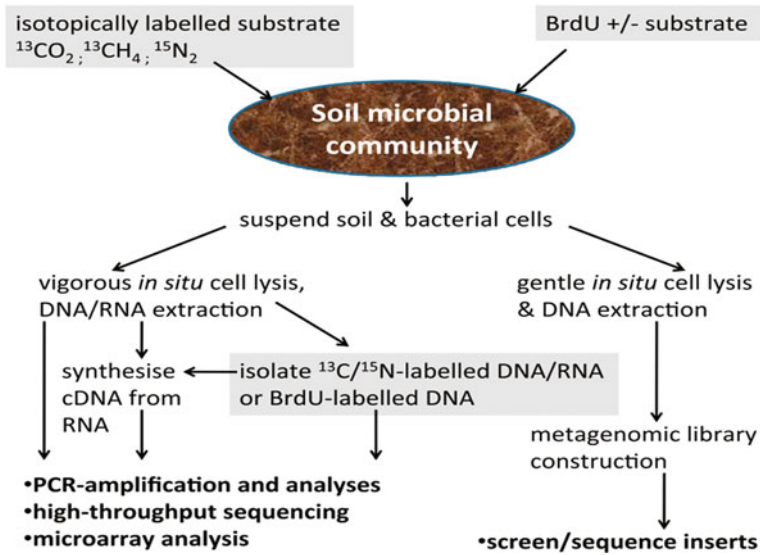


Fig. 6.5 Flow chart showing DNA and RNA extraction from soil. The soil microbial community will include bacteria, archaea, fungi, protists, and microfauna

sites that were incubated at either 4 or 25 °C, we found that BrdU was useful for assessing both the total and growing communities (Figs. 6.5 and 6.6). The data reveal that, in contrast to the profiles generated from actively reproducing bacteria that absorbed BrdU, whole-genome DNA profiles do not cluster according to soil type or incubation temperature. Other possible uses for BrdU include identifying members of disease-resistant plant-related active microbial communities in soil and identifying bacteria connected with arbuscular mycorrhizal fungus hyphae (Artursson and Jansson 2003). Researchers from many institutions collaborated on the project (Hjort et al. 2007).

6.8 The Potential of Microarrays in Soil

Microarrays have been constructed in recent years using sequence data from publicly available bacterial genomes deposited in databases like GenBank at the NCBI. Recent literature on the use of microarrays in soil microbial ecology (Sessitsch et al. 2006). The PhyloChip, designed by Gary Andersen and his colleagues at the Lawrence Berkeley National Lab in the United States, is a phylogenetic microarray that uses databases of 16S rRNA gene sequences. With a total of 500,000 probes, this high-density array has the potential to identify w9000 unique species or taxa. To reflect the ever-evolving nature of DNA databases, these microarrays are routinely

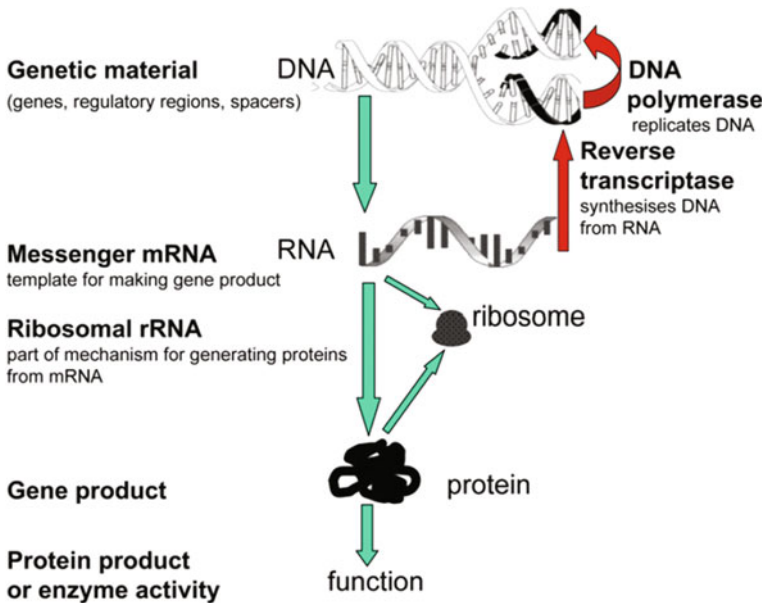


Fig. 6.6 The relationship between DNA, RNA forms, protein functions, and pathways

upgraded to include the most recent data. Some researchers believe that these techniques, as opposed to traditional cloning/sequencing procedures, may show a greater diversity of bacterial phyla and species in soil (DeSantis et al. 2007). In addition to showcasing the richness of ecosystems, these arrays can provide insight into how various taxa are represented within a given treatment, such as rhizosphere vs. bulk soil (DeAngelis et al. 2009). When it comes to identifying the people who make a community tick, rRNA may soon supplant genomic DNA as the gold standard. While it can provide some insight into the composition of known microbial communities, the array does not include all of the functional genes involved for the major activities in soil. The GeoChip, created by Jizhong Zhou at the University of Oklahoma in the United States, is based on functional gene sequences, whereas phylochips are based on non-functional gene sequences. By employing GeoChip 2, for instance, researchers have shown that the number and variety of genes involved in organic C degradation in soils is on the rise. GeoChip 2 has the potential to uncover over 10,000 genes in over 150 different functional areas. To wit: (Zhang et al. 2007). Functional genes in several species may be identified using probes, yielding information on biodiversity. The already outstanding identification rate of 47,000 genes across 292 gene families utilising the new and improved GeoChip 3 is expected to rise as additional sequencing data becomes available. The RNA that is collected from soil can provide information about which genes were active right before sampling, which is something that DNA cannot do. Microarrays, on the other hand, rely on already known DNA sequence information, therefore they are unable to identify new groups

that may become critical to particular functions. Because so little is known about the taxonomic diversity and functional gene sequences of soil fungi, microarrays for soil fungal communities are also now unavailable.

6.9 High-Throughput Sequencing

Traditional cloning and Sanger sequencing take a long time, thus only a small number of samples may be processed at once. Automatic diagnostic techniques (sometimes known as “lab-on-a-chip” technology) are now under development. Utilizing these techniques, researchers may now analyse hundreds of samples to study microbial functional diversity. Thanks to recent advancements in massively parallel high-throughput pyrosequencing techniques, hundreds of thousands of sequences may be processed in parallel. At now, the “GS FLX Titanium Series” machine can sequence one million fragments with an average read length of 400 bp; however, this will increase to 800 bp in the near future. Information on the most recent models may usually be found on company websites, and other commercial high-throughput sequencing equipment offer a more extensive variety of shorter sequences. In comparison to the Applied Biosystems Inc. SOLiD3 system, which promises to sequence 10–15 Gb every 6–7 day run with reads of up to 50b, the Illumina Inc. Genome AnalyserIIX, as of this writing, offers to sequence 20–25 Gb per 9–10 day run. With the fast development of new technologies, new approaches like nanopore sequencing are being created (Mardis 2008). In a 2009 study (Stoddart et al. However, weighing the benefits and drawbacks of each approach is outside the scope of this study and is being done on several online forums. In order to alter the DNA that has been collected from soil, PCR amplification with gene-specific primers can be performed. The relative abundance of bacterial and archaeal 16S rRNA genes (the mean from four soils was w140,000 bacterial; w9000 archaeal sequences) and the total taxonomic diversity of these species in soil were determined using universal 16S rRNA gene primers, for example (Roesch et al. 2007). This approach has the limitation that it assumes all prokaryotes have 16S rRNA gene sequences comparable to the primers used in the PCR amplification stage. However, many bacteria and archaea are able to be cultured despite deviating from the standard primer sequence for this gene. But the technology will develop into a more useful tool as more and more critical pieces may be scheduled with each iteration. Recent advances in high-throughput sequencing have rendered direct and unbiased sampling unproductive because of their inability to unearth uncommon but ecologically significant categories without prior selection. In the future, only the bioinformatic processing of sequence data is anticipated to limit the potential expansion of sequencing, which is now useful and will become much more so as more advanced methods are developed (Pop and Salzberg 2008).

6.10 Conclusion

There are obstacles to exploring the microbiota due to the large number and variety of soil microbiota and the varied nature of the soil, irrespective of the site being polluted or otherwise. Results from small samples that may be used for the study have made traditional protocols unappealing; nevertheless, cutting-edge methods increase the number of soil samples included. Understanding the fundamental mechanisms of each method for investigating microbiota diversity and activity is crucial for mitigating the demerits and capitalizing on the strengths of the various approaches.

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Chapter 7

Biodecoloration of Synthetic Reactive Red and Reactive Black Dyes by Using *Aspergillus niger* and *Pleurotus ostreatus*



P. Muthukumaran, S. Sridhar, and J. Aravind

Abstract The present study is alternative for removal of synthetic dyes particularly reactive red 120 and reactive black 5 dyes by *Aspergillus niger* and *Pleurotus ostreatus* by shake flask system for 7 days. The efficiency of decoloration of reactive red and reactive black, at room temperature (28 ± 2 °C), by *Aspergillus niger* are 78% and 66% respectively. Similarly, *Pleurotus ostreatus* shows 64 and 77% of decoloration for reactive red 120 and reactive black 5 respectively. The optimization of the decolorization efficiency was then carried out for the fungal colonies. The various physical parameters such as pH (3, 4, 5 and 6), temperature (30, 37 and 45 °C) and C:N ratio (1:2, 1:1, 2:1) were optimized to obtain maximum decolorization of the commercial reactive dyes by *Aspergillus niger* and *Pleurotus ostreatus*.

Keywords Reactive red 120 · Reactive black 5 · *Aspergillus niger* and *Pleurotus ostreatus* · Decolourisation

7.1 Introduction

Textile dye effluents pose environmental hazards because of color and toxicity. All dyes used in the textile industry are designed to resist fading upon exposure to sweat, light, water, many chemicals including oxidizing agents and microbial attack. In literature, there have been reported many fungal strains that are able to degrade dyes by their capacity to produce a range of enzymes (extra cellular enzymes like laccase, phenol oxidase, aryl oxidases, peroxidase, etc.). To name a few, fungal strains used in dye degradation are white rot fungus, *Phanerochaete chrysosporium*, *Geotrichum candidum*, *Trametes versicolor*, *Bjerkandera adusta*, various species of *Penicillium*, *Pleurotus ostreatus*, *Pycnoporus cinnabarinus*, *Pyricularia oryzae*,

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Aspergillus niger, etc. which can decolorize and/or biosorb diverse dyes (Martorell et al. 2017; Kaushik and Malik 2009a).

Evaluation of indigenous soil fungi for decolourization of textile dyes, Microbial decolourization of textile dyes through isolates obtained from contaminated sites (Raju et al. 2008; Kaushik and Malik 2009b) and decolorisation of reactive red 11 and 152 azo dyes under aerobic conditions reported by Kodam and Gawai (2006). Colour removal of textile dyes by culture extracts obtained from White- rot fungi was reported by Mehmet et al. (2009).

Large literature exists regarding the potential of these fungi to oxidize phenolic, non-phenolic, soluble and non-soluble dyes (Heinfling et al. 1998). In particular laccase from *Pleurotus ostreatus*, *Schizophyllum commune*, *Sclerotium rolfsii* and *Neurospora crassa*, seemed to increase up to 25% the degree of decolorization of individual commercial triarylmethane, anthraquinonic, and indigoid textile dyes using enzyme preparations (Siripong et al. 2009). This study focuses on the biological decolourization of textile effluents through microbial isolates obtained from contaminated sites. The fungal colonies were isolated from the effluent samples and the soil from the contaminated sites (Feng et al. 2014).

7.2 Materials and Methods

The dye used in this study was reactive red 120 and reactive red 5 dye obtained from Department of Textile Technology, Kumaraguru College of Technology, Coimbatore. The effluent was collected from a textile unit located in Erode, Tirupur and Salem district, Tamil Nadu, India.

7.2.1 Sample Collection

The dye effluent samples and soil sample from the dye contaminated sites were collected from various textile industries located in Perundurai, Tirupur and Salem district, Tamil Nadu. The isolation of fungal colonies from these samples was then carried out.

7.2.2 Isolation of Fungal Colonies from the Collected Effluent and Samples

The effluent dye samples were serially diluted and fungal colonies are isolated by using Potato Dextrose Agar (Hi-Media) plates by pour plate techniques and plates were incubated at room temperature (28 ± 2 °C) for 5 days (Ahmed et al. 2018).

7.2.3 Decolourization Studies

The isolated bacterial and fungal colonies were then studied for their ability to decolorize the effluent dyes. This assay was measured in the terms of reduction percentage using a spectrophotometer (Dos Santos et al. 2007).

Dye removal (%) was calculated as

$$\text{Dye removal (\%)} = \frac{\text{Initial absorbance} - \text{Final absorbance}}{\text{Initial absorbance}} \times 100$$

The dye removal % was calculated after 48 h and the fungal colonies with high decolorization efficiency were selected.

7.2.4 Decolorization Studies on the Isolated Fungal Colonies by Using Minimal Media

The decolorization studies were carried out for the isolated fungal colonies on a specialized media called Bushnell & Haas medium of the following composition (g/L): NH_4NO_3 , 0.5; $\text{MgSO}_4 \cdot 7\text{H}_2\text{O}$, 0.1; K_2HPO_4 , 0.5; NaCl, 1 g/L; glucose, 10; yeast extract, 2% and pH 6. Similarly, two synthetic dyes namely Reactive red 120 and reactive black 5. These dyes were prepared at a concentration of 50 mg/100 mL and the decolorization assay mixture consists of 10% of dye. The samples were withdrawn after 7 days and were centrifuged at 10,000 rpm for 15 min. The Absorbance of the supernatant was measured by using UV-visible spectrophotometer (ELICO, India).

7.2.5 Optimization of Decolorization of the Synthetic Dyes by Fungal Strains

The optimization of the decolorization efficiency was then carried out for the fungal colonies. The effect of pH (3, 4, 5 and 6), Temperature (30, 37 and 45 °C), C:N ratios (1:2, 1:1, 2:1) were studied and these parameters are optimized to obtain maximum decolorization of the commercial dyes using the fungal colonies.

7.3 Results and Discussion

7.3.1 Isolation of Fungal Colonies from the Effluent and Soil Samples

Totally about 13 different effluent and soil samples were collected from Perundurai, Tirupur and Salem district, Tamil Nadu. About 8 different fungal colonies were obtained from the samples. Out of 8 different fungal strains two are identified as *Aspergillus niger* and *Pleurotus ostreatus*.

7.3.2 Decolorization of the Synthetic Dyes Using the Selected Fungal Cultures

The screened fungal strains were then tested for their ability to decolorize the synthetic dyes. Two synthetic dyes namely reactive red and reactive black were used for these studies. The synthetic dye removal % of the screened by two fungal strains (Table 7.1).

7.3.3 Optimization of the Decolorization Efficiency for the Selected Fungal Strains

The optimization of the decolorization efficiency was then carried out for the screened fungal strains such as *Aspergillus niger* and *Pleurotus ostreatus*. The effect of various physical parameters such as pH, Temperature, C:N ratio and shaking conditions were studied. The optimum conditions for the decolorization of the synthetic dyes was determined (Kameche et al. 2022).

Table 7.1 Synthetic dye removal % of the fungal cultures—after 7 days

Culture name	Dye removal % Reactive red 120 (520 nm)	Dye removal % Reactive black 5 (574 nm)
<i>Aspergillus niger</i>	78	66
<i>Pleurotus ostreatus</i>	64	77

Table 7.2 Effect of temperature on the decolorization efficiency—after 7 days

Culture name	% Dye removal Reactive red 120 (520 nm)	% Dye removal Reactive black 5 (574 nm)
Incubation temperature: 30 °C		
<i>Aspergillus niger</i>	78	40
<i>Pleurotus ostreatus</i>	64	77
Incubation temperature: 37 °C		
<i>Aspergillus niger</i>	51	35
<i>Pleurotus ostreatus</i>	34	55
Incubation temperature: 45 °C		
<i>Aspergillus niger</i>	16	22
<i>Pleurotus ostreatus</i>	21	17

7.3.4 Effect of Temperature on the Decolorization Efficiency

The effect of temperature on the decolorization of the commercial dyes was analyzed at three different incubation temperatures 30, 37 and 45 °C. The decolorization % was calculated for the samples incubated at different temperatures and the following table gives the decolorization % after 7 days for the two fungal strains (Agrawal and Verma 2019) (Table 7.2).

The maximum dye removal was found to occur at the incubation temperature of 30 °C. Thus, the optimum temperature for decolorization for the three synthetic dyes by the fungal strains was found to be 30 °C (Ali and El-Mohamedy 2012).

7.3.5 Effect of pH on the Decolorization Efficiency

The effect of pH on the decolorization of synthetic dyes by the fungal strains was analyzed. The decolorization media was prepared at different pH 3, 4, 5 and 6 and the assay mixture decolorization % was calculated. The following table gives the synthetic dye removal % of the four fungal strains at different pH (Sosa-Martínez et al. 2020) (Table 7.3).

The maximum dye removal was found to occur at pH 6. Thus, the optimum pH for the decolorization of the synthetic dye by the fungal strains was determined to be 5.

Table 7.3 Effect of pH on the decolorization efficiency—after 7 days

Culture name	% Dye removal Reactive red 120 (520 nm)	%Dye removal Reactive black 5 (574 nm)
pH 3		
<i>Aspergillus niger</i>	17	15
<i>Pleurotus ostreatus</i>	22	10
pH 4		
<i>Aspergillus niger</i>	36	45
<i>Pleurotus ostreatus</i>	31	48
pH 5		
<i>Aspergillus niger</i>	78	61
<i>Pleurotus ostreatus</i>	64	77
pH 6		
<i>Aspergillus niger</i>	56	45
<i>Pleurotus ostreatus</i>	38	48

7.3.6 Effect of C:N Ratio on the Decolorization Efficiency

The effect of the C:N ratio on the decolorization efficiency was determined by using the decolorization media with different Carbon:Nitrogen source ratios such as 1:2, 1:1, 2:1. The following table gives the dye removal % of the fungal strains at different C:N ratios (Table 7.4).

The maximum dye removal % occurred at the C:N ratio of 1:2. Thus, the optimum C:N ratio for synthetic dye decolorization was determined to be 1:2 for the fungal strains (Mahmoud et al. 2017).

Table 7.4 Effect of C:N ratio on the decolorization efficiency—after 7 days

Culture name	% Dye removal Reactive red 120 (520 nm)	% Dye removal Reactive black 5 (574 nm)
1:1		
<i>Aspergillus niger</i>	11	14
<i>Pleurotus ostreatus</i>	12	22
2:1		
<i>Aspergillus niger</i>	29	52
<i>Pleurotus ostreatus</i>	36	21
1:2		
<i>Aspergillus niger</i>	56	85
<i>Pleurotus ostreatus</i>	48	51

7.4 Conclusion

Environmental pollution caused by the release of a wide range of compounds as a consequence of industrial progress has now assumed serious proportions. Textile dyes are one of the most prevalent chemicals in use today. With increasing usage of the wide variety of dyes in the industries pollution from the effluents has become increasingly alarming. Microbial decolorization and degradation is an environmentally friendly and cost-effective process. The fungal strains were isolated from the effluent and soil samples from the contaminated sites. These strains were subjected to initial screening through the decolorization studies on effluent dyes. The screened strains were then tested for their ability to decolorize three synthetic dyes Reactive Red 120 and Reactive Black 5 then final screening was done based on these studies. After final screening, 2 fungal strains were selected. The fungal strains were identified as *Aspergillus niger* and *Pleurotus ostreatus*. The various physical parameters such as temperature, pH, C:N ratio were optimized for the decolorization of these two synthetic reactive dyes by *Aspergillus niger* and *Pleurotus ostreatus*.

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Chapter 8

Feasibility and Performance Evaluation of *Mentha Aquatica* in Treating Domestic Wastewater



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Abstract For life to exist, water is essential. Several conventional methods and combinatorial approaches are in practice to clean wastewater, which needs to be maintained. Additionally, wasted water must be treated and used for various purposes. Domestic wastewater is defined as water used for activities such as dish washing, laundry, and house cleaning. Different kind of ways to treat household wastewater. It can be purified using various techniques, including physical, chemical, biological, and natural processes. Physical methods require expert or qualified personnel for maintenance and control, along with the best equipment for use. Each tactic uses various techniques. The chemicals used in chemical wastewater treatment methods can cause harm to people. However, biological treatment requires both aerobic and anaerobic systems. *Mentha aquatica* was used in this study as a natural source for treating domestic wastewater. The chemical oxygen demand (COD), biological oxygen demand (BOD), and turbidity measurements were used to evaluate the water quality. It was noted that the chemical oxygen demand and biological oxygen demand readings had significantly decreased on days 11 and 21 of treatment. Thus, the observed data confirm *M. aquatica* as a potential replacement for current wastewater treatment methods.

Keywords Wastewater purification · Aquatic plants · Natural methods · BOD and COD reduction · Removal efficiency

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8.1 Introduction

Germ and hazardous contaminants found in sewage are capable of causing serious harm to the ecosystems and living beings through infections, severe illnesses and sometimes lethal (Muthukumar et al., 2022, 2021a, b; Mahesh et al., 2022a, b). A high concentration of pollutants in wastewater decreases the water's quality and its nutrient compositions and assists in the spread of the disease (Priyadarshini et al., 2022; Anandaraj et al., 2018; Kanmani et al., 2017). An estimated 70% of the country's pure water sources are said to be tainted with bacteria. Additionally, India's water quality scores 120 of 122 countries, which is quite low. Due to microbiologically polluted water, several diseases, including cholera, diarrhoea, dysentery, hepatitis A, typhoid, and polio, can spread. This contamination causes significant changes in variables including pH, biological oxygen demand (BOD), chemical oxygen demand (COD), and the number of heavy metals in the water. Heavy metals are released into the environment from the gamut of process industries like food and pharmaceuticals, cosmetics, leather, and other chemical processing industries (Mahesh et al., 2022b; Manikandan et al., 2022). Water contamination can result in fatal complications from the release of industrial chemicals, poor maintenance, oil spills, sewage overflow, and other causes (Muthukumar et al., 2021a, b). The contaminated water is highlighted as the cause of the increased rate of ailments. The WHO estimates that water contamination causes more than 4 lakh diarrhea-related deaths each year. Before this contaminated water from diverse sources may be used for domestic tasks like drinking, irrigation, farming, gardening, and toilet flushing, it must first be treated.

To clean up contaminated wastewater, numerous physical, chemical, biological, and natural techniques are available (Magar and Ansari, 2017; Gupta et al., 2012). This research aims to develop a natural approach for cleaning contaminated household water that can be applied to many tasks, including cleaning, gardening, and other things. Natural purifying agents, such as plants, are thought of as environmentally friendly ways to purify water because they unintentionally filter water (Foroughi et al., 2010; Dahija et al., 2019). Plants from both the land and the sea can be utilised as filters. Aquatic plants including hyacinth, azolla, duckweed, cattail, lotus (Abd Rasid et al., 2019) and poplar are frequently employed for wetland water remediation because of their high accumulation of heavy metals, high tolerance, or rapid growth and high biomass output (Musa and Idrus, 2021; Mustafa and Hayder, 2021). This filtration process has no negative side effects because it is a natural process. The use of water hyacinth for wastewater treatment in California, Florida, Massachusetts, and Texas are examples of this (Télléz et al., 2008; Lateef et al., 2013). Chemical methods involve adding purifying chemicals to the water. Chemical procedures include the removal of heavy metals, chelation, adsorption, and chlorination (Kumar et al., 2019; Raskin et al., 1997). Physical procedures, including filtration, distillation, and sedimentation, are also available (Dayana and Suresh Babu, 2021; Leiknes, 2009).

Depending on the industrial, agricultural, and municipal wastewater releasing operations, the contaminants found in the wastewater may vary. Pollutants are categorized into three major classes as listed inorganic, organic, and biological contaminants. Heavy metals that are toxic and carcinogenic are the most prevalent inorganic water pollutants (Balakumar et al., 2022; Lakherwal, 2014). Other compounds that can be quite harmful include nitrates, sulfates, phosphates, fluorides, chlorides, and oxalates. Fungicides, insecticides, and herbicides are the sources of the hazardous organic pollutants. Other polynuclear hydrocarbons (PAHs) that can contaminate materials include phenols, polychlorinated biphenyls, halogenated aromatic hydrocarbons, formaldehyde, poly-brominated biphenyls, biphenyls, detergents, oils, and greases. Regular hydrocarbons, alcohols, aldehydes, ketones, proteins, lignin, drugs, and other compounds are also present in the wastewater. Various bacteria that flourish in sewage may be the root of many diseases. Some of the potentially harmful microorganisms include bacteria, fungi, algae, plankton, viruses, and other worms. These pollutants in the water remain as suspended, solvated, or colloidal particles (Sudarsan et al., 2015). Aquatic plants that filter water already exist can eliminate these contaminants. According to a study (Avelar et al., 2014) evaluating the performance of the species *Mentha aquatica* in constructed wetlands of horizontal subsurface flow with regard to the removal of coliforms bacteria in an effluent from the primary treatment of sewage, the results are inferred to show positive results. As a result, the treatment of domestic waste water can be considered.

8.2 Materials and Method

8.2.1 Sample Collection

A *Mentha aquatica* plant sample was obtained from a nearby nursery and corroborated at the Central Council for Research in Siddha. The residential wastewater that was obtained after washing clothes and dishes was collected in a glass tank that measured 5 feet by 5 feet.

8.2.2 Wastewater Analysis

The amounts of pH, BOD, COD, turbidity, chlorine, nitrate (NO_3^-), and sulfate were measured in the collected residential wastewater sample. A pH meter is used to measure the pH, and a LAQUA twin nitrate meter is used to measure the nitrate (Saleh et al., 2020; Priya et al., 2012).

8.2.3 Determination of COD

The amount of oxygen, measured in milligrams, needed to stabilize or oxidize the oxidizable compounds contained in one litre of effluent under certain circumstances is known as the chemical oxygen demand of an effluent. A 2.5 mL tube was filled with 2.5 mL of the sample, 1.5 mL of potassium dichromate (0.25 N), a spatula of mercuric sulfate (HgSO_4), and 3.5 mL of COD acid. The tube was then heated to 150 °C in a COD reactor for two hours. After cooling, the sample was titrated with an indicator and standard ferrous ammonium sulfate (0.1 N). The last point is reddish brown in colour. The same procedure was used to create a blank sample using 2.5 mL of distilled water (Islam et al. 2014).

8.2.4 Determination of BOD

Prepare the deionized water-based 2000 mg/L BOD standard solution. WTW company provided BOD standard powder. The medium for the serial BOD concentrations was the solution. Apply the following equation to the designated BOD medium while adding the BOD solution at concentrations ranging from 0.0 to 1000 mg/L. Inoculation with natural or modified sludge is carried out after the addition of 164 mL of baseline medium and 50 mg/L of standard solution (C1) to 2,000 mg/L of C2 V2 in a BOD container (Inoculums). The OxiTop S12 digital electronic detectors are used to recover every bottle, and they are then placed in a rack shaker in an incubator for five days at 20 °C (Devi and Dahiya, 2008).

8.2.5 Turbidity Determination

Try to wait until the samples are at room temperature before analysing them if the turbidity is less than 40 units. Mix the sample thoroughly for even distribution of the solids. Pour the substance into the turbid meter tube once all the air bubbles have disappeared. The turbidity can be calculated using either the instrument scale or the proper calibration curve.

8.2.6 Phosphate Determination

Before adding it to the cell, the reaction mixture was boiled and cooled. The supporting electrolyte was made up of saturated ammonium acetate and acetic acid (5 mL each), was added to the combination of 10 mL. As opposed to a standard calomel electrode, the applied voltage was changed to +0.8 V during the interval.

The galvanometer spot was set to zero and the sensitivity control was set to an appropriate level. Up to 0.1 mL of the equivalence point's distance, the titrant thiomalic acid solution was added from the burette in 0.5 mL increments, and then for the remaining 2 mL in 0.5 mL increments. The gas stream was circulated through the solution for about a minute (more diluted solutions may take up to three minutes), then it was passed over the surface, and finally the galvanometer deflection was read to help with the precipitation of the copper ions after each addition (current). The volume of the solution has been changed in the current readings to account for the addition of the reagent. An amperometric titration curve was constructed using the obtained result, and the equivalency point was then determined by reading off the curve (Gaballah et al., 2019; Salem, 1996; Nivetha et al., 2016).

8.2.7 Chlorine Determination

A quick and incredibly accurate method for measuring free chlorine has been developed using differential pulse voltammetry and an Au working electrode. As per the literature, this standardized protocol is highly promising for monitoring the level of remaining free chlorine in drinking water. The electrochemical experiments were carried out in a separate compartment cell. Pt wire, an Ag/AgCl (saturated KCl) disc, and a 1.6 mm diameter Au disc were used as the working, reference, and counter electrodes, respectively. The Au working electrode's surface was pre-treated by being polished with an aluminium oxide suspension, followed by a meticulous sonication in water and a rinsing with Milli-Q water. A solution of 0.1 mol/dm³ potassium chloride served as the supporting electrolyte. The DPV measurements were performed using the BD-101 electrochemical analyzer with a C-1B cell stand. For the DPV measurements, the pulse amplitude was set at 25 mV, and the scan rate was 20 mV/s. Potassium chloride was added to a 20 cm³ water sample until the final concentration was 0.01 mol dm⁻³. For all measurements, the thermostat was set at a temperature of around 25 °C (Ahmed 2011; Mangione and Carlyle, 2012).

8.3 Results and Discussion

For samples derived from home wastewater, levels of BOD, COD, turbidity, pH, chloride, phosphate, and nitrate were monitored. These factors were assessed over the course of 21 days, and the findings were tallied. To evaluate the effectiveness of the water mint for treating wastewater, the acquired findings were compared. Three different sets of values are noted. Before adding the water mint plant to the samples, the initial set of values for the aforementioned parameters was measured on day 0. Using the tests conducted on Day 11, the second set of data was determined. All the previously indicated parameters have clearly dropped. Additionally, a greater

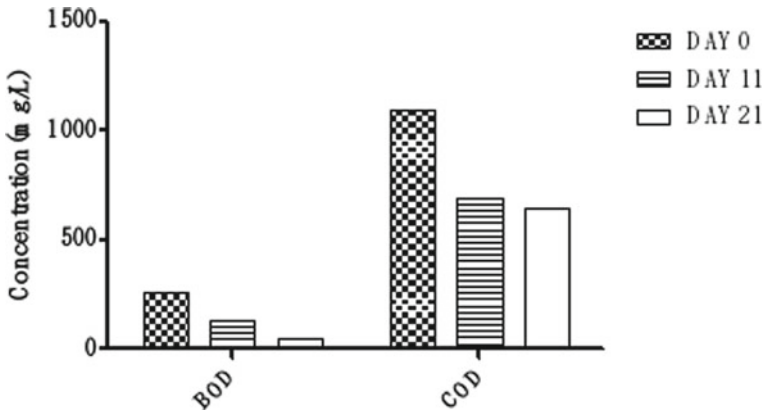


Fig. 8.1 BOD and COD values of domestic wastewater treated with *Mentha aquatica*

reduction was noted on the 21st day. The results and observations are discussed in this section.

8.3.1 BOD and COD

From day 0 to day 21, there was a gradual decrease in the capacity of water mint to remove BOD and COD. The BOD range initially is 263 mg/L, lowers to 136 mg/L on day 11, and finally drops to 49 mg/L on day 21 because of the plant’s great capacity to absorb toxins from the residential wastewater. On day 0, the COD levels were roughly 1100, 697 mg/L on day 11, and 647 mg/L on day 21. The effectiveness of BOD removal on the eleventh day was 48.288%, exceeding the 44.4% figure of BOD removal recorded for Lotus. (Jyoti 2017). The BOD elimination efficiency was almost 64% after 21 days, which was somewhat closer to the value of 66.6% recorded for Lotus. Similar to this, after 11 days, COD elimination efficiency was 36.64%, and after 21 days, it was 7%. After 20 days, Lotus’ COD elimination efficiency was reported to be 32%. This leads us to the conclusion that the first 11 days are when the COD for water mint is actively reduced (Fig. 8.1).

8.3.2 Turbidity

The domestic wastewater was turbid on day 0, which indicates that heavy metal pollution has caused the water to be significantly contaminated and seem to be a greyish black in colour. On day 0, a turbidity measurement of 319 NTU was taken. The turbidity gradually dropped to 36.8 NTU by the observation period’s eleventh day, and by the twenty-first day it had totally vanished to 4.7 NTU. On days 11 and

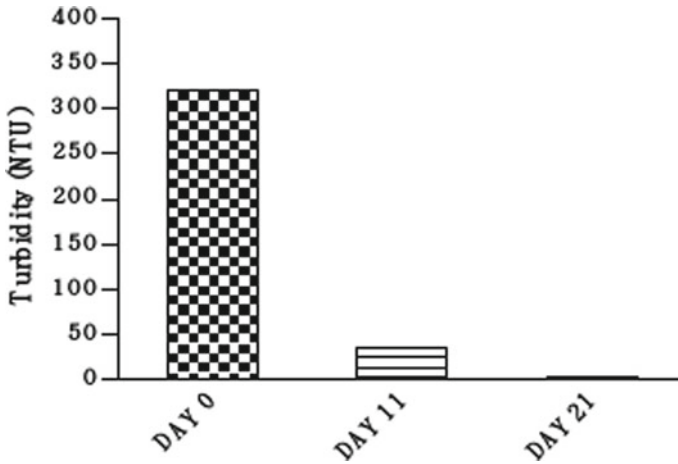


Fig. 8.2 Turbidity values of domestic wastewater treated with *Mentha aquatica*

21, respectively, the efficiencies of turbidity reduction were 88.46 and 87.23%. This was consistent with Lotus claimed TDS removal efficiencies, which were 87% and 86% after 10 days and 20 days, respectively (Jyoti 2017) (Fig. 8.2).

8.3.3 pH Value

From day 0 to day 21, the pH efficiency is shown in Fig. 8.3 to have gradually decreased. Due to the plant's exceptional ability to absorb contaminants from household wastewater, the pH range initially ranged from 8.34 to 7.66 on day 11 and 7.31 on day 21. At 25 °C, the pH of the home wastewater was determined. This consistent pH drop is consistent with the pH changes observed in water caltrop, which was employed to treat sewage from local governments (Kumar and Chopra 2018).

8.3.4 Removal Efficacy of Chlorides, Phosphorus and Nitrates

The chloride as Cl⁻ starting value was around 1250 mg/L on day 0, and it swung to a range of 1799 mg/L on day 11, before decreasing to 1550 mg/L on day 21. The value changed because of the substance's water solubility and lack of biodegradability. The pollutants steadily decreased as the amount of chloride was in the wastewater (Fig. 8.4). Water mint was found to have 13.84% efficacy in removing chloride after 21 days, but Lotus, Water Caltrop, and Hydrilla have not yet been reported on.

Fig. 8.3 pH value of domestic wastewater treated with *Mentha aquatica*

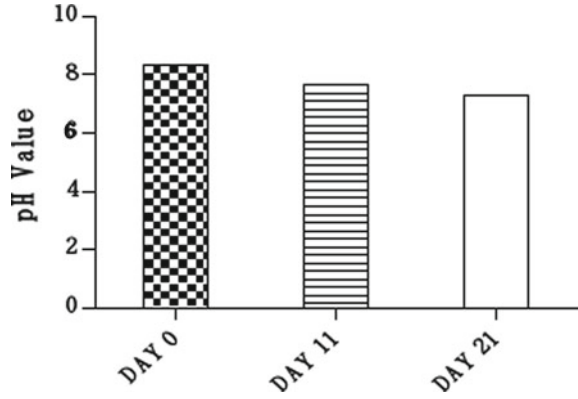
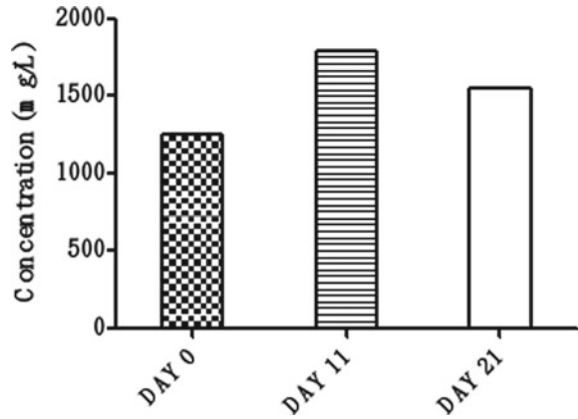


Fig. 8.4 Chloride concentration of domestic wastewater treated with *Mentha aquatica*



The overall phosphorous concentration of the detergents affects pH. On day 1, the total phosphorous level was roughly 0.32 mg/L, and by day 11, it varies between 0.17 and 0.17 mg/L. This serves as an example of the full phosphorous removal process. Phosphorus falls to 0.13 mg/L on the last day of observation. At the end of 11 days, the total phosphorous removal efficiency was 46.88%, and 23.53% at the end of 21 days. Lotus and Hydrilla had total phosphorous removal efficiencies of 66% and 65%, respectively, after 10.5 days (Kanabkaew and Puetpaiboon, 2004) (Fig. 8.5).

On the eleventh day of observation, Fig. 8.6 depicts the fluctuation in nitrate removal from home wastewater. On day eleven of observation, the value rose to 31 mg/L from an initial value of about 29 mg/L. The result had dropped to 27 mg/L on the twenty-first day of monitoring, notwithstanding a fluctuation on the eleventh day, which was seen. This variation in the nitrate concentration can be ascribed to the water mint’s ability to remove ammonia from home wastewater, which necessitates the conversion of NH_3 to NO_3^- by active nitrification. Due to the de-nitrification process, we later see a decrease in the concentration of nitrate. Similar results were reported by Selvarani et al. (2015) for Duckweed.

Fig. 8.5 Total phosphate concentration of domestic wastewater treated with *Mentha aquatica*

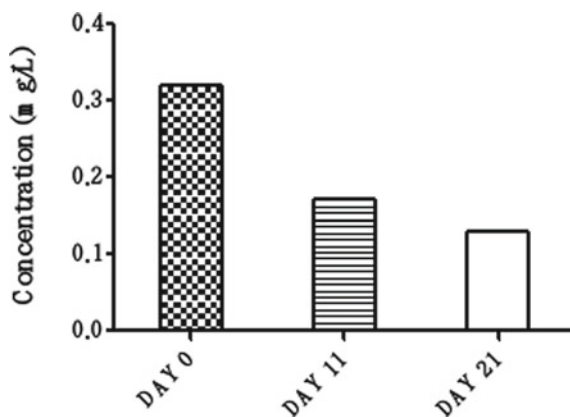


Fig. 8.6 Nitrate concentration of domestic wastewater treated with *Mentha aquatica*

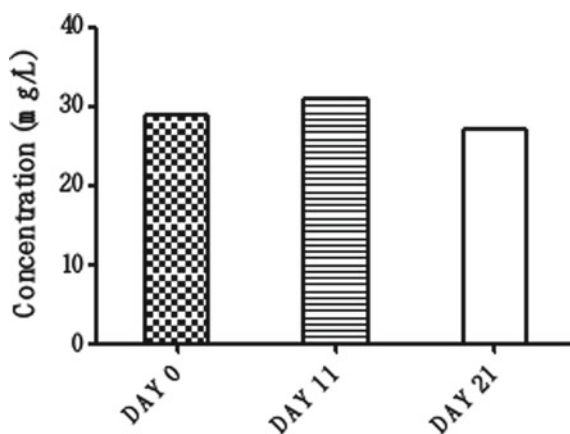


Table 8.1 clearly shows that all parameters have significantly decreased after receiving treatment with *Mentha aquatica* for 21 days.

Table 8.1 Cumulative result of parameters of *Mentha aquatica* treated domestic wastewater

Parameters	Units	Day 0	Day 11	Day 21
pH value	–	8.34	7.66	7.31
Turbidity	NTU	319	36.8	4.7
Chlorides as Cl ⁻	mg/L	1250	1799	1550
BOD	mg/L	263	136	49
COD	mg/L	1100	697	647
Total phosphorus	mg/L	0.32	0.17	0.13
Nitrate as NO ₃ ⁻	mg/L	29	31	27

8.4 Conclusion

The study thus reveals the advantage of using water mint to clean domestic wastewater. The findings showed a sharp reduction in effectiveness from day 0 to day 21 for the criteria of BOD, COD, turbidity, and phosphorus. The BOD and COD levels fell from 263 to 49 and 1100 to 647 mg/L, respectively. Turbidity was reduced by 4.7 NTU, from 319 NTU. The pH readings ranged between 8.34 and 7.31. The effectiveness of the other criteria might also be impacted by environmental variables like nitrate and chloride. The value of the chlorides ranged from 1250 to 1550 mg/L, and the value of the nitrate dropped from 29 to 27 mg/L. The analyses cited above show that water mint is quite good at filtering out pollutants from home wastewater. Using water mint to filter water is convenient and eco-friendly.

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Chapter 9

Screening of Bacterial Isolates from Coal Mining Region in Chhattisgarh



Rajni Kumari, K. Harshan, Prashanth Rajan, Anand Prem Rajan, and Thomas Theodore

Abstract Microorganisms that can thrive under intense conditions have generated considerable interest in recent days on account of their exceptional physiology, nature, and essential industrial applications. This could be because of the special sequences that they possess in their DNA which can produce proteins with unrivalled biocatalytic properties. The present study aims at determining the microbial diversity of mine soil collected from Indian Coal Mines, South Eastern Coal Field Ltd (SECL), Korba district, Chhattisgarh. The soil samples were evaluated based on their color, texture and structure. Bacteria were isolated using iron salts-purified agar (ISP) media by serial dilution method and spread-plate method. The bacterial isolates were of yellow and brown colours. Microscopic observation revealed that the isolates were Gram-negative, rod-shaped bacteria. Bacteria were subjected to glucose test, lactose test, fructose test, triple sugar iron agar test, mannitol motility test, catalase test, oxidase test, citrate test, hydrogen sulphide test, phenylalanine deaminase test, indole test, methyl red test, and Voges–Proskauer test. Molecular assay of 16S rRNA sequencing revealed the isolate to be *Enterobacter cloacae*.

Keywords Microbial diversity · Coal-mine bacteria · *Enterobacter cloacae*

Misc Extremophilic *Enterobacter cloacae* screened and reported for the first time in the coal mines of Chhattisgarh

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9.1 Introduction

Potentially poisonous elements (PTEs) from diverse anthropogenic sources have found a home in soil. Since the beginning of time, it has served as a reservoir for a sizable number of PTEs due to the unlawful and negligent dumping of household, agricultural, mining, and industrial (Palansooriya et al. 2020; Zhou and Wang 2019). Coal harbors bacteria of multiple genera. Bacteria are found ubiquitously in the environment: at great heights, in deepest sea sediments, at extremes of heat and cold, and in highly contaminated sites. These bacteria found in coal mines are extremely rugged in nature, hence may prove to have numerous industrial applications. They also perform coal bio-beneficiation (Mukherjee et al. 2018) along with a role in many other biogeochemical processes. Coal mine soil has numerous strains of *Methanogen* sp. and *Thiobacillus* sp. (Brock 1978). The first bacterium *Thiobacillus ferrooxidans* was isolated from acidic bioleaching environments, (Colmer and Hinkle 1947), (Donati and Sand 2007). It is difficult to describe the exact role and significance of each organism in these active populations, as the intimate connection between sulphide biohydrometallurgy and microbial physiology is not completely understood (Belly and Brock 1974).

Overburden of coal mine spoil signifies disturbed habitat for the soil organisms due to its low pH and high internal temperature (Juwarkar et al. 2014). The quantity of Gram-negative bacteria from coal mine spoils of different geographical regions has also been reported (Marsh and Norris 1983). It was discovered that the Gram-negative bacteria were resistant to acidic environments as well as Fe, Cd, and Cr. They were recovered from acid mine drainage (AMD) of rat-hole coal mines (Ka-ot and Joshi 2022).

Heavy metal resistant bacteria were reported in Bokaro coalfield, Jharkhand soil samples as *Acinetobacter gyllenbergii* (KM029963; NK-7), *Enterobacter ludwigii* (KM029957; NK-1), *Enterobacter ludwigii* (KM029959; NK-3), *Enterobacter ludwigii* (KM029960; NK-4), *Enterobacter cloacae* (KM029962; NK-6), *Enterobacter cloacae* (KM029964; NK-8), *Klebsiella pneumonia* (KM029958; NK-2), *Klebsiella oxytoca* (KM029961; NK-5) (Gandhi et al. 2015).

In Singareni Collieries, Andhra Pradesh BDRC1 (*Bacillus* sp.) (KJ643909.1), BDRC2 (*Bacillus* sp.) (KJ643910.1), and BDRC3 *Bacillus* sp. (KJ643911.1) were the bacterial species that dominated the soil (Machulla et al. 2005). The bacteria found in mining areas are good for bioremediation of aquatic areas by accumulating metals in their cells via a process known as biosorption (Mishra and Ma 2005).

The largest area of acid sulfate soil materials in Finland which caused large-scale fish mortality in 2006 due to acid run-off. *Acidithiobacillus ferrooxidans* and *Acidithiobacillus thiooxidans* were isolated from acid sulfate soil materials (Arkesteyn 1980).

“Potential acid sulfate soil materials” are metal sulphide, containing anoxic residues found various coastal areas of Australia and the Baltic (Rickard and Luther 2007).

Bacterial molecular studies are done by molecular sequencing. Isolates from Australian potential acid sulphate soil (PASS) and acid sulphate soil (ASS) had 16S rRNA gene sequences that showed similarity to the iron-oxidizing bacteria *Alicyclobacillus* species and *Thiomonas* species. The isolates from both PASS and ASS plates during molecular phylogenetic studies resulted in 40 operational taxonomic units (OTUs) (Ohba and Owa 2005).

Acidithiobacillus thiooxidans was first discovered in 1947, in an acid mine drainage in Japan. Nevertheless, experience from low-temperature sulphide mineral environments and acid mine drainage (AMD) has shown a significantly simpler mixed population in soil. A single acidophile capable of low-temperature growth on Fe^{2+} and/or inorganic sulphur compounds (ISCs) has been reported (Baker-Austin et al. 2007). The common bacteria *Acidithiobacillus ferrooxidans* (*A. ferrooxidans*) and *Acidiphilium multivorum* (*A. multivorum*), which mediate Fe^{2+} oxidation and Fe^{3+} reduction under an acidic environment, respectively, are found throughout acid mine drainage (Jin et al. 2020; Wang et al. 2020). The molecular evolution of boreal PASS and ASS isolates were reported from the Risofladan experimental field, Finland (Kupka et al. 2009).

Coal sources in India are mostly dispersed along different valleys such as Pench-Kanhan valley, Damodar valley, Wardha-Godavari valley, Sone-Mahanadi valley, etc. The greater part of the coal stores is situated in the south-eastern region of India bound between 24° North latitude and 78° East longitude in West Bengal, Chhattisgarh, Southern Uttar Pradesh, Eastern Madhya Pradesh, Odisha, and Jharkhand. Altogether, there are 57 coal mines present in these areas. The entire Gondwana coal-fields range up to $64,000^\circ\text{km}^2$. Overall, 194 billion tons of coal reserves are predicted including non-coking coal reserves of approximately 165 billion tons (Anand and Prasad 2013).

Coal contains four different forms of sulphur, namely, organic sulphur-containing compounds of calcium and barium, pyritic sulphur (FeS_2), elemental sulphur, and sulphates of iron. Among these, organic and pyritic sulphur is the most commonly known. Acidic mine water is produced by bacteria *Thiobacillus thiooxidans* and *Thiobacillus ferrooxidans* that are present in coal deposits (Natarajan 1978). Chemoautotrophs such as *Thiobacillus ferrooxidans* oxidizes ferrous to ferric iron and forms sulphuric acid resulting in acid mine drainage (AMD):



Microbial action is also affected by the coal types (lignite, bituminous, sub-bituminous, and anthracite) along with the matrix and contents of the coal matrix (Natarajan 1978).

Different mesophilic organisms, thermophiles and fungi were found to exist in coal-mining waste dumps (Tichý et al. 1998). These species have been found to play an active role in microbial desulphurization. Various strains of heterotrophs such as *Acinetobacter*, *Beijerinckia* sp., *Pseudomonas* sp., *Rhizobium* sp. execute organic sulphur oxidation.

Our in-depth literature survey has brought to light that the bacteria of Chhattisgarh coalfields are still unexplored. No industrially useful bacteria have ever been reported so far from these virgin soils, which prompted us to design and carry out this present research work.

9.2 Materials and Methods

9.2.1 Study Area

South Eastern Coalfields Limited (SECL), Korba, Chhattisgarh is a major coal-producing company of India. The company has its head office at Bilaspur. It has a total of 92 mining sites. There are 70 underground mines and 22 open-cast mines.

9.2.2 Sample Location Area

See Fig. 9.1.

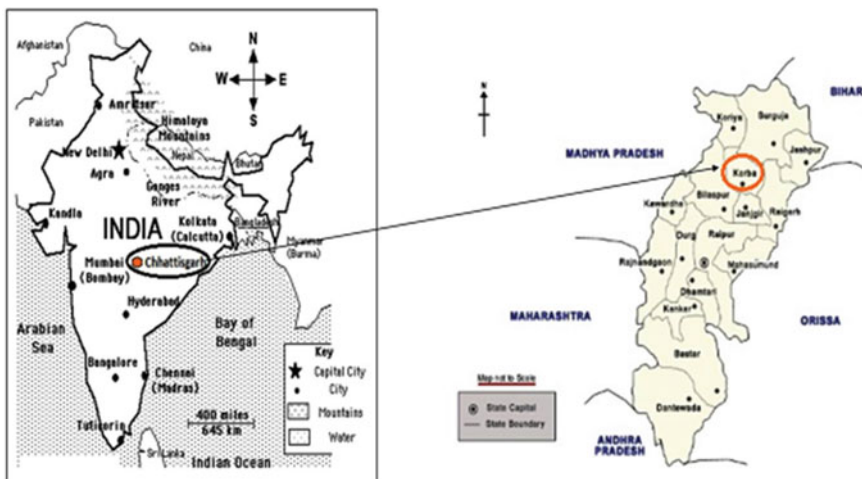


Fig. 9.1 Map of India and Korba district in Chhattisgarh, the location of South Eastern Coalfields Limited

Fig. 9.2 Sample collection area from a coal mine of South Eastern Coal Field Limited (SECL), Korba District, Chhattisgarh



9.2.3 Sampling Site

The mining soil samples were collected at 50 feet depth (22.35° N; 82.68° E) from South Eastern Coal Field Limited (SECL), Korba District, Chhattisgarh. The soil samples were collected in sterile ziplock bags and brought to our laboratory. The samples were examined for their physicochemical characteristics such as colour, texture, density, porosity, and pH (Fig. 9.2).

9.2.4 Growth Media and Culture Conditions

9.2.4.1 Iron Salt Purified Media (ISP)

Bacterial isolation from soil samples was carried out using Iron salts-purified agar (ISP) media.

ISP media was prepared using three sterilized solutions:

Solution A: Ferrous sulphate ($\text{FeSO}_4 \cdot 7\text{H}_2\text{O}$ —33.4 g/L) was added to 300 mL of distilled water and the pH was adjusted to 2.5 using 6 M sulphuric acid (H_2SO_4). The solution was filter sterilized.

Solution B: Ammonium sulphate ($\text{NH}_4)_2\text{SO}_4$ —6.0 g/L), potassium chloride (KCl—0.2 g/L), magnesium sulphate ($\text{MgSO}_4 \cdot 7\text{H}_2\text{O}$ —1.0 g/L), calcium nitrate ($\text{Ca}(\text{NO}_3)_2$ —0.02 g/L) were dissolved in 550 mL double-distilled water. The pH of the solution was adjusted to 3.0 and autoclaved.

Solution C: Purified agar was soaked in 150 mL of double-distilled water and autoclaved. Solution B was mixed with solution C with gentle mixing and then solution A was added to this mixture. The mixture was then poured into Petri plates and allowed to solidify.

9.2.4.2 Colony Forming Units (CFU)

5.0 g of soil samples were taken in Erlenmeyer flasks containing 50 mL of deionized double-distilled water. The plates were incubated at 28 °C in the dark for 48 h. The colony forming units (CFU) was calculated using the formula:

$$\text{CFU} = \frac{\text{Number of colonies}}{\text{Volume of solution taken}} \times \text{Dilution factor}$$

9.2.4.3 Bacterial Isolation

Soil samples mixed in ISP media were shaken rigorously and then subjected to serial dilution followed by spread plating. In order to isolate pure culture, isolated bacteria were grown in ISP media (Manning 1975). Cultures were streaked directly on ISP media to obtain isolated colonies. Individual colonies were cultured in broth at 30°C in Erlenmeyer flasks on a rotary shaker at 150 rpm.

9.2.4.4 Microscopic Examination

Gram staining was done and observed under 10× and 100× magnification.

9.2.4.5 Biochemical Characteristics of the Bacteria

The biochemical characteristics of the bacteria were studied. The extra-cellular enzymatic activities were performed by gelatin hydrolysis and starch hydrolysis. The intracellular enzyme activities were determined by the urease test, catalase reaction, indole production test, methyl red test, Voges-Proskauer test, citrate utilization test, and triple sugar iron test. Different carbon sources were used to study the effect of carbon source on the growth of bacteria.

9.2.4.6 Primers

From the purified genomic DNA, the 16S rRNA gene was amplified using polymerase chain reaction (PCR) with universal primers 27F [5'—AGA GTT TGA TCC TGG CTC AG—3'] and 1492R [5'—ACG GCT ACC TTG TTA CGA CTT—3'] (Mageswari et al. 2012).

9.2.4.7 16S rRNA Sequencing

The PCR reaction mixture of a total volume of 20.0 μL , containing 6.0 μL of nuclease-free water, 1.0 μL of forward primer, 1.0 μL of reverse primer, 2.0 μL of DNA, 10.0 μL of PCR master mix was prepared in a 200 μL PCR tube. The amplification was carried out in a thermal cycler and consisted of 30 cycles. It was performed under the following conditions: initial denaturation at 94 $^{\circ}\text{C}$ for 1 min, denaturation at 94 $^{\circ}\text{C}$ for 1 min, annealing at 49 $^{\circ}\text{C}$ for 1 min, extension at 72 $^{\circ}\text{C}$ for 1 min, final extension at 72 $^{\circ}\text{C}$ for 10 min. The PCR product (10.0 μL) was analyzed using 1.5% agarose gel stained with 5% 10.0 mg/mL ethidium bromide.

9.2.4.8 Phylogenetic Analysis

The analysis of the evolutionary pattern of genes was achieved by the construction of the phylogenetic tree. MEGA v5.2.2 (Molecular Evolutionary Genetics Analysis version 5.2.2) software was used for the construction of the phylogenetic tree. This software has the advantages of collection of databases, alignment of large sequences, and phylogenetic tree construction. This information can be used for evolutionary analysis in basic biology, biomedicine, and other evolution studies. The alignment of DNA sequences was used to perform the trace file editor option. MEGA 5.2.2 has derived models for the evaluation of nucleotide and amino acid substitutions. The trees are constructed using neighbor joining, maximum likelihood, maximum parsimony with bootstrap, and branch length. It computes distances by pair-wise method within and between groups. The maximum likelihood method is used for referring to evolutionary trees to select the best-fit substitution models for amino acid as well as nucleotide sequences to represent the evolutionary rate (Tamura et al. 2011).

9.3 Results and Discussion

The soil samples were brown to yellow in color with a pH of 7.2. The normal temperature during collection was 37 $^{\circ}\text{C}$.

9.3.1 Colony Forming Units (CFU)

Very few colonies were obtained in the test samples. CFU was found to be 3.0×10^{-4} CFU/mL for 48 h.

Table 9.1 Colony morphology of the isolated bacteria from the coal mines of South Eastern Coal Fields Limited (SECL), Korba District, Chhattisgarh

	Bacteria	Colony morphology characterization		Selective media	Bacterial shape	Elevation	Size (mm)
		Colour	Shape				
1	Isolate 1	Yellow	Circular with irregular margin	ISP media	Rod	Convex	1
2	Isolate 2	Grey	Circular with irregular margin	ISP media	Rod	Flat	1
3	Isolate 3	White	Circular with regular margin	ISP media	Coccus	Pulvinate	2

9.3.2 Bacterial Isolation

Three distinct bacterial strains were obtained from the test samples. The colony morphology characterization of the isolates is provided in Table 9.1. From the obtained isolates, the pale-yellow colony forming bacteria was found to be the most dominant. Hence, pale-yellow colonies were chosen for identification and characterization due to their abundance in initial plating (Jamal et al. 2016).

9.3.3 Colony Morphology

Colony morphology is a significant feature for the isolated bacteria. Margins of the colony were irregular but circular with slight elevation for isolate 1. The pale yellow pigmentation may be due to the oxidation of ferrous ions in the ISP media.

9.3.4 Microscopic Examination

The isolates were found to be Gram-negative rods around 2.6 μm long and 1.0 μm wide. The cells were motile and possessed flagella.

Table 9.2 Biochemical characteristics of the isolated bacteria from coal mine soil

Biochemical test	Result
Glucose test	Positive
Lactose test	Negative
Triple sugar iron agar	Positive
Mannitol motility test	Positive
Catalase test	Positive
Oxidase test	Negative
Citrate test	Positive
Hydrogen sulphide test	Negative
Phenylalanine deaminase test	Negative
Indole test	Negative
Methyl red test	Negative
Voges-Proskauer test	Positive

9.3.5 Biochemical Assay

The biochemical characterization was performed for Isolate-1 as a vital part in the characterization of the isolated bacteria. Table 9.2 consolidates the result of reactions of isolate-1 during various biochemical tests.

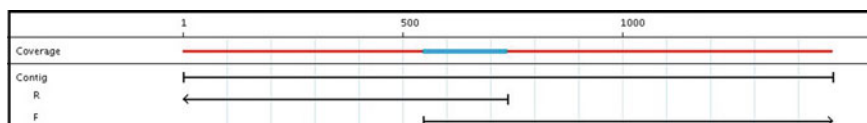
The results of biochemical tests show that isolate-1 may be from the *Enterobacteriaceae* family.

9.3.6 16S rRNA Bacterial Sequencing

Molecular analysis was done to confirm the bacterial species. The assembled sequence is given in Fig. 9.3.

The obtained sequence was found to be:

```
AAACAGCTCAGATTGAACGCTGGCGGCAGGCCTAACACATGCAAGTCGAACGGTA
ACAGGAAGCAGCTTGCTGCTTCGCTGACGAGTGGCGGACGGGTGAGTAATGTCTG
GGAAACTGCCTGATGGAGGGGGATAACTACTGGAAACGGTAGCTAATACCGCATAA
CGTCGCAAGACCAAAGAGGGGGACCTTCGGGCCTCTTGCCATCGGATGTGCCCAG
ATGGGATTAGCTAGTAGGTGGGGTAACGGCTCACCTAGGCGACGATCCCTAGCTGG
```

**Fig. 9.3** The assembled sequence for isolate-1

TCTGAGAGGATGACCAGCCACACTGGAAGTGGAGACACGGTCCAGACTCCTACGGG
 AGGCAGCAGTGGGGAATATTGCACAATGGGCGCAAGCCTGATGCAGCCATGCCGC
 GTGTATGAAGAAGGCCTTCGGGTTGTAAAGTACTTTCAGCGGGGAGGAAGGTGTT
 GAGGTTAATAACCTCAGCAATTGACGTTACCCGCAAGAAGAAGCACCGGTAACCTC
 GTGCCAGCAGCCGCGGTAATACGGAGGGTGCAAGCGTTAATCGGAATTACTGGGC
 GTAAAGCGCACGCAGGCGGTCTGTCAAGTCCGATGTGAAATCCCCGGGCTCAACC
 TGGGAAGTGCATTTCGAAACTGGCAGGCTAGAGTCTTTGTAGAGGGGGGTAGAATTC
 CAGGTGTAGCCGTGAAATGCGTAGAGATCTGGAGGAATACCGGTGGCGAAGGGCG
 CCCCCTGGACAAAGACTGACGCTCAGGTGCGAAAGCGTGGGGAGCAAACAGGAT
 TAGATACCCTGGTAGTCCACGCCGTA AACGATGTGCACTTGGAGGTTGTGCCCTTG
 AGGCGTGGCTTCCGGAGCTAACGCGTTAAGTCGACCGCCTGGGGAGTACGGCCGC
 AAGGTTAAAACCTCAAATGAATTGACGGGGGCCCGACAAGCGGTGGAGCATGTGG
 TTTAATTCGATGCAACGCGAAGAACCTTACCTACTCTTGACATCCAGAGAACTTAGC
 AGAGATGCTTTGGTGCCTTCGGAACTCTGAGACAGGTGCTGCATGGCTGTCGTCA
 GCTCGTGTGTGAAATGTTGGGTTAAGTCCCGCAACGAGCGCAACCCCTTATCCTTT
 GTTGCCAGCGGTCCGGCCGGGAACTCAAAGGAGACTGCCAGTGATAAACTGGAGG
 AAGGTGGGGATGACGTCAAGTCATCATGGCCCTTACGAGTAGGGCTACACACGTGC
 TACAATGGCGCATACAAAGAGAAGCGACCTCGCGAGAGCAAGCGGACCTCATAAA
 GTGCGTTCGTAGTCCGGATTGGAGTCTGCAACTCGACTCCATGAAGTCCGAATCGCT
 AGTAATCGTGGATCAGAATGCCACGGTGAATACGTTCCCGGGCCTTGACACACCG
 CCCGTCACACCATGGGGAGTGGGTTGCAAAAAGAAGTAGGTAGCTTAAACCTTCGGG
 AGGGCGCTTACCACCTTTGTGATTCATGACTGGGGTGAGC

The analysis of the sequence fragments was done by using Blast algorithm at NCBI (www.ncbi.nlm.nih.gov/nuccore/KC990822.1). The complete 16S rRNA gene sequence of the isolated *Enterobacter cloacae* strain DRA1 genomic sequence has been deposited under the accession number KJ947344.

9.3.7 Phylogenetic Analysis

The phylogenetic analysis result of *Enterobacter cloacae* genes is presented in Fig. 9.4. The phylogenetic tree of 4 bacterial genes showed an independent evolution of *Enterobacter cloacae* compared to *Citrobacter murliniae*. *Citrobacter murliniae* is resistant to metal ions (Boechat et al. 2017). In the bootstrap test, the percentage of replicating trees for 4 bacterial sequences is represented next to the branches. The genes of *Enterobacter* showed homology with that of *Citrobacter*, *Leclercia*, and *Salmonella* bacteria. The results show that the evolution of Isolate-1, i.e., *Enterobacter cloacae* including these 3 bacteria originated from the same ancestor as they are present in one clad (Fig. 9.5).

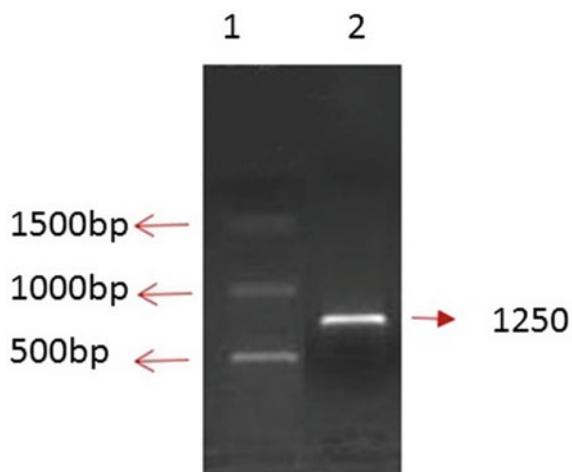


Fig. 9.4 16S rRNA sequencing of the isolated bacterial colony (expected size 1250 bp) Lane 1. 1 kb DNA ladder, Lane 2. Amplified DNA product

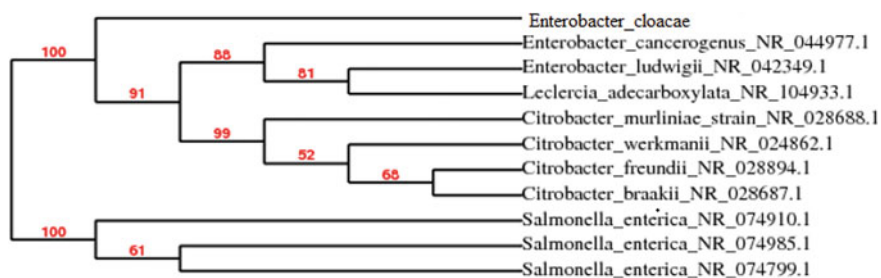
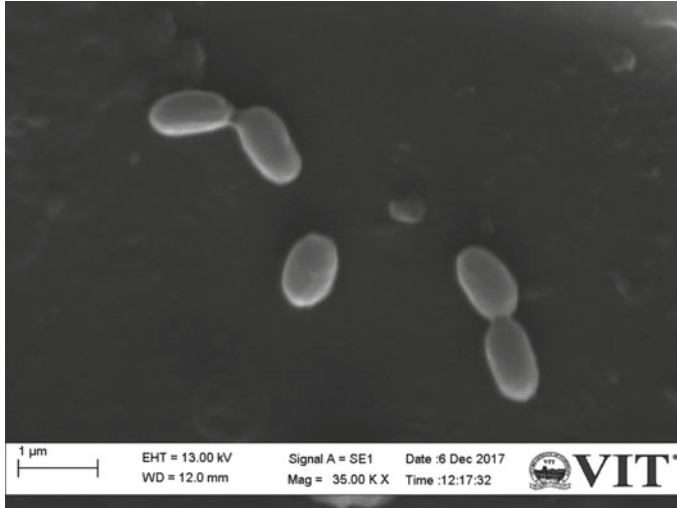


Fig. 9.5 The phylogenetic tree for *Enterobacter cloacae* gene in coal mine bacteria

The optimal temperature of culture for Isolate-1, i.e., *E. cloacae* was observed to be 30–37 °C and the optimal pH was observed over a wide range (pH 5–9). The bacteria could suitably be utilized for bioremediation at optimized temperature and pH conditions (Panda and Sarkar 2012).

Enterobacter strains grown in ISP medium has been reported to show faint growth, which demonstrated that they might not be strict chemolithotrophs, but rather they could still use this media even without any hydrocarbon source. This stress-tolerance ability may be credited to their abundance in the coal mine (Arnold et al. 2004).

9.3.8 Scanning Electron Microscopy (SEM)



The Scanning electron micrograph shows that the isolate-1, namely, *Enterobacter cloacae* was 1 μm in length.

9.4 Conclusion

In conclusion, *Enterobacter cloacae* was found to be the most abundant bacteria out of the three isolates obtained from the mining soils of South Eastern Coal Field Limited, Korba District, Chhattisgarh. This bacterium was isolated and grown on ISP media which exhibited pale yellow colonies. Biochemical studies and molecular studies confirmed it to be *Enterobacter cloacae*. As this strain is isolated from a harsh mining environment, it mimics the chemolithotrophs. It has a vast industrial potential for bioremediation, e-waste treatment, metal leaching, and bioleaching applications. Further studies need to be carried out on the growth kinetics of the isolate and its industrial applications.

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Conflict of Interest On behalf of all the authors, the corresponding author states that there is no conflict of interest.

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Chapter 10

Analysis of *Hedychium Flavum* Waste Powder as a Potent Heavy Metal Adsorbent



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Abstract *Hedychium flavum*, a medicinal, edible, and ornamental plant, is widely cultivated in India. Essential oils from the plant are used for various medical purposes and the leftover plant residue after oil extraction is discarded as a waste. An attempt was made to explore the use of this waste to remove heavy metals from wastewater was analysed. Result shows that the waste powder can be used effectively for metal removal. *Hedychium flavum* sequence was submitted to the NCBI database to procure accession numbers MN988619.

Keywords *Hydicum flavum* · RSM · Optimization software · Metal removal · Modelling

10.1 Introduction

Agricultural residues are a renewable resource that can help alleviate our planet's pressing environmental issues sustainably and economically. Numerous studies have shown that this type of agricultural waste, both cheap and abundant, has great promise as an adsorbent for removing various contaminants from waste streams. Hazardous metal contamination in water has long persisted as a severe concern to human health and aquatic life due to the persistence of heavy metals in nature. Heavy metals are released into water sources typically caused by leakage of chemical pollution, mineral processing, and many other forms of manufacturing activity. Although trace amounts of various transition metals, e.g., copper, nickel, and lead, are required for live organisms, their excessive levels may induce detrimental effects, including neurological diseases, respiratory failure, birth deformities, or even death in extreme

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cases. Conventional treatment procedures, such as oxidation/reduction, coagulation/flocculation, reverse osmosis/membrane filtering, ion exchange, and electrochemical approaches, are either too expensive or too inefficient to be widely used for the decontamination of heavy metals.

However, the adsorption process using activated carbon as an adsorbent is particularly favored because of its high efficiency, simple operation, immunity to harmful chemicals, reusability, and scope of use. Adsorption process efficiency is very sensitive to adsorbents' properties and the choice of optimal operating conditions. The characteristics of the input materials and the method of manufacture often govern the former. In contrast, the latter demands selecting an optimal combination of several influencing parameters, typically established by extensive experimentation. However, the number of experiments required can be drastically cut down by using optimization techniques. One of the most important multivariate techniques used in analytical optimization is response surface methodology (RSM), which is a collection of mathematical and statistical methods based on fitting empirical models to the experimental data collected from a set of predefined trials. To determine which experiments need to be conducted in the investigated experimental region to assess the independent variable effects and their interactions with the fewest possible trials, BBD is the most frequently used matrix design among those with a wide range of potential properties and characteristics.

The *Hedychium* plant genus is ornamental and ethnomedicinal, encompassing around 90 species, disseminated across South-East Asia. *Hedychium* species are a potent origin of numerous components with medical and allied industrial applications. After extraction of oil and other components the wastes of the plant parts are discarded. The current work focused on the use of *Hedychium* waste on the metal adsorption capabilities (Raj et al. 2013; Thomas et al. 2015).

10.2 Materials and Methods

10.2.1 Materials

Processed *Hedychium* waste was collected from a local industry. The waste was washed with distilled water and sun dried.

10.2.2 Adsorption Experiments

Adsorption experiments were carried out in the flasks containing 50 mL of aqueous solution of chromium metal ion. The heavy metals levels were confirmed by AAS analysis. The removal efficiency and the adsorption capacity were calculated as follows:

Table 10.1 Design parameters

Factor	Name	Units	Minimum	Maximum
A	pH		1	5
B	Adsorbent dosage	g/L	0.100	1.0
C	Cr concentration	mg/L	50	500

$$\text{Chromium removal (\%)} = [(C_i - C_f)/C_i] * 100 \quad (10.1)$$

where C_i and C_f are the initial and equilibrium M (II) concentrations (ppm) respectively.

10.2.3 RSM-BBD Experimental Design

In order to obtain responses across the entire factor space and determine the region of optimal or near-optimal response via a sequence of designed experiments, response surface methodology is widely applied due to its potency as a powerful tool that combines the benefits of mathematical and statistical techniques. To identify the best conditions for the adsorption process, the current research used response surface methodology (RSM) along with Box Behnken design (BBD). The adsorption capacity of the adsorbent was used as the output variable, with the concentration of metal ions (Chromium) (A), the dosage of the adsorbent (B), and the pH (C) serving as the input factors (Table 10.1).

Analysis of variance (ANOVA) of the quadratic polynomial regression models was utilized to identify the significance of input variables as well as the relationship between the responses and the influential factors. In this study, the ANOVA was calculated using Design-Expert version 11.

10.3 Results and Discussions

Experimental design was created to explore the contribution of three influential factors: pH, metal ion concentration and adsorbant concentration (Factor a, b, c) on the sorption capabilities of *Hedychium flavum* waste powder on chromium metal (Response R) [Table 10.2]. The response metal sorption % was calculated using Eq. 10.1.

To prove the model fitness, the analysis of variance (ANOVA) was done (Table 10.3). The R (%) response was statistically significant with F-value of 56.61 and p-value of 0.0001. The results elucidated that adsorbant dosage (*Hedychium flavum* waste powder) considerably impacts the R (%) response followed by AC: pH with the combination of metal concentration). There was good congruity of the outcome

Table 10.2 Experimental runs and metal sorption response

	Factor A	Factor B	Factor C	Response R
Run	A: pH	B: Adsorbent dosage g/L	C: Cr concentration mg/L	%removal
1	5	0.55	50	80
2	5	0.1	275	41
3	1	0.55	50	62
4	3	0.55	275	72
5	3	1	50	88
6	5	1	275	90
7	5	0.55	500	69
8	3	0.55	275	73
9	3	0.55	275	73
10	3	0.1	50	40
11	3	0.55	275	73
12	3	1	500	85
13	3	0.55	275	73
14	1	0.1	275	55
15	1	0.55	500	82
16	3	0.1	500	53
17	1	1	275	91

of the 17 experimental runs which can be observed in the normal plot (Fig. 10.1). The relationship between actual and predicted values of R (%) response is illustrated in Fig. 10.2.

For optimization of the variables, the BBD-RSM was utilized, and 17 runs of experiments were created. By the quadratic model (Eq. 10.2) the metal sorption percentage (R %: response) for optimized values of the factors can be calculated using Eq. 10.2.

$$\begin{aligned} \%removal = & 72.8 + -1.25 * A + 20.625 * B + 2.375 * C + 3.25 * AB \\ & + -7.75 * AC + -4 * BC + 1.6 * A^2 + -5.15 * B^2 + -1.15 * C^2 \end{aligned} \quad (10.2)$$

The 3D surface and contour plots of the interaction of pH with adsorbant dosage and metal concentration on the metal sorption capacity are depicted in Figs. 10.3, 10.4 and 10.5). For maximal adsorption of the metal using the sorbent the optimal pH was 2.7 and the adsorbant dosage was 0.89 g a maximal removal of 97.5% was achieved which was validated with an independent run.

Table 10.3 ANOVA table RSM BBD design

Source	Sum of squares	df	Mean square	F-value	p-value	
Model	3934.07	9	437.12	56.61	< 0.0001	significant
A-pH	12.50	1	12.50	1.62	0.2439	
B-adsorbent dosage	3403.12	1	3403.12	440.74	< 0.0001	
C-Cr concentration	45.13	1	45.13	5.84	0.0463	
AB	42.25	1	42.25	5.47	0.0519	
AC	240.25	1	240.25	31.11	0.0008	
BC	64.00	1	64.00	8.29	0.0237	
A ²	10.78	1	10.78	1.40	0.2760	
B ²	111.67	1	111.67	14.46	0.0067	
C ²	5.57	1	5.57	0.7212	0.4238	

Design-Expert® Software

%removal

Color points by value of %removal:

40 ■ ■ ■ 91

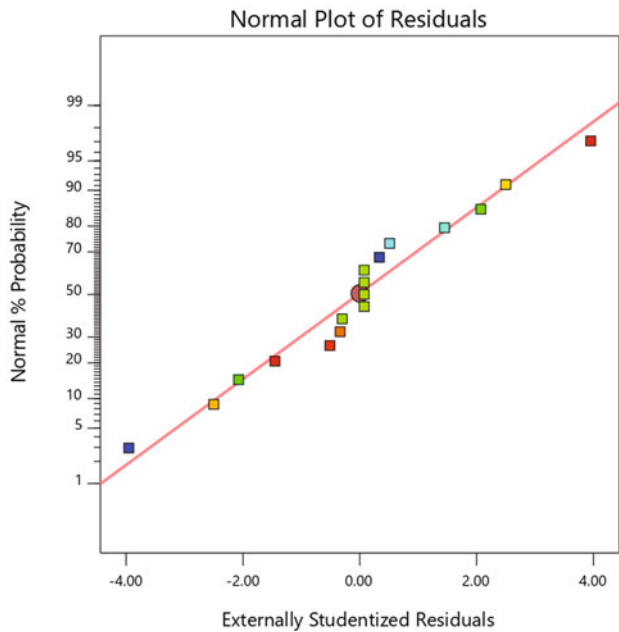


Fig. 10.1 Normal plot

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Color points by value of %removal:

40 91

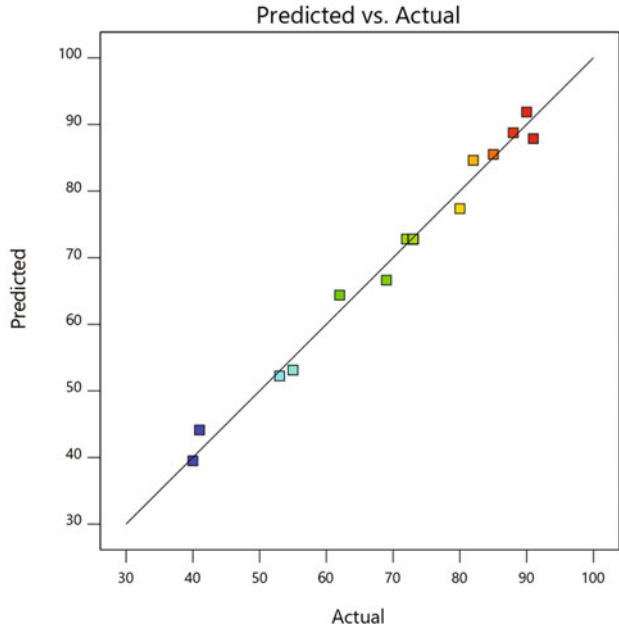


Fig. 10.2 Actual versus predicted

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Factor Coding: Actual

%removal

- Design points above predicted value
- Design points below predicted value

40 91

X1 = A: pH
X2 = B: Adsorbent dosage

Actual Factor
C: Cr Concentration = 275

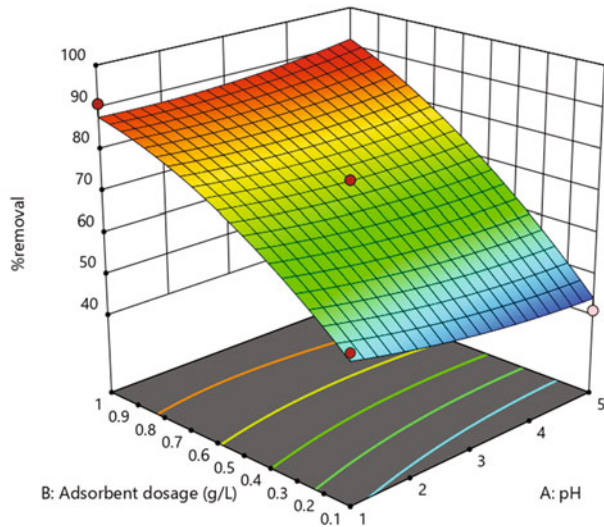


Fig. 10.3 Adsorbent dosage and pH impact on metal sorption.

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Factor Coding: Actual

%removal

- Design points above predicted value
- Design points below predicted value

40 91

X1 = A: pH
X2 = C: Cr Concentration

Actual Factor
B: Adsorbent dosage = 0.55

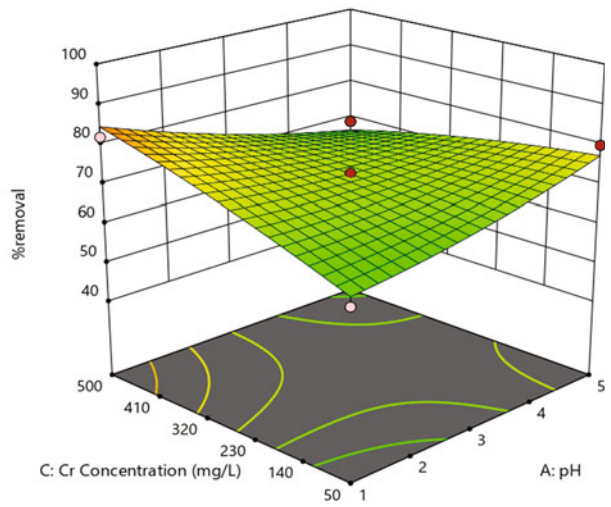


Fig. 10.4 Metal concentration and pH impact on sorption

Design-Expert® Software
Factor Coding: Actual

%removal

- Design points above predicted value
- Design points below predicted value

40 91

X1 = B: Adsorbent dosage
X2 = C: Cr Concentration

Actual Factor
A: pH = 3

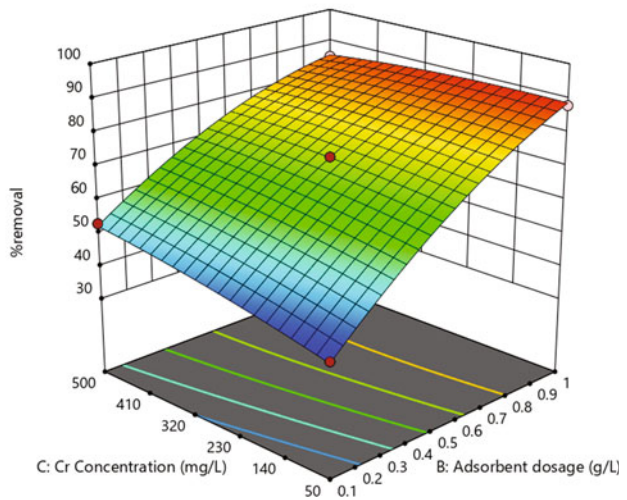


Fig. 10.5 Metal concentration and adsorbent dosage impact on sorption

10.4 Conclusion

This study is the first to show that the waste from *Hedychium flavum* can be a good adsorbant to remove metals from waste streams as a maximal removal of 97.5% of metal removal can be achieved under the optimal pH of 2.7 and adsorbant dosage of 0.89 g.

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Chapter 11

Impact of *Acinetobacter Baumannii* on Dye Degradation and a Molecular Analysis Study



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Abstract *Acinetobacter baumannii* was isolated from polluted soil. An attempt to study the *A. baumannii* to degrade dye was explored. It was found effective against azo dye and was able to completely degrade the dye under 48 h in a shake flask. Molecular analysis on the isolate *A. baumannii* was performed together with *Hedychium flavum*, and the sequence was submitted to the NCBI database to procure accession number MT192652.1. Response surface Methodology-Box-Behnken design (RSM-BBD) was used to optimize the condition and achieve 98–99% dye decolorization.

Keywords *Acinetobacter baumannii* · Dye degradation · Optimization · Molecular analysis

11.1 Introduction

The demand on every industry to provide enough consumer goods for the world's now-staggering 8 billion people was brought on by the Industrial Revolution, which aided in the population expansion that led to that number. The textile sector is vital to meeting the needs of a growing global population. An estimated one million tonnes of dyes and pigments are produced and consumed each year, with a sizable amount of these synthetic colors ultimately finding their way into the environment via wastewater discharges. The environment and human health are in grave danger due to this. Therefore, much water is consumed in the textile industry during the dying process, and the resulting effluent contains a lot of organics and hazardous chemicals. Furthermore, the dissolved oxygen levels of aquatic life are negatively impacted when this wastewater is dumped into water bodies without treatment. Thus, even at 1 ppm, dyes can contribute to diminished oxygen solubility and sunlight dispersion,

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which are essential for aquatic life (Arora 2014; Barikbin et al. 2017). As a result, these effluents must be treated to remove the colors before being released into waterways. Recent reports have highlighted the public health risk posed by wastewater discharges containing the dye, which can cause eye burns, nausea, vomiting, and diarrhea. Numerous treatment technologies have been developed for decolorizing and degrading dyes in wastewater; hence, it is crucial to find an effective treatment method for wastewater containing dyes. Adsorption, reverse osmosis, ultrafiltration, chlorination, and biological approaches, including aerobic and anaerobic treatments, are also examples of such processes. However, most of these conventional pollution control methods only relocate the pollutants from one phase to another rather than transforming them into a less harmful product (Unnikrishnan et al. 2018).

Compared to conventional approaches, bioremediation is a safer, cheaper, greener, and more practical option for dealing with textile waste (Fletcher et al. 2021). Using microbes in the form of bacteria, fungi, algae, yeast, and mixed culture for bioremediation is an exciting new field (Li et al. 2019). The bacteria have an advantage in treating textile effluents due to their capacity to adapt and break down textile effluents at high concentrations. Bacterial methods for degrading synthetic dyes, especially azo dyes, may be helpful (Mishra et al. 2020). Research and development cannot proceed without isolating, identifying, and preserving various dye-degrading bacteria. To decolorize and break down azo dyes, scientists have experimented with a wide range of oxidoreductive enzymes (Srinivasan et al. 2017). Thus, optimizing the significance and interaction among protein ligands can be done cheaply through molecular docking and simulation. Dye decolorizing bacterial systems for real-time wastewater treatment in the textile industries can be effectively screened using this combinatorial approach by first employing bioinformatics tools, then conducting analysis in a wet lab (Gao et al. 2018).

The application of response surface methodology (RSM) has been validated as an efficient way to optimize processes involving several interrelated variables. When designing, modeling, and optimizing a process, the RSM, known as a Box Behnken design (BBD), has proven to be effective in estimating the correlation coefficient for a specific model, resulting in lower experimental errors and better data fitness.

This study focuses on the dye degradation capability of the isolated bacteria, enhancing its potential via optimizing process parameters like the inoculum load, temperature, and dye concentration.

11.2 Materials and Methods

11.2.1 Material

Coimbatore, Tamil Nadu, India, is home to a sizable textile sector, where we sourced our azo dye. The dye has a molecular weight of 967.5 g/mol and an absorption wavelength of 530 nm.

11.2.2 Sample Collection and Isolation of Dye-Degrading Bacteria

Under aerobic circumstances, samples of textile effluent were gathered in Coimbatore, Tamil Nadu, India. Samples that had been adequately diluted were plated on Luria–Bertani (LB) agar that contained azo dye. For additional research, the bacterial strain that could stand up to 10 g/L of the dye was chosen. The chosen bacterial isolates' capacity to discolor was confirmed (Shah 2016).

11.2.3 16S rRNA Sequencing and Submission

Using a kit, genomic DNA was isolated from the bacterium (Gislin et al. 2018). In order to find closely identical sequences, the 16S rRNA sequence was then entered into a National Center for Biotechnology Information (NCBI) blast similarity search engine.

11.2.4 Decolorization Studies

One milliliter of the isolated bacterial strain was added to a flask containing 50 mL 500 mg/L of azo dye amended in LB broth and incubated at 37 °C for 3 days. With the aid of an ultraviolet–visible spectrophotometer, the degree to which a dye lost its color was determined by observing changes in the dye's absorbance at 530 nm (Satar and Husain 2011). The percentage rate of dye decolorization, E, was calculated using the formula:

$$E = [(A_0 - A_t) / A_0] * 100\% \quad (11.1)$$

where A_0 is the preliminary absorbance and A_t is the ultimate absorbance.

11.2.5 Immobilization of Potential Isolate

Calcium alginate beads were used to immobilize the potential isolate. Beads made of calcium alginate are durable and may be re-used after being immobilized by encapsulation. Hot water was added until wholly dissolved, followed by adding 30 g/L of sodium alginate and 5% of the bacterial culture while stirring frequently. The cell/sodium alginate combination was extruded drop by drop into a chilled, sterile calcium chloride solution using a syringe to produce calcium alginate beads with a diameter of 2 mm. CaCl_2 and sodium alginate are combined to create calcium

Table 11.1 Variables chosen for RSM optimization

Factor	Name	Units	Type	Minimum	Maximum
A	Temperature	Celsius	Numeric	25	35
B	Immobilized inoculum	g/L	Numeric	0.1	0.5
C	Incubation time	hours	Numeric	12	36

alginate beads. For increased hardness and durability, calcium alginate beads were resuspended in a calcium chloride solution and chilled in the refrigerator for 24 h.

11.2.6 Optimization of Process Variables

Using the Response Surface Methodology (RSM), statisticians and mathematicians can forecast ideal circumstances with the fewest runs. We studied the connections between temperature, incubation period, and inoculum volume using a three-level factorial Box-Behnken design based on the one factor at a time (OFAT) technique (BBD). The experimental intervals used in the BBD are shown in Table 11.1.

11.3 Results and Discussion

11.3.1 Isolation of Potential Bacteria and Identification

Decolorization effectiveness was evaluated after isolating a bacterial strain that can survive in solutions containing up to 10 g/L of azo dye. It was observed that the bacteria entirely removed maximal color up to 98% (Prasad and Aikat 2014; Permpornsakul et al. 2016). 16S rRNA sequencing helped identify the isolate as *A. baumannii*. *A. baumannii* is a gram-negative bacterium that can be detected in soil and water samples on rare occasions. Few studies have explored the potential of this bacteria for decolorizing dyes. GenBank now contains the sequence in the entry as Accession no. MT192652.1. Results from a phylogenetic study are shown in Fig. 11.1 (seq 1) (Unnikrishnan et al. 2018).

```

1 cgagcggggg tagggagcctt gctactggtc ctagcggcgg acgggtgagt aatgcttatg.
61 aatctgccta ttatggggg acaacatctc gaaagggatg ctaataccgc atactgcta.
121 cgggagaaag caggggatct tcggaccttg cgctaataga tgagcctaag tcggattagc.
181 tagttgtggt ggtaaaggcc taccaaggcg acgatctgta gcggtctga gaggatgatc.
241 cgccacactg ggactgagac acggcccaga ctctacggg aggcagcagt ggggaatatt.
301 ggacaatggg ggaacacctg atccagccat gccgcgtgtg tgaagaaggc cttatggtg.
361 taaagcactt taagcgagga ggaggctact gtaattaata cctatggata gtggacgtta.
421 ctgcgagaat aagcaccggc taactctgtg ccagcagccg cgtaataca gagggtcgga.
481 gcgttaatcg gatttactgg gcgtaaagcg tgcgtaggcg gcttttaag tcggtatgta.

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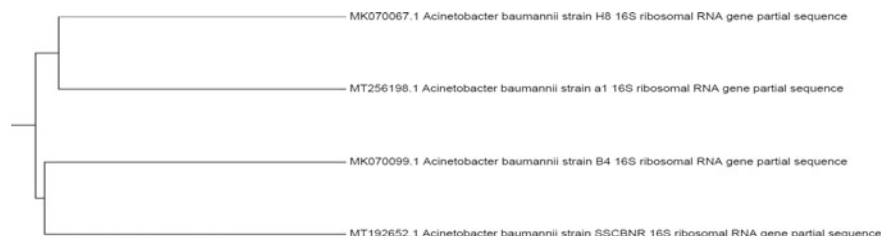


Fig. 11.1 Phylogenetic tree for *Acinetobacter baumannii* (Acc.No. MT192652)

541 aatccccgag cttacttgg gaattgcatt cgatactggt gagctagagt atgggagagg.
 601 atggtagaat tccaggtgta gcggtgaaat gcgtagagat ctggaggaat accgatggcg.
 661 aaggcagcca tctggcctaa tactgacgct gga.

Sequence 1: 16 s RNA partial Sequence of *Acinetobacter*.

11.3.2 Optimization Using Box-Behnken Design

Optimizing decolorization processes with BBD is standard practice. For example, in 2013, Papadopoulou et al. explored the potential of *Pleurotus pulmonaris* AMRL 177 with BBD for decolorizing Remazol brilliant blue, Fazli et al. (2010) employed BBD to enhance *Ganoderma* species-based reactive blue 19 decolorization. Time, temperature, and inoculum volume were used in the BBD experiment. Table 11.2 details the design, while Table 11.3 provides an ANOVA for the response surface model.

$$\begin{aligned} \%removal = & 82.4 + 0.625 * A + 18 * B + 10.875 * C + 0.25 * AB \\ & + -1.5 * AC + 0.25 * BC + -1.95 * A^2 + -7.7 * B^2 + -3.45 * C^2 \end{aligned}$$

When R1 represents the response, A, B, and C represent temperature, immobilized inoculum amount, and incubation period, respectively, while the other variables represent the interaction of one variable against another. Design expert software was utilized to simulate the design and analyze the results of the testing runs. The model's F-value of 58.26 suggests that it is significant at p0.0001 (Table 11.3). The most critical model terms in the current model are B, C, B2, and C2. Based on the normal-plot, actual vs. predicted plots, the actual and expected outcomes (response- dye decoloration) were good (Figs. 11.2 and 11.3). The outcome was equivalent to that of Unnikrishnan et al. (2018).

It is also possible to predict and compare the interaction of the design parameter on each other and their effect on the dye decolorization evident from the 3-D contour plots (Figs. 11.4a, b, and c) (Papadopoulou et al. 2013).

Table 11.2 Experimental runs with dye decolorization outcome

	Factor 1	Factor 2	Factor 3	Response 1
Run	A: Temperature	B: Immobilized Inoculum	C: Incubation time	% Removal
	Celsius	g/L	Hours	
1	25	0.3	12	61
2	30	0.1	12	45
3	25	0.3	36	88
4	30	0.3	24	82
5	35	0.5	24	91
6	25	0.1	24	55
7	25	0.5	24	93
8	30	0.1	36	64
9	35	0.1	24	52
10	30	0.3	24	83
11	35	0.3	12	69
12	30	0.3	24	83
13	30	0.5	12	78
14	30	0.3	24	82
15	30	0.3	24	82
16	30	0.5	36	98
17	35	0.3	36	90

Table 11.3 ANOVA analysis information

Source	Sum of Squares	df	Mean Square	F-value	p-value	
Model	3891.11	9	432.35	58.26	<0.0001	Significant
A-Temperature	3.13	1	3.13	0.4211	0.5371	
B-Immobilized inoculum	2592.00	1	2592.00	349.26	<0.0001	
C-Incubation time	946.13	1	946.13	127.49	<0.0001	
AB	0.2500	1	0.2500	0.0337	0.8596	
AC	9.00	1	9.00	1.21	0.3072	
BC	0.2500	1	0.2500	0.0337	0.8596	
A ²	16.01	1	16.01	2.16	0.1853	
B ²	249.64	1	249.64	33.64	0.0007	
C ²	50.12	1	50.12	6.75	0.0355	

Design-Expert® Software

%removal

Color points by value of

%removal:

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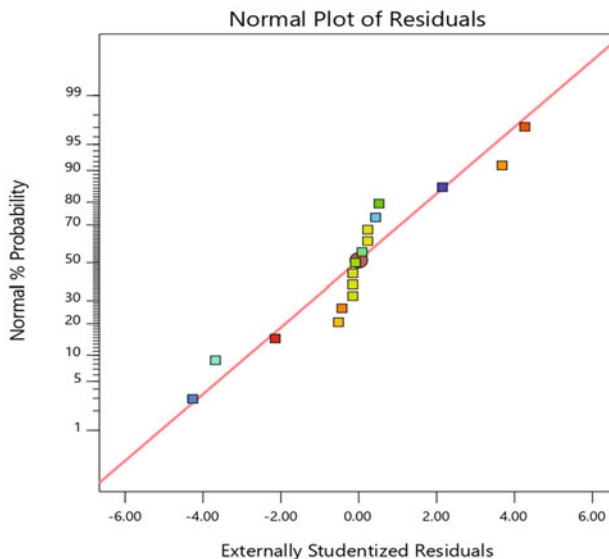


Fig. 11.2 Normal plot

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Color points by value of

%removal:

45 98

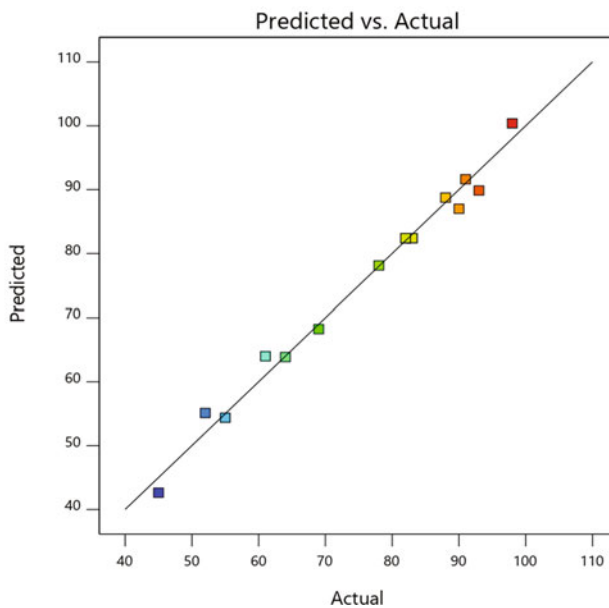


Fig. 11.3 Predicted versus actual plot

(a)

Design-Expert® Software
Factor Coding: Actual

%removal

● Design points above predicted value

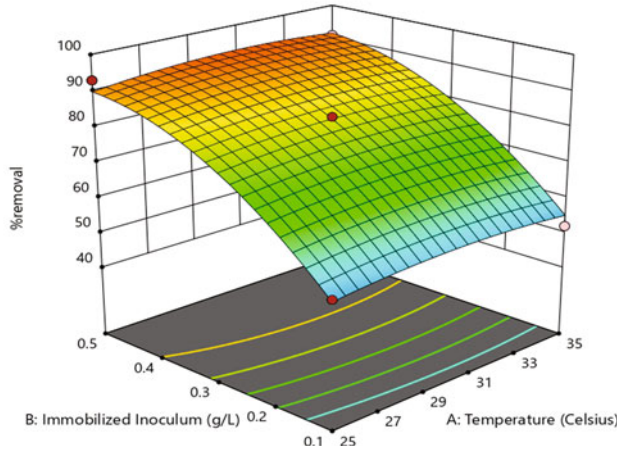
○ Design points below predicted value

45 98

X1 = A: Temperature
X2 = B: Immobilized Inoculum

Actual Factor

C: Incubation time = 24



(b)

Design-Expert® Software
Factor Coding: Actual

%removal

● Design points above predicted value

○ Design points below predicted value

45 98

X1 = A: Temperature
X2 = C: Incubation time

Actual Factor

B: Immobilized inoculum = 0.3

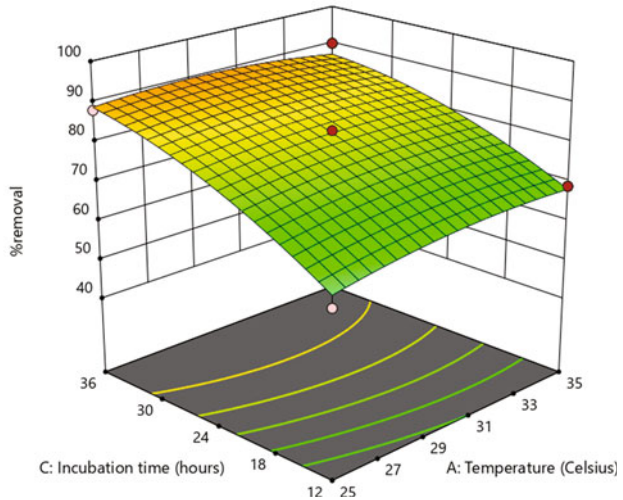


Fig. 11.4 a 3D contour plot of temperature versus immobilized inoculum on dye decolorization b 3D contour plot of temperature vs. incubation time on dye decolorization c 3D contour plot of incubation time vs. immobilized inoculum on dye decolorization

(c)

Design-Expert® Software
Factor Coding: Actual

%removal

● Design points above predicted value

○ Design points below predicted value

45 98

X1 = B: Immobilized Inoculum

X2 = C: Incubation time

Actual Factor

A: Temperature = 30

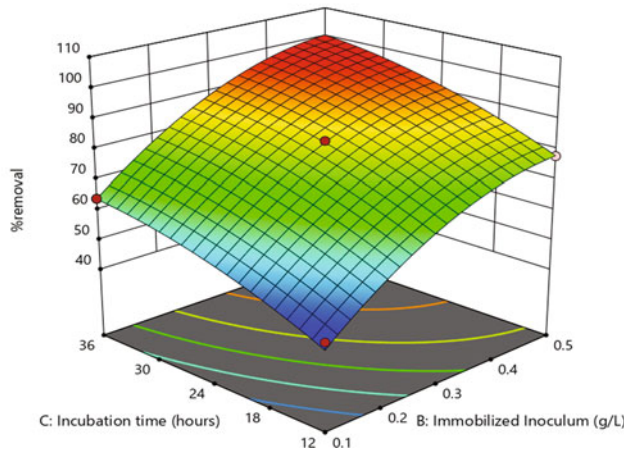


Fig. 11.4 (continued)

11.3.3 Model Validation and Experimental Confirmation

The RSM BBD output was validated with an independent run to achieve peak dye decolorization. For this dye decolorization, the experimental decolorization rate (99.15%) was achieved for an optimized condition of 28 °C with an inoculum load of 450 mg/L and an incubation period of 34 h.

11.4 Conclusion

Acinetobacter baumannii dye decolorization capabilities were evaluated and optimized to the experimental condition via RSM-BBD. As a result, 99% of dye discoloration was achieved for optimized conditions of 28 °C temperature, inoculum load of 450 mg/l, and the maximal dye discoloration was achieved in 35 h. Hence this isolate can be used for larger-scale exploration of dye degradation under optimal conditions.

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Chapter 12

Photocatalytic Degradation of Reactive Orange M2R Using Green Route Synthesized Copper Oxide Nanoparticles and Its Optimization Studies



Easvaran Ramya, Alagu Thirumurugan, Natanamurugaraj Govindan, Jayaseelan Aravind, and Sriramulu Gobikrishnan

Abstract The photocatalytic degradation approach for industrial dyes in the presence of appropriate nanocatalyst offers a possible way for the complete removal of various pollutants from the aqueous environment. Therefore, in this study, we have made an attempt on use of bio-synthesized copper oxide (CuO) nanoparticles harnessed by neem extract for the degradation of reactive orange M2R (RO-M2R). The significant catalytic degradation of dye was found to be pH10, catalytic dosage at 10 ppm, and dye concentration at 3 gL⁻¹ and irradiation time at 5 h through one factor at a time (OFAT). Further, Box Behnken design was applied to optimize and investigate the interaction between these four variables for the degradation of dye. It was predicted that maximum degradation was 84.04% under optimum conditions of pH at 11.84, catalytic dosage at 2.20 gL⁻¹, dye concentration at 7.49 ppm and irradiation time of 4 h. The validity of the optimal level predicted under the RSM was established by an independent experiment.

Keywords CuO nanoparticles · RSM · Box Behnken design · Photocatalytic degradation

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12.1 Introduction

Water pollution has become a very sensitive issue in the world due to inorganic and organic pollutants. The primary source of contamination of water is the organic dyes discharged by textile industries. There are more than 10,000 commercially available dyes with over 7×10^5 tons of dyestuff produced. In which, concerning 36,000 dye/year are consumed by textile industries across the world. 20% of the dyes in the world production are lost during the dyeing process and is discharged in the textile wastewater (Reza et al. 2017; Esplugas et al. 2002; Houas et al. 2001). In particular, the principal and most adaptable classes of azo dyes are presently utilized in the dyeing of different materials such as textiles, leather, plastics, and cosmetics. The azo dyes containing effluents are released into rivers, lakes, and ground waters for the period of dying process that contains mutagenic and carcinogenic health hazards. As a result, it leads to severe environmental problems due to their good stability under ambient conditions (Mathur et al. 2012).

Organic pollutants are challenging to biological degradation; also, conventional treatment processes are not succeeded to attain their complete elimination of dye molecules. Even though, for a decade, researchers have attempted various methodologies such as membrane filtration, chemical coagulation/flocculation, ion exchange, precipitation, adsorption, biological degradation, and ozonation, etc. Still, these methods are not capable enough and have more limitations. UV light is generally applied to excite a semiconductor surface to produce photoinduced holes or reactive oxygen species (ROS), such as hydroxyl and superoxide radicals in the photocatalytic advanced oxidation technology. Then, the ROS interrelate with organic compounds and guides to their oxidation and overall degradation (Jantawasu et al. 2009). Moreover, for a decade nanocatalyst development has highly attracted researchers to work towards the degradation of industrial wastewater pollutants. Apart from dye degradation studies by nanoparticles, Magnetic Moringa Oleifera (MMO) have also been used as a coagulant for palm oil waste water treatment (Noor et al. 2022).

There are various studies reported on dye degradation by metallic nanoparticles. Chaibakhsh et al. (2016) have reported on photocatalytic degradation of neutral red dye using titanium oxide (TiO_2) nanocatalyst and its optimization by Box-Behnken design. Recently, Uddin and Baig (2019) have studied on the removal of methyl orange dye using cobalt oxide (Co_3O_4) nanoparticles and Dawoud et al. (2020) have also reported on photocatalytic degradation of organic dye of Rhodamine B by using silver (Ag) doped Zirconium oxide (ZrO_2) nanoparticles. Recently, CuO NP's have been prepared from various sources such as *Canthium coromandelicum* leaves extract (Selvam et al. 2022), *Punica granatum* (Kaur et al. 2022), *Pistacia vera* L. hull (Ltaief et al. 2021), *Dictyota dichotoma* endophytic fungi (Kumar et al. 2022), *Elaeagnus indica* (Indhira et al. 2022), *Solanum tuberosum* (Hamami and Javanbakht 2021) were used as a catalyst for textile dye degradation.

Most of the reports on dye removal by magnetic nanoparticles have been performed by 'one variable at a time' (OFAT) approach that perfectly presumed that all variables are independent and do not represent the combined effect of all

the variables involved. Moreover, this method is a prolonged one and also requires a higher number of experiments to determine optimum levels, which may or may not be trustworthy (Farooq et al. 2017). These limitations of the conventional methods can be reduced by response surface methodology (RSM), using this statistical design of experiment.

Thus, herein, biological route synthesized CuO nanoparticles assisted photocatalytic degradation of RO-M2R dye, and their optimization studies were performed using Box-Behnken design to maximize the percentage of dye degradation.

12.2 Materials and Methods

12.2.1 Photocatalytic Degradation Confirmation

RO-M2R was obtained from Precision Scientific Co. Coimbatore, India. The photocatalytic degradation experiment of RO-M2R was performed under sunlight irradiation. The photocatalyst CuO NP's 2 gL^{-1} was added to 50 ppm of RO-M2R dye solution at neutral pH to establish the adsorption equilibrium between the dye molecule and CuO NP's. The resulting solution was exposed to sunlight for 4 h, and control was maintained without addition of nanoparticles. These suspensions were measured against absorbance at 495 nm. The percentage of degradation was calculated from the following equation;

$$\% \text{ of Degradation} = (C_0 - C/C_0) \times 100$$

where, C_0 = Initial concentration of dye, C = Final concentration of dye.

12.2.2 Preliminary Experiment

Preliminary experiments were conducted by one factor at a time (OFAT) approach to identify the ranges of four parameters such as pH, dye concentration, catalytic dosage and irradiation time for the degradation of RO-M2R. Statistical design of experiment consists of a variation of several trials at a time when compared to one factor at a time approach in which only one factor is changed and keeping all other factors constant.

12.2.2.1 Evaluation of Process Parameters by OFAT

The dye solution was taken out in reaction vessel, and the pH (3, 4, 5, 6, 7, 8, 9, 10, 11 and 12) was adjusted by using a pH meter (Elico LI 120, India). Similarly, the

different concentrations of RO-M2R working solutions (10, 20, 30, 40, 50, 60, 70, 80, 90 and 100 ppm) were prepared and kept the irradiation time for 4 h to study the effect of dye concentration on dye degradation. Likewise, the different concentrations of CuO nanoparticles (0.5, 1, 1.5, 2, 2.5, 3, 3.5, 4, 4.5 and 5 g/L) were varied and the reaction vessel was kept under sunlight for 4 h. Similarly, the different reaction periods (0, 50, 100, 150, 200, 250, 300, 350, 400 and 450 min) were chosen to check the percentage of degradation. Finally, 2 g/L of CuO nanoparticles was added to various ranges of pH and concentration of RO-M2R dye solution to establish the adsorption equilibrium between the dye molecule and CuO nanoparticles.

12.2.3 Response Surface Methodology for Optimization

Response surface methodology (RSM) was utilized to optimize the four parameters were catalytic dosage, pH, dye concentration and irradiation time. The four parameters were selected as independent variables, and the degradation percentage of RO-M2R was the response variable. Box-Behnken model was preferred to study the combined effect of four independent variables, and Design-expert 11 version was used to explain the response surface. The model was statistically examined. Analysis of variance (ANOVA) employed Fischer's *F*-test to evaluate the overall significance of the model, correlated probability values, and coefficient of determination to estimate the goodness of fit of the regression model. The apt polynomial equation was further conveyed the three-dimensional plots (3D) form, which depicted the interactions graphically. Finally, the central and interaction effects, the 3D plots were analyzed to check the accuracy of the model (Kanmani et al. 2013).

The following second-order polynomial equation explains the connection between the dependent and independent variables:

$$Y = \beta_0 + \beta_1A + \beta_2B + \beta_3C + \beta_{11}A^2 + \beta_{22}B^2 + \beta_{33}C^2 + \beta_{12}AB + \beta_{13}AC + \beta_{23}BC \quad (12.1)$$

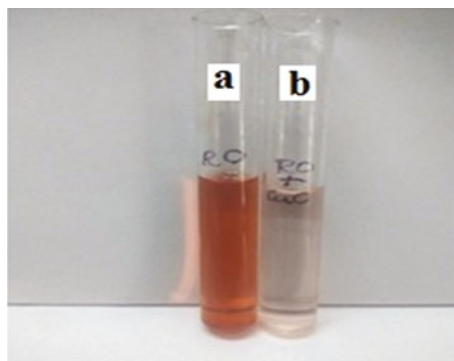
Here the *Y* is the predicted response, β_0 is the intercept, $\beta_1, \beta_2, \beta_3$, are the linear coefficients, $\beta_{11}, \beta_{22}, \beta_{33}$, are the squared coefficients, and $\beta_{12}, \beta_{13}, \beta_{23}$ are the interaction coefficients.

12.3 Results and Discussion

12.3.1 Confirmation of Photocatalytic Degradation

The previously synthesized CuO nanoparticles (Thirumurugan et al. 2017) were used for the photocatalytic degradation of RO-M2R under sunlight. These

Fig. 12.1 Confirmation of photocatalytic degradation of RO-M2R using CuO NP's under sunlight



nanoparticles were already proved its photocatalytic degradation of 4-nitrophenol as a model (Ramya et al. 2019) and photocatalytic degradation of reactive red 120 (Thirumurugan et al. 2017). Thus, in this study, the photocatalytic degradation of RO-M2R was studied by using green synthesized CuO NP's under sunlight, as shown in Fig. 12.1. After the addition of nanoparticles, the orange color of the solution decreased significantly, become colorless with 74% degradation, and no changes were observed in control. The dye degradation efficiency of nanoparticles may be due to differences in size, shape and catalytic activity (Moon et al. 2015). Kumar et al. (2013), has also reported that metal nanoparticle synthesized by green approach was effectively degraded the dye with periodically increase in time. There are various parameters affecting the photocatalytic degradation, such as pH, dye concentration, catalytic dosage, irradiation time, reaction temperature, light intensity and the presence of electron acceptors. In particular, we investigated the effect of pH, dye concentration, catalytic dosage, and irradiation time through one factor at a time (OFAT) approach to study the impact of each factor ranges for the photocatalytic degradation of RO-M2R.

12.3.2 Evaluation of Process Parameters by OFAT

Effect of pH is one of the main factors for the degradation of industrial dye. To investigate the effect of pH, the experiments were carried out at different pH ranges from 3 to 12 at 2 gL^{-1} of CuO NP's, 50 ppm of RO-M2R and irradiation time of 4 h. The result showed that by increasing the pH of the dye solution, the dye degradation was getting increased, as shown in Fig. 12.2. It was observed that degradation rate increases, with an increase in pH and the maximum photocatalytic degradation, was found to be at pH 10. The cause can be due to efficient formation of hydroxyl radicals in alkaline medium. Excess of hydroxyl anions increases the formation of hydroxyl radicals as the main oxidizing function responsible for dye degradation (Gopalapp et al. 2012). Similarly, Devadi et al. (2014) has observed the maximum degradation

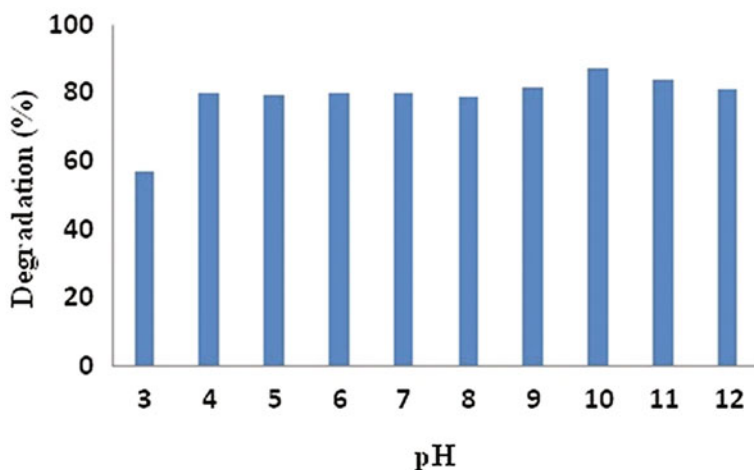


Fig. 12.2 Degradation of RO-M2R at different pH

of methyl blue at pH 11 and at under acidic condition the degradation efficiency get decreased due to the columbic repulsion. Also, Daneshvar et al. (2003) reported that in the presence of ZnO NP's, the photodegradation was considerably improved at high pH (pH = 11), While the lower at low pH is due to the photodecomposition of ZnO to Zn^{2+} takes place in acidic and neutral solutions and that efficient formation of hydroxyl radicals occurs in alkaline solution.

The effect of dye concentration for degradation of RO-M2R was varied between 10 to 100 ppm where pH was kept at 10, catalytic dosage at 2 gL^{-1} and irradiation time for 4 h. The degradation of RO-M2R at different concentration of dye is shown in Fig. 12.3. The result showed that up to 40 ppm the percentage of degradation was approximately the same and after 40 ppm the degradation rate was reduced. While maintaining a constant catalyst dose, it caused the fixed number of catalysis sites to be saturated rapidly, which is due to increasing dye concentration in the solution (Mohan and Pittman 2006). In addition, the light transmittance in the solution may be decreased due to the increased dye concentration; decreased light infiltration can diminish the activation rate of nanoparticles and delay the formation of OH radicals, results in decreased photocatalytic degradation.

Figure 12.4 shows the effects of different concentrations of nanoparticles as catalysts on degradation. The increase in concentration up to 3 gL^{-1} contains a high amount of active site and thus producing more free radicals which uninterruptedly increase the degradation of RO-M2R, and beyond 3 gL^{-1} there was a decrease in degradation rate. According to Akpan and Hameed (2009), this was due to the formation of turbid in the solution, which reduces the penetration of sunlight. Hence, it blocks the radiation to enter into the RO-M2R solution. Recently, Mortazavian et al. (2019) has reported in a study that the amounts of catalyst particles might aggregate leading to a reduced number of active sites because the increase of TiO_2 particles in the solution might supply more active sites for the molecules of dye to be adsorbed

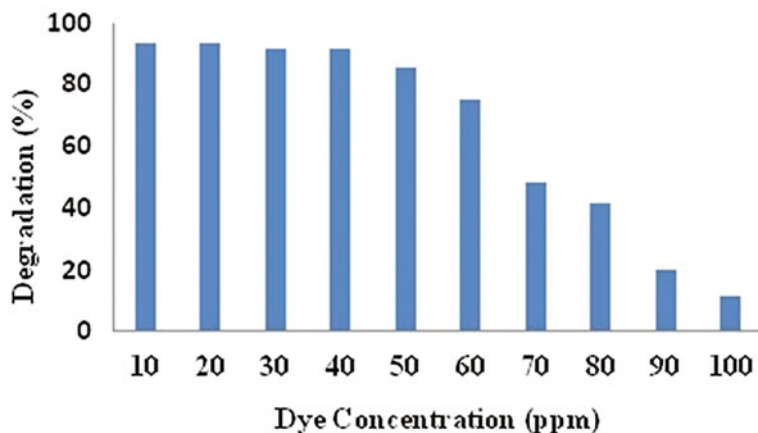


Fig. 12.3 Degradation of RO-M2R at different dye concentration

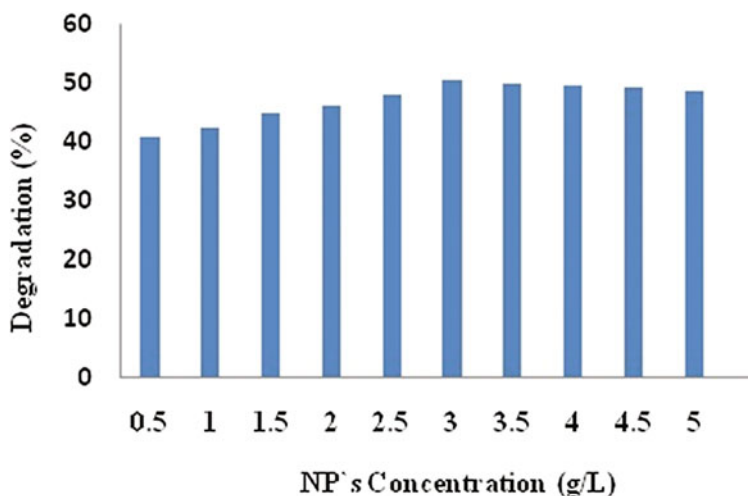


Fig. 12.4 Degradation of RO-M2R at different concentrations of CuO NP's

and degraded. This phenomenon might be one of the reasons for decreasing the degradation rate beyond 3 gL^{-1} of CuO NP's as a catalyst. Very recently, Hashemi et al. (2022) reported the optimization and assessment of photocatalytic degradation of methyl orange using silver nanoparticles synthesized through green route. As a result, observed that the methyl orange pollutant degradation was increased while increasing the concentration of nanocatalyst.

The effect of irradiation time on degradation was investigated by keeping catalyst amount at 3 gL^{-1} , pH at 10, and dye concentration at 40 ppm and varying the irradiation time of the dye solution under sunlight. Figure 12.5 shows that degradation rate

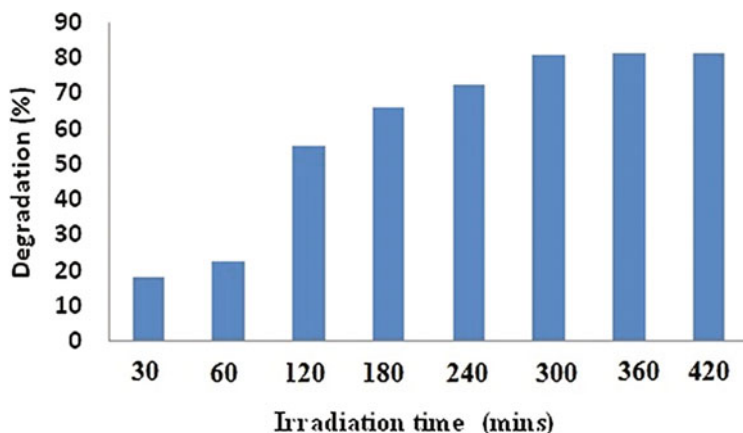


Fig. 12.5 Degradation of RO-M2R at different time intervals

increased as time increased, and this is due to the increased number of photons. The maximum degradation efficiency was observed at 5th h, and after that the degradation remains the same.

12.3.3 Optimization Using Response Surface Methodology

Box Behnken Design (BBD) was employed to evaluate the effect of variables on the degradation of RO-M2R in the presence of sunlight. The influence of operating parameters on the photocatalytic degradation efficiency of RO-M2R were assessed, four main factors were chosen: pH, dye concentration (ppm), catalytic dosage (gL^{-1}), irradiation time (hrs). A total of 29 experiments were used using Design expert software. The experimental outcomes of degradation of dye by nanoparticles were analyzed through RSM as tabulated in Table 12.1 to obtain an empirical model.

Based on the outcomes, an empirical relationship between the responses and independent variables were attained for the dye degradation expressed by the quadratic polynomial equation, as shown below.

$$\begin{aligned} \text{\% of Degradation} = & 59.19 + 14.14 * A - 4.52 * B + 0.19 * C - 2.13 * D \\ & + 0.38 * AB - 0.74 * AC + 0.80 * AD - 0.71 * BC \\ & - 1.27 * BD + 1.74 * CD + 4.08A^2 + 0.048B^2 + 1.94C^2 + 1.80D^2 \end{aligned}$$

Response surface methodology integrated the interaction effects of variables and helps us in simultaneously optimizing many process parameters within a minimum number of trial runs. Such statistically aided experimental designs can guide to considerably better performance of the nanoparticles. The P value acts as a tool

Table 12.1 Box-Behnken design of response surface methodology with real values and their responses

S.NO	A: pH	B: Nanoparticles concentration (g/l)	C: Dye concentration (ppm)	D: Irradiation time (hr)	Degradation (%)
1	8	2	22.5	5.5	54.5
2	12	2	22.5	5.5	80.09
3	8	4	22.5	5.5	47.1
4	12	4	22.5	5.5	74.2
5	10	3	5	4	66.85
6	10	3	40	4	65.18
7	10	3	5	7	58.51
8	10	3	40	7	63.8
9	8	3	22.5	4	53.82
10	12	3	22.5	4	80.22
11	8	3	22.5	7	49.25
12	12	3	22.5	7	78.84
13	10	2	5	5.5	67.3
14	10	4	5	5.5	55.51
15	10	2	40	5.5	69.2
16	10	4	40	5.5	54.57
17	8	3	5	5.5	48.65
18	12	3	5	5.5	80.65
19	8	3	40	5.5	49
20	12	3	40	5.5	78.04
21	10	2	22.5	4	64.73
22	10	4	22.5	4	60
23	10	2	22.5	7	62.35
24	10	4	22.5	7	52.55
25	10	3	22.5	5.5	60.85
26	10	3	22.5	5.5	58.02
27	10	3	22.5	5.5	57.83
28	10	3	22.5	5.5	61.34
29	10	3	22.5	5.5	57.89

for assessing the significance of each coefficient and is indicative of the interaction strength of all independent variable. Low values of P at ($P < 0.05$) indicate the high significance of the corresponding coefficients. In general, larger t , F and smaller P values indicate that the corresponding coefficient terms are significant.

From the ANOVA Table 12.2, the model F -value of the model (41.08) and values of $P > F (< 0.0500)$ implied that the model was significant statistically. The predicted

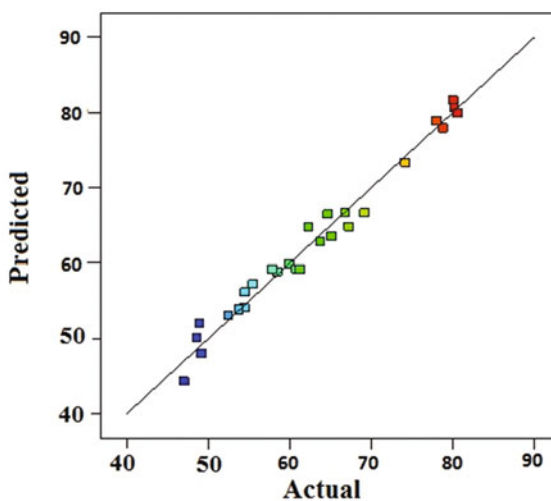
Table 12.2 Analysis of variance (ANOVA) and regression analysis for selected model

Source	Sum of squares	Degree of freedom	Mean squares	F value	P value	
Model	2855.57	14	203.97	41.08	<0.0001	Significant
Residual	69.51	14	4.97			
Lack of fit	57.23	10	5.72	1.86	0.2877	Not significant
Pure error	12.29	4	3.07			
Cor total	2925.09					

$R^2 = 0.9762$; $Adj.R^2 = 0.9525$; $Predicted R^2 = 0.8808$

R^2 of the 0.8808 is in reasonable agreement with the adjusted R^2 of 0.9525. Adequate precision measures signal-to-noise ratio (S/N ratio) and detect which experimental parameters generate signals that are large in comparison to the noise. The observed adequate precision ratio was 23.29, greater than 4, which indicated the selected model could be employed to precisely navigate the design space as reported by Chaibakhsh et al. (2016). The accuracy of the model is shown in Fig. 12.6, which evaluates the measured values against the predicted responses of the model for the degradation of RO-M2R. It can be observed that correlation between the experimental data and predicted values ($R^2 = 0.9762$) showed the data fit well with the model in the range studied.

Fig. 12.6 Predicted plotted against the actual experimental values for photocatalytic degradation of RO-M2R



12.3.3.1 Analysis of Response Surface Plots

Three-dimensional surface can be used as a graphical representation of the regression equation to find the optimum levels of the variables studied and widely used to achieve a better understanding of the interactions between variables within the ranges.

Figure 12.7 represents the interaction effect of pH, nanoparticles concentration and dye concentration and irradiation time on the degradation of dye from aqueous solution. There was a significant degradation observed by increasing pH till 12. Similarly, nanoparticles concentrations also influencing the degradation rate of dye molecules up to 2.20 gL^{-1} then degradation rate declines. Whereas, dye concentration and irradiation time were not influencing the degradation rate as pH and nanoparticles concentration. Recently, Chaker et al. (2021), has studied a statistical modelling optimization approach for efficient photocatalytic degradation of azo dye using cerium-doped mesoporous ZnO by central composite design. Their results indicated that all the variable chosen had an impact on the photocatalytic process and pH was the least factor against the effect of catalyst concentration.

The three-dimensional plots based on the interactions between the variables showed an increase in RO-M2R dye degradation as each variable increased to an optimum level, beyond the optimum level; decline rate could be observed (Fig. 12.7a–f). The optimal values obtained from the three-dimensional were also almost equal to the results obtained by the regression analysis (Eq. 12.1). From the model, a maximum dye degradation of 84% under the optimum condition of pH 11.8, nanoparticle concentration of 2.20 gL^{-1} , dye concentration of 7.49 ppm and irradiation time of 4 h were predicted. The experiment was conducted in triplicate to validate the prediction of the model, and the maximum degradation of RO-M2R was found to be 83.41% which was close to the predicted maximum dye degradation of the model.

12.3.4 Mechanism of Photocatalytic Degradation by CuO NP's

For the above observed results, a possible mechanism for the degradation of RO-M2R dye by CuO nanoparticles in aqueous solution can be proposed. A possible photocatalytic mechanism of CuO nanoparticles for the degradation of aqueous RO-M2R is shown in Fig. 12.8. Photocatalytic degradation of organic dyes through the utilization of CuO nanoparticles arises when the photocatalyst absorbs light of a suitable wavelength from sunlight. The light energy from the sun is exciting and promotes electrons in the valence band to conduction band of photocatalyst. As results, creation of positive charges (holes) and negative charges (electrons) on the valence and conduction bands respectively, leads to the formation of electron-hole pairs. This hole oxidizes the water molecule into hydrogen gas and hydroxyl radicals (OH^*), whereas electron reduces the oxygen molecules to form superoxide radicals ($\text{O}_2^{\bullet-}$). Thus radicals formed radicals, attack the dye molecule over the surface of

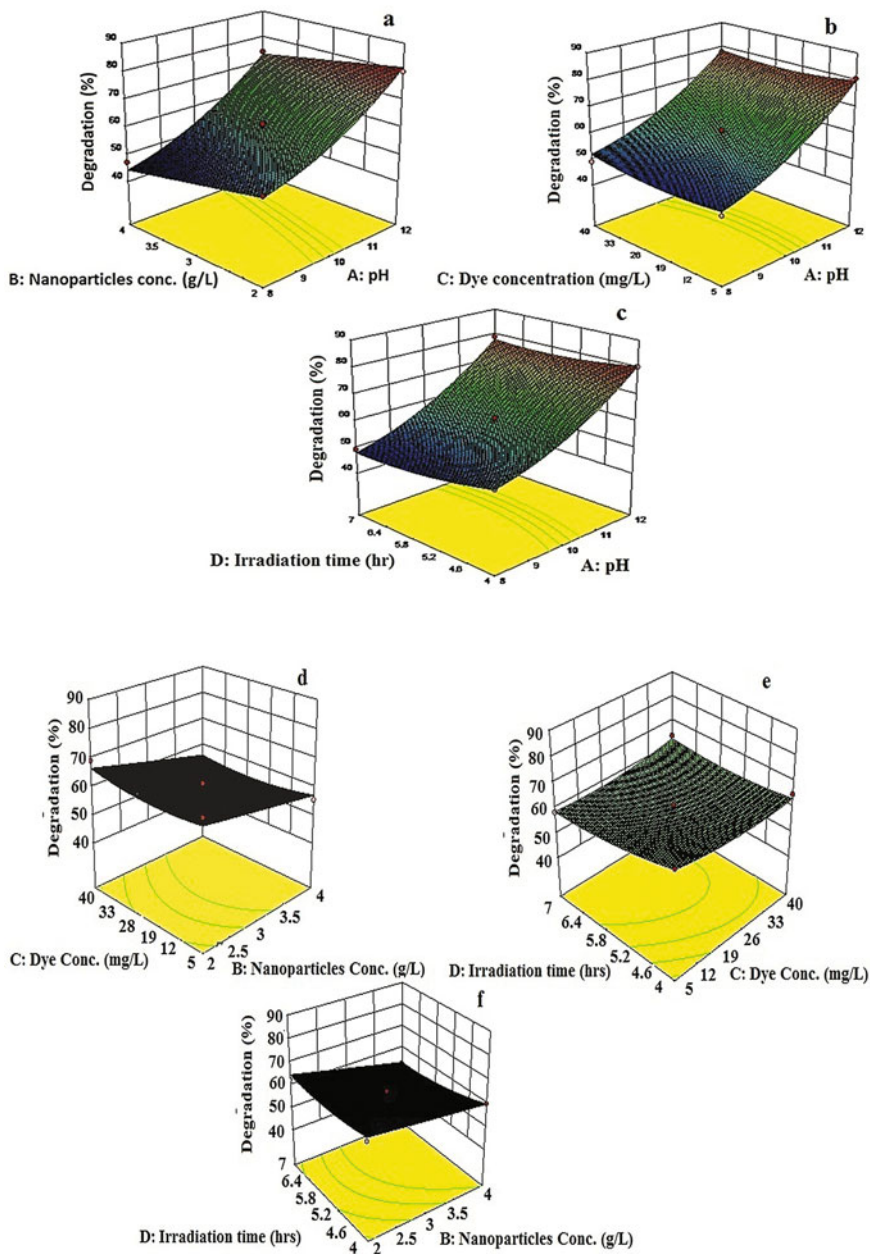


Fig. 12.7 Three dimensional surface plot showing the effect of **a** nanoparticles concentration (gL^{-1}) versus pH **b** dye concentration (mgL^{-1}) versus pH **c** Irradiation time (h) versus pH **d** Dye concentration (mgL^{-1}) versus nanoparticles concentration (gL^{-1}) **e** Irradiation time (h) versus dye concentration (mgL^{-1}) **f** Irradiation time (hrs) versus nanoparticles concentration (gL^{-1}) on degradation of RO-M2R

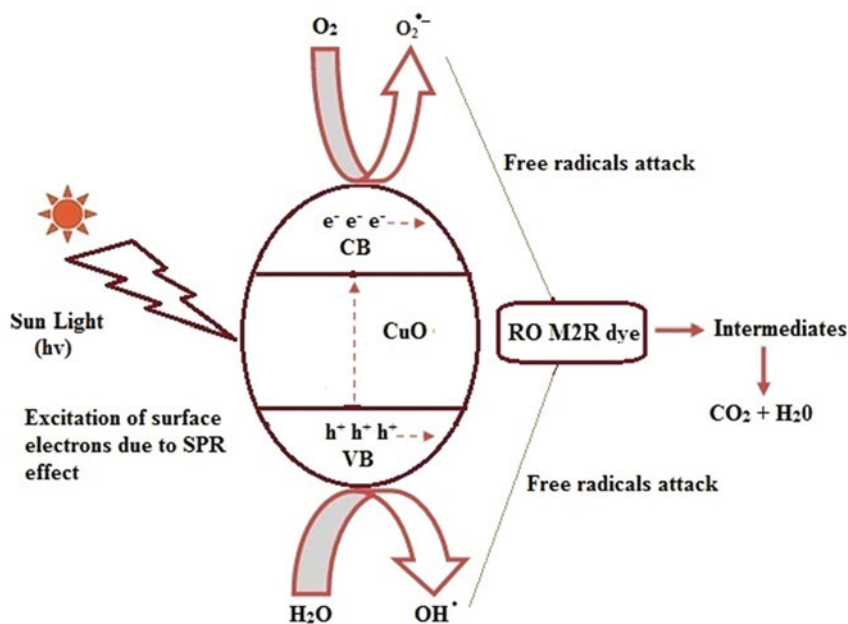


Fig. 12.8 A possible mechanism of photocatalytic degradation by CuO NP's

the CuO NP's and accomplish the dye degradation, and these formed radicals are also responsible for inactivation of bacteria through their cell walls degradation (Dalrymple et al. 2010).

12.4 Conclusions

We conclude that the photocatalytic degradation of RO-M2R by CuO NP's as a catalyst in the aqueous medium was developed a quadratic model using Response Surface Methodology as a functional relationship between catalytic dosage, pH, dye concentration and irradiation time to determine the optimum condition for the degradation of RO-M2R. The optimized degradation of RO-M2R was achieved with 83.41% by Box-Behnken design. Therefore, CuO NP's could be utilized as an efficient photocatalyst for the degradation of dye-containing waste water from textile industries.

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Chapter 13

Vermicomposting—An Effective Method for Sustainable Agriculture and Environmental Impact



R. Vinoth Kumar, Farheen Fathima, R. Jaishree, and G. Priya

Abstract Societal waste materials from the landfill are considered as one of the major contributors affecting the ecosystem and environment. Composting and vermicomposting are the commonly used methodologies to treat and convert these waste materials into the useful manure by the collective action of earthworms and other beneficial microbes. Waste management is considered as an integral part of a sustainable society, thereby necessitating the conversion of these biowastes into alternative management processes such as vermicomposting. Vermicompost produced by the activity of earthworms is rich in nutritive organic macro and micronutrients, vitamins, growth hormones, beneficial soil microbes and enzymes such as proteases, amylases, lipase, cellulase and chitinase etc. Such microbes and enzymes continue to disintegrate organic matter even after they have been excreted from the earthworms. In this review, we have summarized the importance of vermicomposting, effect of process-enhancing supplements, the detoxification process of industrial wastes by earthworms and the role of vermicompost in plant growth and development. Further, the recent advancements in innovative vermicomposting technologies employed for small- and large-scale production are highlighted. Collectively, it highlights vermicompost as a powerful crop nutrient in sustainable agriculture.

Keywords Biowaste · Beneficial microbes · Composting · Earthworms · Plant growth and development · Process-enhancing supplements · Technological innovations · Waste management

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13.1 Introduction

Urbanization and advances in industrial processes have increased the amount of garbage produced globally. To mitigate pollution originating from the garbage, developed nations have implemented practical measures like limiting waste production at the source, increasing recycling, and reuse. Globally, at least 2 billion tons of non-hazardous waste materials are generated each year (Chalmin and Gaillochet 2009). Particularly, in low- and middle-income nations, bio-waste is the major source of municipal and industrial wastes (Kaza et al. 2018). Circular economy has gained recognition in current scientific literature and among international and national authorities as a key strategic environmental policy framework (Jain et al. 2022). The most popular methods used for treating and converting these waste materials into valuable end-products are composting and vermicomposting (Adhikary 2012; Zhou et al. 2022a, b). Vermicompost research has had impressive positive growth trends in the previous 25 years with >2000 papers published connected to vermicompost and with more nations participating in this subject globally (Ghorbani and Sabour 2021). This study emphasizes on vermicomposting as an effective method for attaining sustainable agriculture.

13.2 What Is Vermicomposting?

Vermicomposting is a bio-oxidation and stabilisation method that uses the earthworms' biological activity to break down organic waste into a nutrient-rich organic biofertilizer. Sewage sludge and solids from wastewater, brewery materials, paper trash, urban leftovers, food and animal wastes, as well as horticultural residues from processed potatoes, dead plants, and the mushroom business, have all been treated successfully using this method (Garg and Gupta 2009; Lim et al. 2015; Datta et al. 2016; Bhat et al. 2018). Earthworm activity results in the production of vermicompost, which is rich in macro- and micronutrients, vitamins, growth hormones, enzymes including lipase, cellulase, and chitinase, as well as beneficial microflora (Sanchez-Hernandez 2019; Singh et al. 2020).

In the vermicomposting process, the microbial community's composition is greatly changed by the earthworm gut transit (Romero-Tepal et al. 2014). The earthworms aerate, fragment the substrate there by increasing the surface area for microbes and modify the microbial activity in organic waste residues for decomposition (Lavelle et al. 2006). The stomach of earthworms secrete a variety of enzymes, including lipases, proteases, cellulases, amylases, and chitinases, to convert the inaccessible form of minerals, such as nitrogen, phosphorus, potassium, and calcium, etc., (in organic waste materials) into forms that are available to plants (Hand 1988). Due to the faster rate of mineralization and humification by earthworms during vermicomposting than during thermal composting, the finished product (vermicompost) has a higher nutritional value (Albanella et al. 1988). In the process of vermicomposting,

earthworms excrete coelomic fluids, which kill the parasites and bacteria present in the waste and produce pathogen- and odour-free end-product, vermicompost (Kale and Krishnamoorthy 1981).

13.3 Classification of Vermicomposting

Vermicomposting can be broadly divided into two categories: primary vermicomposting and secondary vermicomposting (Garg and Gupta 2009). Directly feeding raw materials to the worms is the primary method of vermicomposting. Because all of the nutrients in organic matter are used to raise worms, a primary system has a high stocking density (number of worms per surface area). The drawback is that worms need to be fed frequently and in small amounts.

Prior to being fed to worms, raw materials are composted by fungus and microorganisms in secondary vermicomposting. In piles of secondary vermicompost, auto-heating doesn't happen. Because the organic matter has already undergone some digestion before the worms can consume it, secondary vermicompost systems have relatively high feeding rates (mass of organic matter fed per surface). This implies that secondary bins are supplied in thick layers less frequently.

13.4 Characteristics of Vermicompost

Vermicompost, which is comprised of compost that has been digested, is a great soil enhancer (Garg and Gupta 2009). Worm castings are regarded as a higher value product since they contain significantly more nutrients and microbial life (Table 13.1). Vermicompost includes nutrients that are necessary for plants and other microorganisms, as well as a variety of organic or inorganic minerals like oxygen, nitrogen, and phosphorus. They also have very little heavy metal content. Vermicompost contains a 3:1:2 ratio of NPK in its NPK makeup (Garg and Gupta 2009; Adhikary 2012). This makes it as an excellent nitrogen fertiliser for plants, particularly those with green leaves and those that develop quickly. Vermicompost can also be applied to soil that lacks organic matter to improve it. It includes 1.5 to 2.2% nitrogen, 1.8 to 2.2% phosphorus and 1.0 to 1.5% potassium on an average. There are micronutrients such as sodium, calcium, zinc, sulphur, magnesium and iron in the organic carbon, which ranges in concentration from 9.15 to 17.98 (Garg and Gupta 2009; Olle 2019). The physical characteristics of the soil may also be significantly impacted by worm compost. It was demonstrated that adding 20 tha of vermicompost over the course of two years greatly increased soil porosity and aggregate stability (Table 13.2).

Table 13.1 Population of different microorganisms present in vermicompost (Reproduced from Pramanik et al. 2007)

Treatments	Total bacterial count (X10 ⁸)		Cellulolytic fungal count (X10 ⁶)		Nitrifier population (X10 ⁴)	
	L ₀	L ₁	L ₀	L ₁	L ₀	L ₁
<i>Cow dung</i>						
M ₀	29	32	50	52	4	3
M ₁	20	169	72	55	10	13
M ₂	16	110	59	45	3	28
M ₃	49	49	76	59	49	5
<i>Grass</i>						
M ₀	17	19	34	35	2	3
M ₁	29	31	44	76	12	11
M ₂	25	38	41	52	7	15
M ₃	35	30	51	38	13	9
<i>Aquatic weeds</i>						
M ₀	13	18	41	43	4	4
M ₁	27	33	47	49	9	10
M ₂	29	37	43	63	8	12
M ₃	35	28	51	48	11	9
<i>MSW</i>						
M ₀	9	10	19	17	0.7	1
M ₁	17	19	23	20	1.5	2.1
M ₂	15	23	21	22	1.7	2.9
M ₃	20	17	25	21	2	2.3

L₁ is lime at 5 g/kg and L₀ is control; M₀, M₁, M₂, and M₃ are control and incubation of *Trichoderma viridae*, *Bacillus polymxa* and *Phenerocrete crysosporium*, respectively at 50 ml/kg

Table 13.2 Optimal conditions required for breeding *E. fetida* and *E. andrei* in organic wastes

Parameters	Optimal conditions required
Temperature	15–20 °C
pH	5–9
Moisture content	60–80%
Ammonium content of the waste	<1 mg/g
Salt content	<0.5%
Oxygen	Aerobic

13.5 Sources of Organic Wastes for Vermicomposition

Organic wastes are substances that come from living beings such as plants, animals, and microorganisms and are biodegradable—that is, they can be converted into more basic organic molecules. Depending on the amount of moisture present, they can be found in either a solid or a liquid state in nature (Ansari et al. 2020). Vermicompost is made from a variety of organic waste sources, including both small- and large-scale production. The following are the common sources:

Municipal solid trash, which includes food leftovers, recycled newspaper, and grass clippings. Animal waste includes everything from egg shells to vegetable skin to animal dung.

Plant waste, which includes rotting and dead plants, leaves, flowers, etc.

Food waste includes ruined food goods, fruit peels, and vegetable skin.

13.6 Role of Earthworms in Vermicomposting

Earthworms are nocturnal, terrestrial invertebrates in nature. Globally, there are about 3000 species of earthworms belonging to 9 families, of which 450 species/subspecies have been found in India (Thakur and Yadav 2018). Indian blue worms, *Metaphire posthuman*, and *Lampito mauritii* are a few examples of common Indian earthworms. Except for few species like *Pontodrilus burmudensis*, which thrives in estuary water, the majority of earthworm species are found in soil. These worms eat organic soil surface matter, including bacteria, fungi, dead plants, living protozoa, rotifers, nematodes, and other microbes. The organic materials that earthworms consume are transformed into nutrient-rich inorganic and organic compounds, which are then excreted (Singh et al. 2020).

The commonly used earthworms for the process of vermicomposting are *Eisenia fetida*, *Eisenia hortensis* and *Perionyx excavates*. Red wigglers, or *Eisenia fetida*, are ideal for vermicomposting because of their quick digestion and efficient breakdown. They may be readily maintained, and require moisture and shade to grow. Compared to vegetable and mixed food waste, *Eisenia fetida* favoured tea waste. Vermicomposting was more productive by nearly 40% in tea trash (Garg et al. 2006; Yadav and Garg 2011; Mohapatra et al. 2019). The European nightcrawler, *Eisenia hortensis*, is larger than other earthworms and is generally grey or brown in colour. They are prevalent in heavily forested areas. They can withstand a broad range of temperatures and reproduce quickly. They are hence recommended in equatorial areas.

Native to sub-Saharan Africa, *Perionyx excavatus* is a species of nightcrawler. They efficiently aerate and decompose organic materials. Because they can withstand a wide variety of temperatures and moisture levels, they are ideal for vermicomposting. Additionally, they serve as fish bait. *Perionyx excavatus*, sometimes known as Indian or Malaysian Blues Worms, is a native species to Southern Asia.

They effectively break down organic debris, and are simple to cultivate (Martin and Eudoxie 2018; Ananthavalli et al. 2019).

13.7 Basic Environmental Requirements for Earthworms

The earthworms in the vermicomposting process require the following environmental conditions—temperature, moisture, acidity and alkalinity, aeration, salts and carbon:nitrogen ratio (Peigne and Girardin 2004; Singh et al. 2016) (Table 13.3).

Temperature: Red wigglers grow at their fastest between 27 and 35 °C. Composting worms can withstand temperatures above 85 F for brief periods of time, but they perish if exposed to extreme heat for an extended amount of time. Lower temperatures (below 65F) are more tolerable to them (Liang et al. 2003; Amaravathi and Reddy 2015). As long as the pile's core temperature stays above freezing, composting worms will survive. The worm eggs known as cocoons can endure extreme cold for long periods of time before hatching.

Moisture: Water is a must for worm life at all costs. The ideal range for moisture content for growth is between 70 and 85%. This is a little moister than microbial composting piles, which typically have a moisture content of between 50 and 60% (Liang et al. 2003; Singh et al. 2004). Red wigglers can briefly survive in entirely saturated bins, but if the oxygen supply starts to diminish, they will leave.

Carbon to Nitrogen Ratio (C:N): Compost worms digest just about any type of organic matter, but they prefer higher amounts of carbon than is normal for microbial composting (Ndegwa and Thompson 2000). Organic wastes such as manure, kitchen wastes, coffee grounds, and freshly cut grass contain lots of Nitrogen (N). You need to dilute these materials with byproducts containing high amounts of Carbon (C), such as straw, paper or fallen leaves. The ratio of carbon to nitrogen in worm food is called the Carbon to Nitrogen Ratio (C:N). Worms thrive when their food has C:N greater than 50. The main reason for this is microorganisms outcompete worms at lower C:N ratios. At best, microorganisms take food from the worms' mouth. At worst, they cause the pile to heat, which kills the worms.

Acidity and alkalinity: The pH of soil serves as a gauge for its relative acidity or alkalinity. The ideal pH for composting worm growth and reproduction is 7 (Amaravathi and Reddy 2015; Singh et al. 2016). Their survival rate reduces with pH values of 5 or above. Worms may withstand somewhat greater pH ranges than other less desirable compost pile visitors, such as flies. Fungus development is also aided by a lower pH. When the pH is low, worms will flee, but once it is close to neutral, they will return to eat the fungi.

Aeration: Earthworms lack specific respiratory organs and instead breathe via their body walls. As a result, earthworms are extremely sensitive to anaerobic environments. High concentrations of *E. fetida* have been observed migrating from water-saturated substrates where oxygen has been reduced or where carbon dioxide or hydrogen sulphide have accumulated (Yadav and Garg 2011).

Table 13.3 Role of vermicompost for sustainable agriculture

Crops	Beneficial activities on plants
Banana, cassava and cowpea	Improved in yield and biometric characteristics
Brinjal	Increased germination, number of fruits per plant, mean fruit weight, yield per plant
Capsicum	Improved plant nutrition, growth, photosynthesis and chlorophyll content of the leaves Vermiwash significantly affected the growth and nutrient utilization
Carrot	Increased root length, root volume, root weight and root yield
Cowpea	Increased seed yield, straw yield, biological yield, total root nodules and leghaemoglobin content, protein content in seeds
Chickpea	Increased plant height, plant shoot biomass, number of pods and photosynthetic pigments of chickpea plants
Hyacinth bean	Increased growth and yield characteristics
Marigold	Growth augmentation, increased overall nitrate– nitrogen concentrations in leaf tissues at flowering stage, improved soil fertility Increased bud formation, number of flowers per plant, plant shoot biomass, plant height and increased diameter of the flower
Mulberry	Increased luxuriant growth of bacteria treated rhizosphere and other micronutrients
Muskmelon	Increased germination rate, plant height, leaf area and plant biomass
Okra	Increased pod yield and soil fertility Reduced fruit borer infection Induced resistance in plants against pests and disease
Peppermint	Increased levels of chlorophyll a, chlorophyll b, total chlorophyll and carotenoids
Paddy	Increased growth, yield, nutrient uptake and soil characteristics
Soybean	Increased seed germination, seedling survival, shoot length and root length
Spinach	Increased plant height
Strawberry	Increased photosynthesis rate, free radical scavenging and soil enzymatic activity
Tea	Increased radical scavenging activity in leaf extracts
Tomato	Improved quality and productivity of tomato Protective agent for improvement of salt stress resistance in tomato Accumulation of plant growth hormones and carbohydrates Increased juiciness, titratable acidity, ascorbic acid content, shelf life Decreased number of nematode-induced galls on tomato plants

Ammonia and salts: Earthworms cannot survive in organic wastes having significant concentrations of cations because they are extremely sensitive to ammonia and salts (Tripathi and Bhardwaj 2004; Wu et al. 2019). Additionally, they die in wastes that include a lot of inorganic salts. Ammonia and inorganic salts both have extremely distinct hazardous and benign ranges (1 mg/g of ammonia and 0.5% salts, respectively). However, if the ammonia has been removed through pre-composting or water leaching, organic wastes with high ammonia levels may become acceptable. Earthworm activity and rates of organic waste processing drastically drop outside of these environmental parameters, hence wastes should be preconditioned to make them acceptable for earthworms in order to maximise vermicomposting effectiveness.

13.8 Vermicomposting Process

The vermicomposting process consists of two phases based on the biological and physiochemical mechanisms in the earthworms and microorganisms (Fig. 13.1):

Active phase: The earthworm interacts with by-products to change their physical and chemical properties as well as their microbial make-up.

Maturation phase: The earthworm moves to a new layer of new substrate, creating space for the emergence of bacteria that break down the biological matter that the earthworm has previously processed.

13.9 Impact of Additives, Bulking Agents and Microbes

Commercial composting supplements are frequently marketed as “starters,” “activators,” or “accelerators,” but they strongly differ in composition and typically contain a combination of chemicals, bulking agents, and microbes. In order to speed up the composting process and enhance the leaching, greenhouse gas emissions, and odour of the composting process, additives are frequently applied (Awasthi et al. 2020b). The use of such chemicals may not only affect important process variables but also raise the final product’s agronomic value (Barthod et al. 2018). Increased pH, improved nitrogen retention (both total and nitrate nitrogen), and faster seed germination have all resulted with the addition of 10–15% biochar made from straw, but the bioavailability of heavy metals has decreased (Awasthi et al. 2020a; Zhou et al. 2021; Zhou et al. 2022a). Additionally, in a field vermicomposting experiment,

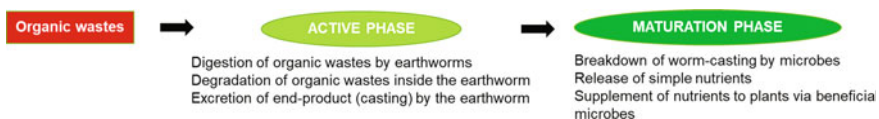


Fig. 13.1 The different phases of vermicomposting by earthworms

biochar was used in conjunction with earthworms to produce vermicompost, which resulted in decreased emissions of ammonia, methane and nitrous oxide (Raza et al. 2022). Similarly, a modest amount of lime (1%), combined with 30% zeolite, may improve the composting process by lowering greenhouse gas emissions and limiting ammonia loss (Awasthi et al. 2016).

Bulking agents are substances that are added to the compost to change its physical composition, including its ability to absorb water, the amount of airflow and space between particles, and the C:N ratio. In order to optimise open air space, C:N ratio, pH, and adjust moisture holding qualities of compost, dry fibrous and carbon-rich materials (such as lignocellulose) are utilised (Sarda et al. 2019). When compared to other comparable bulking agents, sawdust has a stronger potential to minimise leaching and is an efficient regulator of free air space and moisture content (Rich et al. 2018).

Given that peak temperatures during the thermophilic phase can reach up to 70 °C, tolerance to high temperatures is a crucial need for microorganisms that are intended to remain active throughout the composting process. The inoculation of refractory lignocellulose with thermo-tolerant *Actinomycetes* strains such as *Streptomyces* sp. H1, *Mycobacterium* sp. G1, *Micromonospora* sp. G7, and *Saccharomonospora* sp. T9 has demonstrated encouraging results (Wei et al. 2019; Muhammad et al. 2021). The humification process in composted cattle dung was shown to be improved by the addition of thermo-tolerant ammonia oxidising bacteria (Xu et al. 2022).

13.9.1 Effect of Vermicompost on Plant Growth and Development

As macronutrients, micronutrients, and other organic substances found in vermicompost are abundant, they are highly significant for the growth and development of plants (Ceritolu et al. 2018; Annapoorani and Sindhu 2021). Additionally, vermicompost contains enzymes and hormones that promote plant growth (Abbasi and Ramasamy 1999; Hussain et al. 2017). Numerous studies have indicated that bacteria from vermicompost produce plant growth hormones (auxin and cytokinins) (Muscolo et al. 1999; Atiyeh et al. 2002). Many researchers have demonstrated that the final vermicompost made from cow dung, sewage, and paper mill sludge contains significant amounts of humic compounds, which are important for plant growth and development (Senesi et al. 1992; Garcia et al. 1995; Masciandaro et al. 1997; Elvira et al. 1998). The different plant species that have experienced a significant increase in growth include banana, cassava and cowpea (Padmavathiamma et al. 2008), carrot (Chatterjee et al. 2014), Capsicum (Rekha et al. 2018; Khan et al. 2014), cowpea (Khan et al. 2015), chickpea (Yadav and Gard 2015), hyacinth bean (Karmegam and Daniel 2008), mulberry (Mary et al. 2015), muskmelon (Manh and Wang 2014), okra (Oroka 2015; Hussain et al. 2017), peppermint (Ayyobi et al. 2014), paddy

(Jayakumar et al. 2011), tomato (Joshi and Vig 2010; Truong et al. 2018; Benazouk et al. 2018; Rajya Laxmi et al. 2015; Xiao et al. 2016), soybean (Sheikh et al. 2015), spinach (Peyvast et al. 2008), brinjal (Najar et al. 2015), marigold (Atiyeh et al. 2001; Sangwan et al. 2010), strawberry (Singh et al. 2008; Zuo et al. 2018), groundnut (Mycin et al. 2010), french bean (Hussain et al. 2017) and tea (Bagchi et al. 2015).

13.9.2 Advances in Vermicomposting Technology

In order to obtain high-quality natural fertilisers for sustainable agriculture practises and to improve the role of composting and vermicomposting practises throughout the transition to a circular economy, technology development based on research and innovation activities will be essential (Awasthi et al. 2022). China has the highest percentage of global patents (44%) and that novel composting devices are frequently the subject of compost technology-related inventions (Zheng et al. 2020). Comparing reactor composting to other conventional methods, such as static heaps, windrow composting, and membrane-covered composting, an LCA research found that reactor composting is a potential answer in terms of eco-efficiency (Chen et al. 2020; Liu et al. 2022). Further, pre-degrading organic waste before vermicomposting has resulted in reduced vermicomposting time period of about 2 weeks than traditional vermicomposting methods (Kausar and Khwairakpam 2022) (Fig. 13.2).

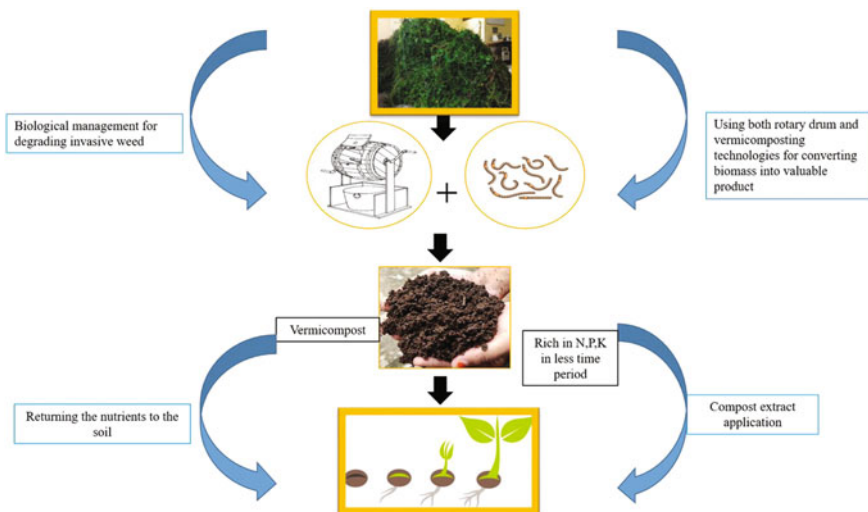


Fig. 13.2 Two-step vermicomposting process for organic waste management (Reproduced from Kausar and Khwairakpam 2022)

13.9.3 Small Scale Vermicomposting with Newer Technology

A complementing strategy to raise citizen responsibility and environmental awareness of bio-waste management and composting difficulties might be the community's involvement in constructing practical composting containers (Lunag et al. 2021). Improved composting toilet design may be a solution to the area's sanitation problems; however, this solution is only appropriate in select "green zone" locations where composting processes are likely to be successful (Lopez-Zavala 2019; Zhou et al. 2022a, b). To fasten the transition to a circular economy, on-site vermicomposting must be integrated into a comprehensive strategy. For instance, extra heat energy produced during the thermophilic phase of on-site composting at the home level could be captured and used to heat a greenhouse in the fall. In keeping with the objectives of the circular economy, Neugebauer et al. (2021) employed artificial neural networks to forecast the temperature inside the greenhouse to demonstrate that the vermicomposting process not only decreases bio-waste but also offers a chance for heat recovery. Future studies on small-scale composting techniques should really support such integrative methods (Zhou et al. 2022a, b).

13.9.4 New Technologies for Large-Scale Composting

When additional precautions like insulating methods, additives, and substrate pre-treatment are taken into account, in-vessel vermicomposting technologies are viable for colder climates for large-scale production (Kumari et al. 2022). The fact that earthworm extraction is frequently done manually is one of the major issues with large-scale vermicomposting methods. According to a novel technology proposed by Walling et al. (2020), earthworms can be sorted using centrifugation with a worm recovery rate of roughly 84%. A combined strategy, such as vermicomposting with room drying for 10 days, is suggested to improve the stabilisation process of dewatered sewage sludge (Huang et al. 2022). Another technical option to improve operating parameters and hasten the biodegradation process is the vermicomposting reactor (Ramprasad and Alekhya 2021). When compared to a conventional container, a smart vermicompost reactor generated compost in half the time and with a 30% faster worm development rate (Ghorbani and Sabour 2021). The development of mathematical models and sensors to optimise a variety of composting/vermicomposting parameters, including artificial intelligence, is anticipated to be the subject of new study.

13.10 Summary

Without proper treatment, industrial pollutants and sludges may pollute the soil and other wildlife, posing serious health risks. Utilizing earthworms as adaptable natural bioreactors, vermicomposting technology allows for effective nutrient recycling and improved plant development while also decomposing organic matter and maintaining soil fertility. Vermicomposting can be used for a variety of organic solid wastes, including those that are domestic, animal, agro-industrial, human, etc. The value of vermicompost is further increased by the fact that it also provides other advantages. Extra worms can be utilised as a soil cleaner and as protein-rich animal feed, providing they are not grown on polluted wastes. The majority of research also showed that the vermicompost's end product may function as an appropriate medium for plant growth since it has a higher concentration of soil enzymes and growth hormones.

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Chapter 14

Remediation of Endocrine Disrupting Compounds and Organic Dye Pollutants Through Biosorbents in a Circular Bioeconomy: Prospects and Constraints



Annu T. Mathew and M. P. Saravanakumar

Abstract The presence of endocrine-disrupting compounds (EDCs) and organic dye pollutants in water may have toxic, hazardous, and carcinogenic effects on exposure. The recent advances in research on biomaterial synthesis, characterization, and applications have attracted a lot of researchers in this area. The physical, chemical, and thermal characteristics of green synthesized materials help to facilitate their interactions with a variety of water pollutants. The use of green synthesized nano-materials as eco-friendly sorbents increases the value of wastes and naturally available compounds utilized during its synthesis. The major advantage of the adsorption process is that it does not involve the formation of additional toxic by-products, unlike other treatments. Several operational parameters like pH, nanomaterial dose, reaction time, pollutant concentration, other contaminants, and temperature may influence the adsorption process. The adsorptive interactions between bio-based material and water pollutants may be through electrostatic, covalent (H-bond and Van der Waals forces), π - π , and n- π bonds. Many isotherm and kinetic models are used to validate the experimental results for adsorption. The simplicity, eco-friendly, and cost-effective nature of pollutant adsorption through biosorbents provides scope for its utility on a large scale. The circular bioeconomy motivates the *close the loop* concept through the generation of the least waste, utilization of created wastes, and maximum utility of naturally available substances. The prospects regarding the application of biosorbents for the elimination of EDCs and organic dyes in a circular bioeconomy are explained and constraints concerning it are discussed in this work.

Keywords Green synthesized materials · Endocrine-disrupting compounds · Organic dyes · Adsorption · Circular bioeconomy

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14.1 Introduction

EDCs are emerging organic contaminants whose consumption is considered to increase over a while in the future. This is due to its wide range of applications, increasing production, and consumption with the improving standards of living. Micropollutants are a class of compounds that may include pharmaceutical compounds, EDCs, pesticides, personal care products, and biocides. The entry of these compounds into an aquatic ecosystem is through industrial effluents, sewage effluents, animal wastes, landfill leachate etc. Even traces (ng/L- μ g/L) of these compounds inside an aquatic ecosystem can prove hazardous (Goswami et al. 2018). Therefore, there is a high risk of exposure and a great need for a detailed study of these toxicants. The existence of these compounds in groundwater, seawater, surface water, and effluents has been detected in various concentrations (Olasupo and Suah 2021). Hence, it becomes important to remove these micropollutants before it transforms into other complex products under the diverse environmental conditions.

In the meantime, dyes can be broadly divided into categories such as acidic, disperse, basic, azo, diazo, and a metal complex. About 20% of industrial wastewater consists of dye pollutants in it. The organic dye release in the environment is a major concern due to potential mutagenic, carcinogenic, and toxic effects (El-Sharkawy et al. 2007). The organic dyes possess chromophores (aromatic rings and azo groups) which make the structure quite stable. Hence, these chemicals become difficult to degrade under natural light, heat, and oxidants in the environment (Qin et al. 2022). Therefore, it is necessary to eliminate dyes from an aquatic environment. Biosynthesis may include the use of plant extracts or sludge or wastes for the generation of nanoparticles (Singh et al. 2018). The extracts of plants or naturally available solutions act as natural reducing and capping agents during nanoparticle synthesis (Jadoun et al. 2021).

The circular water economy aims at providing sustainable production and consumption of water through water reuse and resource recovery. Bioeconomy acts as a catalyst for the transition from a linear economy to a circular one (Ferreira et al. 2018). The linear economy maintains a linear relationship between input and output. It focuses on converting raw materials to final products, distribution, and consumption of these final goods, and disposal of wastes. Bioeconomy is also called circular green economy. An effective bioeconomy must be economical, circular, sustainable, and ecological. The basic strategies towards such a transition involve the least production of waste, and the reuse of the products in a close the loop manner (Aguilar and Twardowski 2022).

14.2 Sources and Effects of EDCs and Organic Dyes in Water

Organic pollutants are released due to anthropogenic activities from domestic, industrial, pharmaceutical, and other sectors. The maximum quantity of various chemicals existing in wastewater treatment plant (WWTP) effluent is mostly present in a range of $1-10^3$ $\mu\text{g/L}$ (Rogowska et al. 2020). The report produced by the Department of Water Affairs for wastewater quality management (2012 Green Drop Report) and potable water quality management (2012 Blue Drop Report) indicates that regulatory limits are not met by many municipalities in the North West Province of South Africa. Therefore, this water is not proper for consumption without appropriate treatment (Wanda et al. 2018).

Carpenter et al. screened for about 200 micropollutants collected from 17 sites along the Hudson River estuary. Hudson river serves as a drinking water source for more than 100,000 people and is also used for recreational and commercial purposes. The researchers identified clusters and sub-clusters of micropollutants based on normalized concentration patterns and spatiotemporal occurrence. The major clusters are named 'core micropollutants', 'sourced from diffuse upstream sources', and 'those having sewage treatment plants as sources'. Totally 127 samples are analyzed. A list of 200 target micropollutants is used to detect 168 in at least one of 127 samples. Out of which 116 are derived from wastewater and 52 are sourced from agricultural activities. The concentration of a majority of micropollutants has been found in the range of 1 to 100 ng/L. Few compounds such as atenolol acid, metformin, and sucralose are found in high concentrations of about several mg/L and are from sewage treatment plant outfall samples (Carpenter and Helbling 2018).

The limits of micropollutants present in freshwater have also been exceeded in many other parts of the world. It was reported that high concentrations of acetaminophen, valsartan, codeine, gabapentin, tramadol, etc. have been detected in River Taff and River Ely of Wales, UK. In southern Spain, pharmaceutical concentrations have been detected from compost, sewage sludge, and sediment samples along the Guadiamar River (Ebele et al. 2017). Considering, developing countries like India which show the presence of pesticidal micropollutants in influents of WWTP indicates, that it is an agriculture-based country. The values of pesticide concentration above safety limits can prove to be life-threatening. Similarly, the presence of caffeine in raw sewage of WWTP in China infers great consumption of coffee and tea in the area (Zhou et al. 2010). Therefore, the type of micropollutant is greatly influenced by habitual nature and other associated factors of that particular place. Hence, due to the hazardous nature and increasing risks posed by more than thousands of materials, watch lists have been created for inter-government monitoring. The list includes several priority substances or a group of materials. Some of the compounds like bisphenol A (BPA) and diclofenac are part of the list as they are extremely toxic (Barbosa et al. 2016).

Meanwhile, dyes find broad applications in leather tanning, food processing, cosmetics, paper and printing, and dye manufacturing units. The dyes are present in

industrial wastes and eventually get discharged into water bodies. The total release of dyes from various processes is still unknown but studies suggest that 100 tonnes of dyes are received by water streams per year (Yagub et al. 2012). The processing of fabric requires about 100–200 L of water per kilogram (Holkar et al. 2016). The regulations for textile wastewater emission depend on the product type generated by the factory. Various countries like Malaysia, India, Algeria, etc. are found to release textile wastewater with different physical and chemical properties. The parameters like pH, biochemical oxygen demand (BOD), chemical oxygen demand (COD), total dissolved solids (TDS), total suspended solids (TSS), turbidity, and color of textile wastewater vary based on the factory processes it undergoes (Dihom et al. 2022). The dyes make the water bodies colored, and their toxicity, structure, and non-degradable nature (sometimes), make even their exposure at small concentrations risky to living organisms (Yagub et al. 2014). Therefore, these water contaminants need to be removed from water and wastewater systems.

Human beings can encounter micropollutants through activities like drinking, bathing, swimming, and recreation in surface water. Sometimes, these toxic compounds get into groundwater, and a lack of proper treatment specific to micropollutant elimination can cause severe health issues (Eggen et al. 2014). Though sometimes they are invisible to the naked eye, even small traces of micropollutants in water are dangerous. From toxicological studies, the presence of micropollutants in drinking water is linked to potential carcinogenic risks which exceed the EPA's negligible risk level (Xue et al. 2021). Additionally, researchers suggest that intermediates generated when a pollutant gets exposed to some organisms or chemicals may be of higher toxicity than parent compounds (Žur et al. 2018).

The presence of diclofenac in quantities less than 1 $\mu\text{g/L}$ can have ill effects on the kidney, liver, and gills of Pacific rainbow trout (*Oncorhynchus mykiss*) (Mehinto et al. 2010). Similarly, chlorpyrifos micropollutants pose a severe threat to living organisms by causing infertility, skin disorders, cancer, etc. (ul Haq et al. 2020). It is reported that parabens induce immunologically mediated systemic hypersensitivity reactions and inhibit estrogenic activity. Also, exposure to N,N-diethyl-meta-toluamide inhibited the enzyme controlling the central nervous system (Keerthanam et al. 2020).

Studies also suggest that the presence of a detrimental type of amines in azo dyes makes them highly toxic (Robinson et al. 2001). The highly soluble and chemically stable nature of dyes makes the wastewater pigmented, posing a severe threat to the bio-system (Asghar and Bhatti 2012). Exposure to dyes can cause allergic skin disorders, in textile workers and people using close-skin fitting clothes made of synthetic fibers. The toxic impacts of prolonged dye contact include chromosomal fractures, respiratory toxicity, carcinogenesis, and mutagenesis (Yagub et al. 2014). The use of dyes like malachite green is banned in many countries due to their highly toxic impact on the aquatic system, yet its illegal utilization is still widespread in some countries (Pathy et al. 2022). Dyes like rhodamine B which are extensively used in the dyeing of paints and fabrics are also reported to endanger the earth due to their hard-to-degrade and high chromatic nature (Xu and Ma 2021). Hence, there is a need

for the removal of these toxic organic compounds from the water and wastewater system.

14.3 Characterization and Use of Biomaterials for Organic Pollutant Removal

The unique properties of green synthesized nanoparticles can be determined by utilizing several tools for characterization. The green synthesized nanoparticles exist in different shapes like spheres, rods, cubes, etc., similar to the nanomaterials obtained using non-green ways. A few examples of morphologies obtained using the green synthesis pathway are mentioned below. The existence of rod-shaped nanoparticles of size 100–150 nm is obtained for hydroxyapatite, green synthesized using sea biowastes (Karvandian et al. 2020). Licorice root extract is utilized to produce hydroxyapatite nanorods in other studies. The aggregated rods appear uniform, similar-sized, and well-crystallized (Ali et al. 2021). In another research, the celery seeds extract acts as a green reducing agent for generating nanocomposites. Silver nanoparticles on the surface of reduced graphene oxide nanosheets are obtained through this process. The TEM images obtained reveal monodispersed, spherical, and fine silver nanoparticles (size 20 nm approx.) on the surface of reduced graphene oxide after microfluidization treatment. The results propose that celery extracts enhance the dispersibility of silver nanoparticles on wrinkled sheets of reduced graphene oxide (Soleymani et al. 2020). Uniform and porous microsphere structures loosely packed by nanocubes are obtained through coconut shells in another study. This structure enhances the active sites and sensitivity (Sivasakthi et al. 2020). Researchers suggest the biosynthesis of Cu₂O nanospheres decorated on graphitic carbonitride using citrus limon leaf extracts through the hydrothermal method. The average particle size of the Cu₂O nanosphere is found to be in the range of 2–10 nm (Induja et al. 2019). Therefore, different morphologies of nanoparticles can be attained using green synthesis methodologies. The derived nanoparticles also possess excellent properties for various field-scale applications.

The surface area of green synthesized nanoparticles is determined using BET analysis. Results indicate that the synthesis of nanotubes using wastes or natural feedstock like coconut oil, rice straw, camphor, palm oil, etc., produces carbon nanotubes of excellent surface area and dimensions (Makgabutlane et al. 2021). For instance- the carbon nanotubes green synthesized using rice straw yield a surface area of 188 m²/g and a size of 15–40 nm (Lotfy et al. 2018). In another research, the surface area and pore volume of green synthesized iron nanoparticles (using eucalyptus leaves) are compared with raw laterite particles. The surface area and pore volume are higher for iron nanoparticles (36.62m²/g and 0.0394cm³/g respectively) than for raw laterite particles (23.18m²/g and 0.0091cm³/g, respectively). This can be attributed to polyphenols in the eucalyptus that act as capping and reducing agents (Sangami and Manu 2017). The surface area of α -Fe₂O₃ synthesized using green tea leaf extract

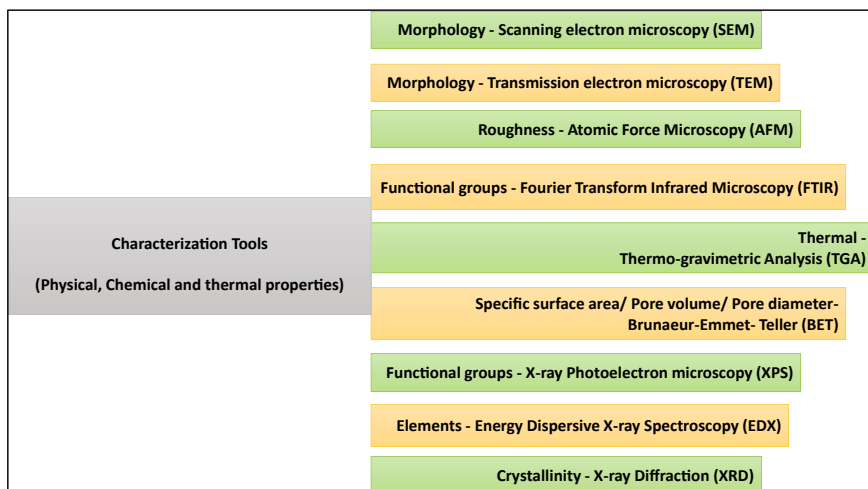


Fig. 14.1 Tools to characterize biosorbents to determine physicochemical properties

through the hydrothermal method is observed to be four times more than traditional commercially available ones. The reactivity is also doubled for green synthesized α -Fe₂O₃ when compared to synthetic nanoparticles (Ahmmad et al. 2013). Therefore, green nanoparticle synthesis helps achieve high surface area and pore volume beneficial for wastewater treatment applications. Hence, the physicochemical properties of bio-based materials found through various tools help to determine the nature and stability of biosynthesized material. The various instrumentation techniques and characterizations that may be used to characterize biomaterial and their origin are presented in Fig. 14.1.

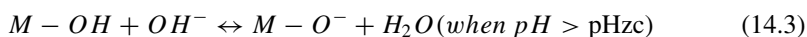
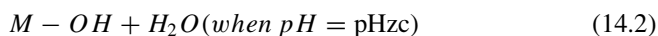
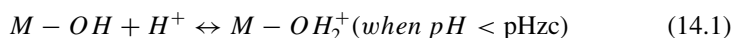
14.4 Adsorption of EDCs and Dyes Using Bio-based Materials

Adsorption is a process employed by most researchers due to its simple operation, high efficiency, and flexibility of design (Crini 2006). The process does not involve the formation of toxic intermediates or by-products unlike other treatments (for example: in the degradation of pollutants in an advanced oxidation process) (Mathew and Saravanakumar 2022a). There are two essential elements in an adsorption process. Firstly, the adsorbent on which the pollutant goes and attaches. Secondly, the adsorbate which is the pollutant itself. Here, the adsorbent is biomaterials and the adsorbate are the EDCs and dyes.

Several factors impact the process efficiency based on the nature of bonds formed between biosorbent and organic pollutants. The presence of other contaminants mostly hinders the removal of EDCs/ dyes in a real water system (Wu et al. 2022).

Still, biosorbents which can perform well in the presence of other pollutants, have been synthesized. The improved performance can be mostly due to better adsorbent properties obtained in the system (Yu et al. 2018). The solution pH, temperature, reaction time, biosorbent dose, and contaminant concentration are a few factors that affect the efficiency of sorbent. The pH of the solution can affect the nature of bonds between biosorbents and pollutants. This interaction is based on multiple factors like the point of zero charge (pH_{zc}) of the adsorbent and the isoelectric point of the pollutant. The biosorbent dose determines the amount of active sites available to pollutant and an increase in the dosage enhances removal till saturation. The pollutant concentration in the solution is a measure of the quantity of a pollutant that needs to be treated by the fixed dose of biosorbent, and this impacts process efficiency. The temperature variation may affect the viscosity of the solution or weaken the bonds for effective adsorption (Mathew and Saravanakumar 2021).

The nature of bonds formed in the system can be analyzed through a deep study of the adsorption mechanism. The types of functional groups on the pollutant and sorbent, with other operational parameters, play a vital role to determine the type of bond in the system. Hence, physisorption or chemisorption can occur in the system. The adsorptive bonds may include electrostatic interaction, π - π bonds, n - π , complexation, H-bonds, hydrophobic interactions, pore filling, etc. Weak physical bonds are formed in the case of physisorption (for example hydrogen bonds), and the formation of strong chemical bonds portrays chemisorption (for example electrostatic bonds) (Heo et al. 2019). The electrostatic bonds of attraction are a result of opposite charges between pollutant and adsorbent. For instance, metal oxides can act as an acid or basic material based on the pH_{zc} of the material. There are $-OH$ groups attached to the surface of metal oxide nanomaterials. The following reactions can occur in such a system that determines the adsorbent charges for electrostatic interactions (Kasprzyk-Hordern et al. 2003).



Bonds like Yoshida H-bonds, dipole-dipole, n - π , π - π , etc., can also be found. The Yoshida H-bonds result from bonds between hydrogen from functional groups (like $-OH$) on the adsorbent and the aromatic rings of the pollutant. The hydrogen on the adsorbent acts as a donor towards hydrogen-accepting groups (for example oxygen and nitrogen) on pollutants forming dipole-dipole H bonds. The n - π bonds are formed when electrons transfer from functional groups (example-carbonyl) of adsorbent to aromatic rings of pollutant. The interaction of π electrons of adsorbent and pollutant leads to the formation of π - π bonds, enhancing adsorption.

The adsorption isotherm curve relates the equilibrium concentration of micropollutant/ organic dye on the surface of the biosorbent to the concentration of these

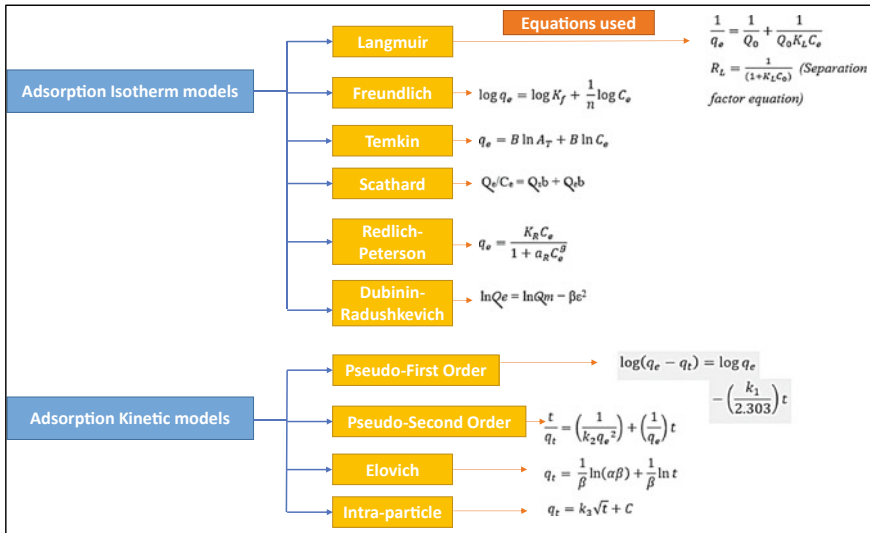


Fig. 14.2 Adsorption isotherm and kinetic models to validate experimental data

pollutants in the reaction solution at a given temperature. Adsorption kinetic models measure the diffusion of pollutants in pores of nanorods and tell about adsorption uptake with respect to time (for constant concentration or pressure). Figure 14.2 shows some adsorption isotherm and kinetic models which helps to validate the adsorption data. The values obtained through experimental analysis are put into equations of isotherm and kinetic models. The isotherm and kinetic plot which yields the highest correlation coefficient and least error function value is the best fit for the set of obtained experimental values (Guillossou et al. 2020).

14.5 Transition to Circular Bioeconomy Through Bio-based Materials

A circular economy is a close the loop approach where the wastes generated in the system are utilized in the system itself. The main strategy toward such a change involves: improving the product’s life span, keeping a product in operation, conserving materials, and preserving the energy embodied in materials (Moraga et al. 2019). The circular bioeconomy depends on several factors like environmental, economic, and social factors. The environmental factors cover areas of setting environmental awareness about the toxic effects of water pollutant exposure and regulations set to make these toxin levels under control. Economic factors involve the GDP, income, and job opportunities created if such an approach is adopted. Social

factors include perspectives like the education of people, age, and access to information. The implementation of a circular bioeconomy requires a lot of planning and proper execution to make these toxins under control in an environmentally friendly and economic manner (Mathew and Saravanakumar 2022b).

The transition to a circular bioeconomy for the elimination of organic pollutants on larger scales may involve several steps. The identification of the process involved in pollutant removal is a key step affecting the transition to a circular bioeconomy. This is because the nature of by-products (formed during the treatment) further determines the possibility of process applicability in a circular bioeconomy. If the by-products formed are hazardous, additional treatment should be employed for the removal of such toxins. This may incur additional expenses on large scale. Meanwhile, employing a treatment process that doesn't generate additional by-products or intermediates is advised. This can help to reduce additional costs incurred in the removal of intermediates/by-products. Hence, adsorption is suggested, as the process does not involve the generation of other by-products (as described in the sections above).

The selection of sorbent used for EDC and dye adsorption is another crucial step for implementing a circular economy. The sorbent should possess certain properties like high efficiency, should be economical, and be synthesized in an eco-friendly manner. The biosorbents can be used to remove organic dyes and EDCs from wastewater. The wastewater obtained after purification can be used to water the plants. These plants/ plant parts act as a source for the synthesis of biosorbents, thereby closing the loop in a circular economy. In such a scenario, the utilization of biomaterial can be an effective option. The treated sludge from wastewater can also enrich the soil, which yields trees, fruits, natural extracts, etc. in a circular bioeconomy. The advantage of utilizing biomaterials is that they can be used in energy storage and generation devices on larger scales. Figure 14.3 shows the sources for the generation of bio-based materials (from sludge and natural components) and their application (pollutant removal and energy devices) in a circular bioeconomy. The major challenge of implementing bio-based materials as adsorbents in a circular bioeconomy is about complete separation of pollutants from sorbents after adsorption.

Several countries (especially European countries) in the world are taking efforts to remove EDCs in effluents. The implementation of lab-scale experiments on large scales can be through considering the costs required for the upgradation of present techniques, availability, and accessibility of resources required. To implement the above in a circular bioeconomy it should be ensured that the biomaterial works for the elimination of multiple types of pollutants (for example EDCs and dyes) on larger scales. Hence, it is crucial to perform pilot scale studies for highly effective bio-based materials obtained at lab scales.

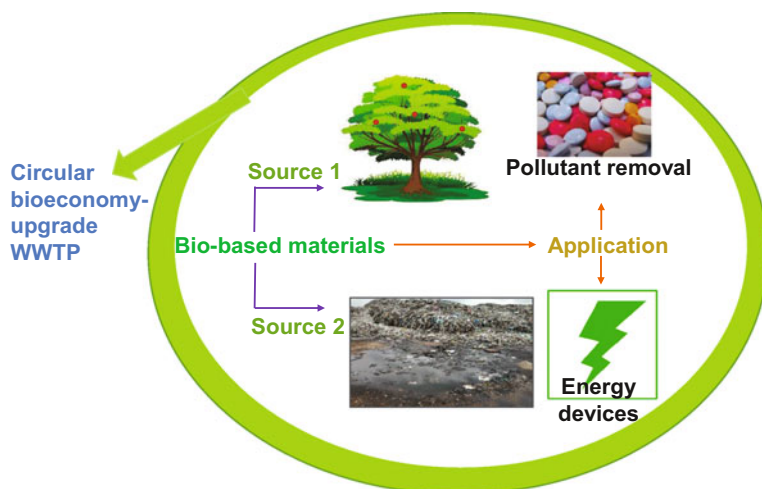


Fig. 14.3 Circular bioeconomy approach: sources and applications of biosorbents

14.6 Conclusion and Future Scope

The biomaterials show excellent morphological, physicochemical, and thermal properties, which help in the adsorption of micropollutants and organic dyes. The bonds formed between biosorbents and contaminants include electrostatic, π - π , n- π , hydrogen, and Vander Waals bonds. The application of biomaterials in energy devices gives scope for its implementation in a circular bioeconomy to upgrade wastewater treatment plants. More pilot-scale studies on energy applications and water pollutant removal using biosorbents should be carried out for a transition to a circular bioeconomy. The properties of biomaterials should be checked using characterization tools after each synthesis, considering the change in properties of sludge/ fruits/ natural components used as pre-cursor. Studies on the disposal/ reuse of pollutant-loaded biosorbents should be encouraged for a complete close the loop approach in treatment plants.

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Chapter 15

Role of Non-metallic Fraction Recycled from Waste Printed Circuit Boards for Producing Sustainable Construction Products



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Abstract A large number of waste printed circuit boards are produced annually, which looms large as a serious threat to our environment. When discarded in a landfill, printed circuit boards (PCB) contain poisonous and dangerous metallic and non-metallic fractions (NMF). This study illustrates the viability of using NMF recycled from waste PCB as one of the constituent materials in the manufacture of building goods like fly ash bricks and paver blocks as a possible route to environmentally friendly construction. The raw materials were combined with NMF in a variety of ratios ranging from 5 to 25%. In comparison to conventional products, NMF-incorporated fly ash bricks and paver blocks shown an improvement in compressive strength. Fly ash bricks made with NMF showed no efflorescence. In comparison to conventional products, NMF-incorporated products were observed to absorb less water. When compared to the control specimens, the NMF blended paver block specimens showed a significant resistance to water penetration.

Keywords Printed circuit boards · Non-metallic fractions · Fly ash bricks · Paver blocks

15.1 Introduction

The Global E-Waste Monitor 2020 estimated total e-waste generated globally about 53.6 million tons in 2019. India generated 3.2 million tonnes of e-waste annually in 2019 and is expected to produce 5 million tonnes by 2021 (Cui and Forsberg 2003). The report by Ministry of Environment, Forest and Climate Change (MoEFCC)

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estimates the recovery of metals such as Cu, Ag, Au and Pd is about 30 million tons per annum from waste PCBs. However, recycling PCBs is made difficult by the high non-metal content (up to 70%). Nevertheless, there is evidence about the possibility of using non-metallic fractions for construction applications. The non-metallic powder mainly consists of resins and glass fibres which do not need to be separated and their combined characteristics are very useful for cement composites (Veit et al. 2006). A few studies have been conducted in India related to the use of NMF in composites (Li et al. 2008; Guo et al. 2018). Janardhanan and Ramasamy (2017) used crushed PCB waste in fly ash based geopolymer concrete and suggested that replacing fine sand with 30% of crushed PCBs showed encouraging results. Kunal Kakria et al (2020) studied the addition of NMF by 5, 10, 15, and 20% in geopolymer concrete as an additive material and found that the addition of 15% NMF enhanced the mechanical performance. Using construction fillers, composite panels, bricks, and decorative cement, among other materials, Mou et al. (2007) investigated the effectiveness of incorporating non-metallic materials recovered from waste PCBs in construction. They came to a satisfactory conclusion when compared to conventional materials. Another study conducted in China by Guo et al. (2008) shown that the glass fibres and resin powder found in non-metallic debris from PCBs are also used to improve asphalt through a composition effect. But these are highly uneconomical since there is no commercial value of PCB waste as it is an industrial waste and due to the high costs of recycling.

Generally, PCB is discarded in dump yards and when they are burnt, which emit chemicals that can be harmful to human health. Exposure to the dangerous compounds found in e-waste can cause serious health problems (Qu et al. 2019). Pollutants that enter our environment, such as dioxins and furans from polyvinyl chloride, lead, beryllium, cadmium, mercury, etc., are known to pose a number of health risks, including those related to reproduction, the immune system, the nervous system, kidney damage, lung cancer, chronic beryllium diseases, and others (Guo et al. 2018). Thus, an efficient system to use them for a better purpose will serve the purpose. Senthil Kumar and Baskar (2014) have presented on E- waste as construction material and the results shows that there is decrease in strength for compressive, tensile and flexure but up to certain percentage of replacement it shows a good result and the concrete with partially replaced E-waste is recommended for non-structural elements which lead to reduction of electronic waste (Senthil Kumar and Baskar 2014). Lakshmi and Nagan (2011) have investigated on durability characteristic of E-plastic waste incorporated concrete such as chloride attack test, sulphate attack test, permeability test, Saturated water absorption, Porosity and Sorptivity. For the above test, the concrete with E-plastic waste shows a good result.

Construction materials with minimum damage to the environment is gaining momentum. Urbanization has made cement one of the most consumed materials for large scale production of concrete. Cement demand has topped a whopping 5 billion metric tons worldwide. The inclusion of large amount of raw materials has led to huge amount of CO₂ release in the atmosphere (Adesina 2020). Exhaustive research is being done across globe to find a supplementary cementitious material (SCM) to tackle the damage to environment. Various SCM's such as the Fly Ash

(FA), Ground Granulated Blast Furnace Slag (GGBS), Silica Fume (SF) and Rice Husk Ash (RHA) (Pal et al. 2003; Shanmuga Priya et al. 2021; Corona et al. 2019), etc., which are derived from industrial processes poses a suitable substitute to cement in concrete. On the other hand, with the increasing awareness about the effects of using naturally available resources in the manufacture of construction products, it is important to try and incorporate various other materials into these products to decrease the overall consumption of natural resources (Corona et al. 2019). With the restrictions on the use of natural aggregates like sand, etc. getting tighter day by day, the use of non-metallic fraction from recycled PCB as a replacement to fine aggregates can be thought of as an alternative.

In the present study, the NMF was used to make construction products like fly-ash bricks and paver blocks as additive, and to find the optimum percentage of NMF that can be added to achieve optimum strength. In comparison to conventional materials and fillers, NMF products exhibit improved mechanical properties and durability. Based on the test results it was concluded that optimum replacement of NMF was found to be 30% for fly ash bricks and paver blocks and maximum strength and durability properties were observed. This study suggested that the use of NMF in construction product can be one of the economic ways for their disposal of e-waste in environment friendly manner which contributes to save environment and conserve natural resources.

15.1.1 Objectives

The objective is to study the properties of NMF and its application in construction products. This research also focuses on the impact of NMF on the properties of the products.

- To collect and characterize non-metallic fraction from waste PCB.
- To assess the suitability of recycled non-metallic fraction in construction products.
- To find the mechanical strength of the NMF incorporated mortar samples and pavement blocks.
- To attain the optimum amount of non-metallic fraction to be added to achieve maximum strength output.

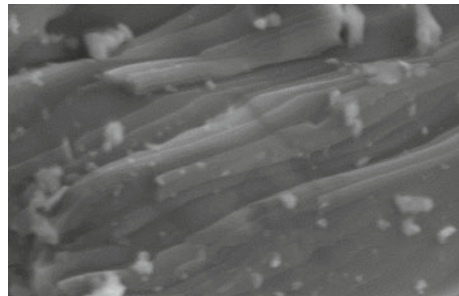
15.2 Materials and Methods

OPC of 53 Grade was used in accordance with IS 12269 (2003). The specific gravity of cement was found to be 3.15. The chemical composition of cement are presented in Table 15.1. A low calcium Class F fly ash [confirming with ASTM C311 (2013)] was used and its chemical compositions are given in Table 15.1.

Locally available natural sand conforming to Zone II as per IS 383 (1970), having maximum size of 4.75 mm was used. The specific gravity of fine aggregate was 2.65

Table 15.1 Chemical composition of class f fly-ash

Properties	Cement	Fly ash
SiO ₂	15.55	48
Al ₂ O ₃	8.40	24.3
Fe ₂ O ₃	5.87	15.6
SO ₃	3.12	0.4
CaO	62.44	3.2
Na ₂ O	0.16	0.8
K ₂ O	1.27	–

Fig. 15.1 SEM image of NMF

and fineness modulus being 2.36, bulk density was found to be 1675 kg/m³ with a water absorption of 1.8%. NMF from e-waste was procured from Victory Recycling, Kanchipuram. NMF being both fibrous and particulate in nature had particle size varying from 0.15 to 1.3 microns. NMF is made up of bundles of thermosetting resins and fibre that are loosely intertwined with glass fibres (Fig. 15.1).

Crushed, graded, angular coarse aggregates, of nominal maximum size 12.5 mm with the specific gravity and water absorption being 2.81 and 0.6%, respectively were used. The bulk density of coarse aggregate was 1550 kg/m³. Fresh potable water, free from organic and acid substances, was used for concrete preparation.

15.2.1 Mixture Proportions for Mortar Mixes

The effects on different percentages of NMF (partial replacement by weight of cement) were evaluated, expecting developments in the strength, durability and microstructural properties. A total of 5 batches of mortar samples were cast. Ordinary portland cement, river sand, NMF and Potable water was used. Mix design was done as per the procedure mentioned in ASTM C311 (2013). The mix proportion in kg/m³ is presented in Table 15.2.

All the materials were kept at room temperature (27 ± 2 °C) to maintain a uniform mixing temperature. A paddle mortar mixer was used to mix all the materials. The

Table 15.2 Mix proportion

Mix id	Percentage replacement		Quantity of materials (grams)			
	Cement	MBS	Cement	NMF	Sand	Water
CC	100	0	500	0	1375	242
M5	95	5	475	25	1375	242
M10	90	10	450	50	1375	242
M15	85	15	425	75	1375	242
M20	80	20	400	100	1375	242
M25	75	25	375	125	1375	242

mixing of materials was done for around 7 min. After mixing all the materials, the mortar was filled in the mould in 3 layers and compacted using the vibrating table for proper compaction. Tap water was used for curing these specimens for 28 days at room temperature (27 ± 2 °C). The specimens were immersed in water throughout the curing process. Figure 15.3 shows the casted cubes (Fig. 15.2).



(a) Raw material



(b) Mixing process



(c) Bricks after curing

Fig. 15.2 Flyash bricks preparation



Fig. 15.3 Paver blocks preparation

15.2.2 Mixture Proportions for Fly-Ash Bricks

The various raw materials that constitute the fly-ash bricks, along with their proportions used in the industry are: Fly-ash–50%, Lime–15%, Cement–10%, Crusher dust–25% and 150–200 ml of water is used per kilogram of the raw mixture. Fly-ash bricks were cast with varying proportions of NMF replacing fly-ash (0, 5, 10, 15, 20 & 25%). The raw materials were mixed and blended in a miller drum. A fly-ash brick weight about 3.3 kgs. Therefore, a batch of 20 kg of this mixture was taken for 6 bricks (each mix) to get samples with desired proportion of NMF.

15.2.3 Mixture Proportions for Paver Blocks

The various raw materials that constitute the fly-ash bricks, along with the mix proportions used in the industry are:

Cement: Fine aggregates: Coarse aggregates–1: 2.5: 3.25.

Water-cement ratio (w/c)–0.4

Samples of M-35 grade paver blocks were cast with varying proportions of NMF replacing Cement (0, 5, 10, 15, 20 & 25%). The raw materials were mixed and blended in a mixing drum. A paver block weight about 5.4 kgs. Therefore, a batch of 33.75 kg of this mixture was taken for 6 blocks (each mix) to get samples with desired proportion of NMF.

15.2.4 Testing Procedures

Fly ash bricks and paver blocks were used to test the compressive strength as per Codal provisions. For every mix, 6 samples were cast. The strength was determined at

Fig. 15.4 Compression test of paver block



Fig. 15.5 Industrial oven with paver block



the age of 7 and 28 days of curing. The test was conducted using AIMIL compression testing machine with a constant load rate of 2.5 kN on the cubes (Figs. 15.4 and 15.5).

15.2.5 Compressive Strength Testing

Samples of fly ash bricks of size 230 mm X 110 mm X 70 mm were cast. Paver Blocks of size 300 mm X 100 mm X 70 mm were cast. The raw materials with varying proportions of NMF are blended in a blender drum with optimum water content. The mixture is then transferred to hydraulic pressing machine where the blend is fed into the molds and then pressed under pressure. The bricks are then

dried under the sun for 24–48 h. These bricks are then cured by sprinkling water for 21 days for fly ash bricks and paver blocks were cured for 7 and 28 days. At the 7th, 21st and 28th daytests were carried out on the sample utilising a compression testing equipment with a 1200 KN capacity. Three samples per mix were tested and their average was considered in the results. The brick specimens were placed in the compression testing machine on their flat surfaces between plywood sheets. Load was applied axially at a uniform rate of 14 N/mm per minute till failure occurred and the maximum load at failure was noted. The compressive strength was calculated using the equation,

$$\text{Compressive strength} \left(\frac{\text{N}}{\text{mm}^2} \right) = \frac{\text{Ultimate load (N)}}{\text{Area of the specimen (mm}^2\text{)}}$$

15.2.6 Water Absorption Testing

The fly ash bricks and paver blocks were tested in accordance with the procedure laid down in IS3495 (Part 2) for the total water absorption. Specimens were dried in a ventilated oven at a temperature of 105 to 115 °C till it attained substantially constant mass. The specimens were cooled to room temperature and its weight (M_1) was obtained. The completely dry specimens were immersed in clean water at a temperature of 27 °C for 24 h. The specimens were removed and any traces of water were wiped out with a damp cloth. The specimens were weighed 3 min after removing from water, and its weight (M_2) was obtained. Water absorption, percent by mass, after 24-h immersion in cold water is given by the following formula:

$$\text{Water absorption (\%)} = \frac{(M_2 - M_1)}{M_1} \times 100$$

15.2.7 Microstructural Analysis

SEM the raster scan image which is generated can target a resolution higher than 1 nm. Before being considered for SEM analysis, the sample was broken down into smaller sections by grinding, after undergoing the compressive strength test. After which, the samples were cleaned and oven-dried at a temperature of about 96 °C. The samples were prepared in a powdered form, which passed through a 75-micron sieve. Furthermore, the uncoated sieved samples were then kept securely over a Canada Balsam coated glass plate. The samples are then analysed through LEICA microscope and visualized with the aid of QWIN software.

15.3 Result and Discussion

15.3.1 Compressive Strength Results—Mortar Samples

Compressive strength values of NMF incorporated mortar samples shown in Table 15.3. Compressive strength showed an increasing trend till 20% replacement. A compressive strength of 24.02 N/mm² was observed in the 28th day mortar sample. On addition of 10% a percentage increase of 1.87 was observed. On 15% replacement a percentage increase of 0.82% was observed and on 20% replacement an increase in strength of 1.45 was observed. Up to 20% of replacement, the compressive strength exhibited an increasing trend. Compressive strength of 25.04 N/mm² was found to be the maximum with 20% replacement. The generation of CSH gel is what causes the rise in compressive strength, and the addition of NMF has improved gel formation and increased strength even more. Table 15.3 provides the compressive strength results (Fig. 15.6).

Table 15.3 Compressive strength results

Mix ID	Sectional area	Compressive strength in N/mm ²	
		7 days	28 days
MCC	2500	15.78	24.36
M5	2500	15.34	24.02
M10	2500	15.47	24.47
M15	2500	15.53	24.67
M20	2500	15.87	25.04
M25	2500	15.27	24.01

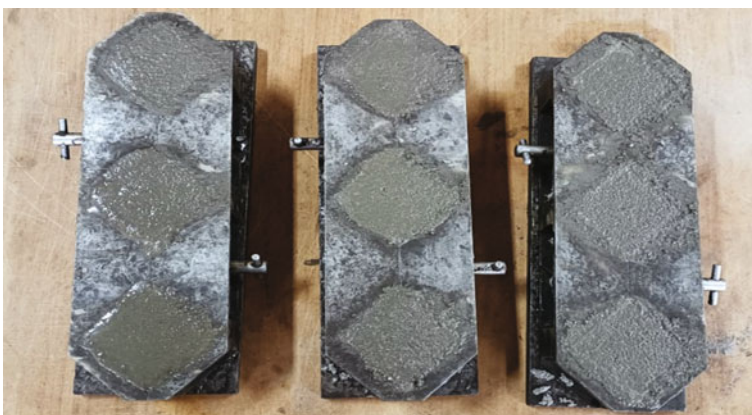


Fig. 15.6 Cubes for compressive strength testing

Table 15.4 Compressive strength for fly ash brick at 21 days

Mix ID	Sectional area (mm ²)	Average load at Failure (kN)	Average strength (N/mm ²)
FAB-CC	25,300	343.34	13.57
FAB-5	25,300	351.67	13.89
FAB-10	25,300	366.67	14.49
FAB-15	25,300	370	15.04
FAB-20	25,300	373.34	16.98
FAB-25	25,300	363.34	16.40

15.3.2 Compressive Strength—Fly Ash Bricks

Average strength values of NMF incorporated fly ash bricks at 21 days. It is observed that on 5% replacement of NMF the strength was increased to 13.89 N/mm², as the percentage replacement increased the strength also increased. On 10% replacement there was a percentage increase of 4.32 of compressive strength. The percentage increase on 15% replacement was 3.79 when compared to 10% replacement. Up to 20% replacement, the compressive strength exhibited an increasing trend. The compressive strength was seen to improve to FAB-20. FAB-20, which replaced 20% of the crusher dust with NMF and had a maximum compressive strength of 16.98 N/mm², was found to have the highest compressive strength. The formation of cementitious gel around them has imparted strength to the material. Due to addition of NMF, the formation of this gel is further enhanced due to the increased number of particles. The fibrous glass particles present in it contributes to the increase in strength. There was a slight decrease in strength on the addition of 25% NMF. The increase in strength is due to the formation of cementitious gel around them. Due to the addition of NMF, the formation of gel is further enhanced due to the increased number of particles. Also the fibrous glass particles present in it contributes to the increase in compressive strength. The results are given in Table 15.4.

15.3.3 Compressive Strength Results—Paver Blocks

Paver block samples were tested at 7 day and 28 day for compressive strength. For each mix three samples were casted and tested on 7 day and 28 days and the average of the three readings were taken and recorded in the Tables 15.5 and 15.6.

Average strength values of NMF incorporated paver brick at 7 and 28 days is shown in Table 15.3 and 15.4. For both curing age the compressive strength values showed an increase with 5% (PAB-CC) to 0% (PAB-25) replacement of NMF. The percentage increase on 5% replacement was 7.42 compared to conventional mix. On 10% replacement it was 4.73 when compared to 5% replacement. A percentage increase of 2.77 was found in 15% replaced mortar sample. 20% replacement has

Table 15.5 Compressive strength for paver blocks at 7 days

Mix ID	Sectional area (mm ²)	Average load at failure (kN)	Average strength (N/mm ²)
PAB-CC	30,000	643.2	14.4
PAB-5	30,000	673.5	15.6
PAB-10	30,000	675	17.2
PAB-15	30,000	678	18.5
PAB-20	30,000	696	19.4
PAB-25	30,000	538.89	17.6

Table 15.6 Compressive strength for paver blocks at 28 days

Mix ID	Sectional area (mm ²)	Average load at Failure (kN)	Average strength (N/mm ²)
PAB-CC	30,000	1051.5	25.6
PAB-5	30,000	1125	27.5
PAB-10	30,000	1128.3	28.8
PAB-15	30,000	1129.5	29.6
PAB-20	30,000	1148.1	30.2
PAB-25	30,000	1050	29.1

shown a percentage increase of 2.03 when compared to 15% replaced sample. The paver blocks has shown an increase in strength till 20% addition of NMF. The increase in strength is due to more finer particles than sand in NMF. Due to the addition of NMF the finer particles fills the voids when vibrated in vibrating table. The fibre like glass particles has also contribute to the increase in strength. Similarly a dip in strength was also observed with 25% replacement incorporation of NMF for both curing age. Optimum strength value was observed at 20% replacement of M sand with NMF with maximum compressive strength.

15.3.4 Water Absorption Results—Fly Ash Bricks

The results of all the mixtures water absorption tests were displayed in Table 15.7. It was shown that when NMF was substituted for fly ash brick at various percentages, as percentage replacement increased water absorption decreased. On percentage increase in replacement of NMF the water absorption was decreasing, on 5% replacement the percentage of water absorbed was found to be 21.6% on 10% replacement it was found to be 18.3%. Water absorption was 16.2% and 15.8% on 15 and 20% replacement of NMF respectively. Lowest water absorption percentage was observed in 20% NMF in fly ash brick. The optimum water absorption percentage was 20%.

Table 15.7 Water absorption results—fly ash bricks

Mix ID	Dry weight (Kg)	Wet weight (Kg)	% Water absorbed
FAB-CC	3.303	4.035	22.2
FAB-5	3.213	3.892	21.1
FAB-10	2.970	3.521	18.5
FAB-15	2.789	3.259	16.8
FAB-20	2.817	3.246	15.2
FAB-25	2.532	2.898	14.5

The existence of finer NMF particles and its pozzolanic feature, which led to a discontinuous pore structure, can be a cause of this positive result.

15.3.5 Water Absorption—Paver Block

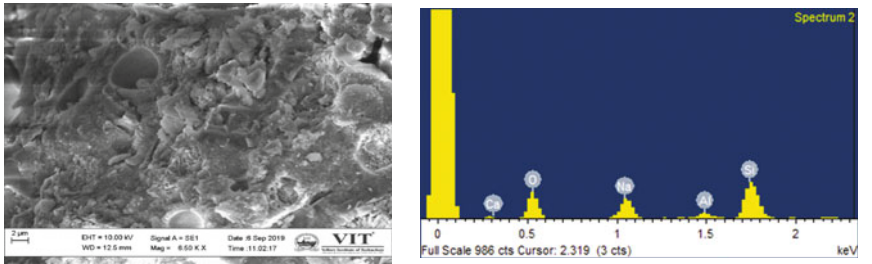
The water absorption performance of the mixes is presented in Table 15.8. Water absorption in paver blocks was found to be decreasing on increasing percentage replacement. 5% replacement has shown a water absorption of 9.33%. Compare to convention mix the water absorption decreased to 7.5%. On addition of 10% NMF the water absorption decreased to 8.27% and it was observed that on 15% and 20% replacement it was 8.04 and 7.50 respectively. Decrease in the water absorption is observed on the addition of 25% NMF with lowest water absorption. Lowest water absorption percentage was observed in 25% NMF in paver blocks. The optimum replacement percentage was 25%.

15.3.6 Microstructural Analysis—SEM

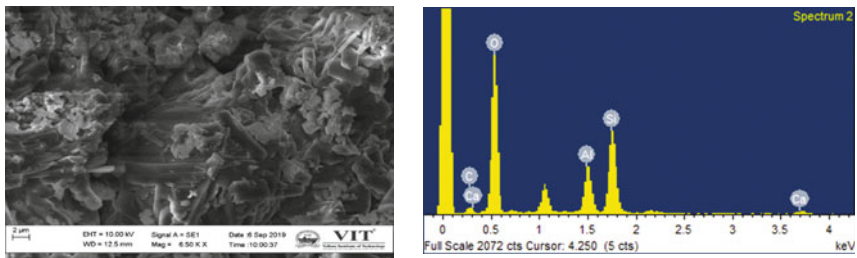
The conventional and 15% replacement of cement by NMF replacement mix were studied through micro-structural analysis and compared with CC to evaluate the

Table 15.8 Water absorption results—paver blocks

Mix ID	Dry weight (kg)	Wet weight (g)	% Water absorbed
PAB-CC	5.327	5.892	10.60
PAB-5	5.035	5.506	9.35
PAB-10	4.962	5.375	8.32
PAB-15	4.627	5.001	8.08
PAB-20	4.280	4.598	7.43
PAB-25	4.105	4.413	7.50



(a) SEM image of Conventional Cement Mortar (CC)



(b) SEM image of NMF

Fig. 15.7 SEM images of CC and NMF

texture and effect of NMF in cement mortar. Figure 15.7a shows the microstructural arrangement of CC, being a uniformly distributed arrangement of materials in absence of any major cracks visually observable. Unreacted particles are visible on the surface of the surface which shows incomplete pozzolanic reaction. Moreover the presence of small amount of Si is reflected in the EDS analysis. Figure 15.7b shows SEM image of NMF and NMF fibers can be seen embedded in the micro-structure of the mix M20. The coexists of calcite and CSH was identified in EDX and white layer of calcite on the top of fibres ie CSH matrix was seen in SEM images. Interstitial voids can be inferred which is found to decrease in the porosity of the materials as reported in waerabsorption test results. An enhanced bonding between the sand and binder is evident as compared to the controlled mix.

15.4 Conclusions

This study shows NMF can be reused in a very efficient manner by using it as a filler material for fly ash bricks and paver blocks. It has been identified that the properties NMF could increase the strength of fly ash bricks and paver blocks. From the various mixes with different proportion of nonmetallic fraction and from various

tests taken to check the strength and durability of the products that have been casted the following inferences were drawn.

- The PCB nonmetallic fraction can be used in construction products.
- The waste PCB nonmetallic fraction improves the compressive strength of fly ash bricks and strength of paver blocks. This increase in strength can be attributed to the glass particles and fibrous materials present in NMF.
- Mortar cubes with 20% replaced NMF showed the maximum compressive strength.
- It is observed that the addition of NMF in fly-ash bricks and paver blocks improved compressive strength up to 20% replacement with cement.
- The optimum replacement of NMF was found to be 25% for fly ash bricks and paver blocks.
- Micro-structural analysis of fly-ash bricks showed the effect of fibrous nature of NMF in increasing the overall performance.
- The NMF addition products can be used for buildings as a sustainable and economical way for disposal of e-waste.

From above experimental results and discussions, it is proved that the replacements are the good alternative materials for reducing crusher dust & fine aggregate. Use of NMF in construction product can be one of the economic ways for their disposal in environment friendly manner which contributes to save environment and conserve natural resources.

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Chapter 16

Bioconversion of Organic Waste for Solid Waste Management and Sustainable Agriculture—Emphasized Impact of Bioelectromagnetic Energy



Parul Vats, R. Y. Hiranmai, and Ajay Neeraj

Abstract Municipal solid waste (MSW) management is a pressing issue in developing nations across the globe. Organic fraction makes up a major fraction of the waste composition. Owing to its high organic matter, high abundance and low concentrations of pollutants, MSW is suitable for bioconversion. Old age-established concepts like composting and vermicomposting offer rearrangement and recycling of the organic components of waste through the cyclic pathways in the natural environment lowering the quantity of waste and upcycling it to value-added organic manure. Food production owing to the green revolution has proved to be unsustainable in the longer run. Sustainable agriculture embraces judicious ecosystem management and discourages the use of synthetic inputs. Fortified quality organic manures can be directed for sustainable agricultural practices achieving natural circularity, safe quality food production and upliftment of the physicochemical and biological quality of soil and crops. Engaging the other hidden half of nature- the human world of intents and thoughts, positive-divine thought-based bioelectromagnetic energy generated by meditation during the farming process- Sustainable Yogic Agriculture (SYA) has experimentally shown enhanced levels of agroecosystem health, microbial activity and crop production. The present work reviews the bioconversion of locally available organic waste to manures for nutrient management of agricultural soil and crop production and their quality enhancement by incorporating a transdisciplinary SYA approach. Nature has multidisciplinary intricate linkages and is highly receptive to human emotions. Transdisciplinary practical solutions can become potential solutions to realize the vision of a sustainable nation.

Keywords Agriculture · Bioconversion · Organic waste · Soil fertility · Sustainability · Yogic farming

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16.1 Waste Generation Scenario

The world population has been on the rise and so have developmental activities which is the source of the enormous amount of waste. Approximately 10,000 tons of solid waste are disposed of in Lagos each day, and a large amount of electronic waste is disposed of from the 500 container ships that discharge waste there each month. In the new atlas of the world's biggest dumps, Mbeubeuss is one of 50 giant sites. It is a contender for the title of the largest dump in the world. As a comparison, there is the massive Agbogbloshie dump in Ghana, which is one of the largest in the world and takes 192,000 tonnes of electronic waste every year. Also, the Awotan dump in Ibadan, Nigeria, which has become a breeding ground for diseases. There is a massive tip in Bekasi, Indonesia, called Bantar Gebang, which takes 230,000 tonnes of industrial, medical, and farm waste a year at Dandora outside Nairobi.

About two billion tonnes of municipal solid trash are produced yearly throughout the world, with daily garbage production averaging 0.74 kilos and ranging from 0.11–4.54 kg. Under a “business as usual” scenario, this quantity is projected to increase to 3.40 billion tonnes by 2050. The three high-income nations of Bermuda, Canada, and the United States currently produce the most garbage per person on average, 2.21 kg per day, whereas low-income nations produce 0.74 kg per day (Kaza et al. 2018). Many cities in the world and major cities in India have been struggling due to mountains of waste. Bantar Gebang landfill in Jakarta (Timmerman 2021); Chile's Atacama Desert (Agrawal 2021); Ghazipur, Okhla and Bhalswa in Delhi (Times of India 2022); Bandhwari in Gurugram (Harigovind 2022); Deonar in Mumbai (Hindustan Times 2022); Mittaganahalli landfill in Bangalore (VL 2021) are some examples highlighted in recent years. About 33 percent of the waste worldwide is openly dumped and not disposed of in an environmentally sound manner (Kaza et al. 2018). The basic amenity of sound solid waste management is not yet properly organized in many parts of the world and exhausts quality of life and natural resources.

MSW generation rates are influenced by economic development, industrialization, public habits, and local climate. Income level and urbanization promote increased consumption of goods and services and hence increase the amount of waste generated (Hoorweg et al. 2012; OECD 2021). High-income countries (16% of the world's population) generate about 34 percent of the world's waste. The total quantity of waste generated in South Asia is expected to increase by two times by 2050 (Kaza et al. 2018).

Planning Commission stated that urban residents produce double the waste as rural residents and India is moving into urban areas. 377 million i.e. about 30% of the total population live in towns, it will be more than 50% by 2050. 81% of the total MSW is generated in urban cities. Waste collection efficiency is between 70 and 90% in major Metro cities but the treatment of waste is very improper. Many cities other than composting and refuse-derived fuel (RDF), which is done in limited quantities (Planning Commission 2014). Urban waste management needs urgent attention and priority across the globe and in India.

16.2 Factors Influencing Municipal Solid Waste Generation

MSW generation has been constantly escalating. Several factors influence the generation of waste. Hoornweg et al. (2012) reported that a population of about 3 billion residents generates 1.2 kg per person per day (1.3 billion tonnes per year). The world population is on rise and currently stands at about 7.9 billion (United Nations n.d.). With the population surge, the world generates 2.01 billion tonnes of municipal solid waste annually, per person per day averages 0.74 kg in the range of 0.11 to 4.54 kg and by 2030, waste generation worldwide is expected to reach 2.59 billion tonnes per year (Kaza et al. 2018). MSW generation is affected by economic conditions, industrialization, urbanization, consumer habits, and local climate. Income level, population and urbanization affect waste generation highly by increased consumption of goods, services and materials consumption, and consequently the amount of waste generation (Kawai and Tasaki 2016; Hoornweg et al. 2012).

As per an assessment report by OECD (2021) using United Nations population trends and World Bank development indicators, in a span of 27 years from 1990 to 2017, 2.5 billion people are added to the world population to 7.5 billion people at present, this creates an impact on consumption and production rates. The gross domestic product across the globe per capita has increased by fifty percent. As a consequence, the annual material consumption has almost doubled (from 37 billion tonnes to 88 billion tonnes) in 2017 from 1992. The average daily materials used per capita increased from 22 kg in 1990 to 33 kg in 2017. Economic conditions of countries have a distinct impact on waste contribution. High-income countries, although only account for 16% of the world's population, contribute to more than 30 percent, of the world's waste. Other than increased consumption, global trade and exchange come along. As pointed out earlier that presently, Bermuda, Canada, and the United States are the biggest generators of waste per capita with 2.21 kg per day, but in the coming future low-income countries are expected to see waste generation increment by more than three times and South Asia will also witness a doubling of waste generation. In these regions, 33% of the waste is not handled in an environment safe manner and is openly dumped (Kaza et al. 2018), thus requiring urgent action in terms of waste handling, management and disposal. Trade influence is high in this aspect. According to OECD (2021) projections, a relative decoupling between economic growth and materials use will take place in the coming decades with the annual economy growth rate of 2.8%, global materials use is set to increase on average by 1.5% every year. This relative decoupling will happen owing to technical advances and structural changes in the economy. Increasingly, countries are expected to shift toward a service-based economy at the global level, which will reduce the material intensity and global material use and improved technological advancements will help save more than 60 billion tons of materials. However, additional policy measures are required to support the benefit obtained from structural and technological changes.

World bank reports that waste composition differs across income levels, reflecting varied patterns of consumption. High-income countries generate lesser organic, biodegradable waste (32% of the total) and more dry recyclable waste like plastic,

paper, cardboard, metal glass etc. (about 50% of the total waste). Middle- and low-income countries have bigger fraction of organic waste (53% and 56% respectively) (Kaza et al. 2018). Across the globe as well, by 2050, 55% population is likely to move into urban regions automatically promoting higher levels of material use in cities (OECD/European Commission 2020). Globally cities produce half of the solid waste and 70% of greenhouse gas emissions (IEA 2016) but cities hold important facilities for resource efficiency and if proper coordination between authorized bodies is executed, cities can progressively move towards circularity of solid waste management (OECD 2021). Low-income countries are still figuring out proper waste disposal mechanisms and global trade of waste has promoted economically efficient recycling, but also increased waste burden in the recipient countries and pose environmental risks if not handled carefully. Trade influences waste management and treatment. Recycling sector is highly influenced by trade. Trade of waste between nations has promoted economically efficient recycling, with that comes the risks of environmental pollution if the management of waste is not under checks and regulations. The trade has increased by around 30% (by weight) and the main categories are papers and metals (Yamaguchi 2021; Garsous 2019). The largest exporters of waste and scrap in 2018 were the United States, Germany, Japan and France. Import of waste was highest in China along with India and Turkey. Regulatory policies and regulatory instruments, recycling targets, product standards, recycled content requirements, lifetime warranties, bans and restrictions and deposit-refund systems are useful policies to increase resource efficiency and economically efficient and environmentally sound waste processing and disposal (OECD 2021).

16.3 Present Status of Waste Generation in India

Urbanization and sloppy waste management directly contribute to garbage production and pollution. An expanding economy produces more solid trash, such as household, commercial, institutional, and industrial garbage, which substantially raises the volume of waste produced from diverse sources. Garbage, junk (package materials), waste from building and demolition, leaf litter, hazardous wastes, etc. are only a few of the wastes produced in a normal metropolitan civilization (Rajput et al. 2009). In August 2009, FICCI carried out a study on solid waste management in Indian cities. Cities that produce more than 1000 tonnes of solid trash per day, such as Ahmedabad, Kanpur, and Pune, each have just one dumpsite, but Faridabad and Jamshedpur, which produce less than 450 tonnes of solid garbage per day, each have two pits.

16.3.1 Waste Generation and Disposal Status of Indian Cities

There are eight cities where 36% (8 out of 22) generated more than 1000 tons of waste per day (Ahmedabad, Delhi, Greater Mumbai, Jaipur, Kanpur, Lucknow, Pune,

Surat). Indore, Ludhiana and Vadodara generated 13.6% of the waste generated in 22 cities (500–1000 TPD). 11 cities (Agartala, Asansol, Chandigarh, Faridabad, Guwahati, Jamshedpur, Kochi, Kozhikode, Mangalore, Mysore and Shimla) generated less than 500 TPD of waste of 22 cities surveyed, 63.6% (14 out of 22) dispose of more than 75% of their rubbish in dump yards (Ahmedabad, Asansol, Chandigarh, Delhi, Faridabad, Greater Mumbai, Jaipur, Jamshedpur, Kanpur, Lucknow, Ludhiana, Mangalore, Pune, and Vadodara). On the whole, 47.05% (8) of the 17 class I cities had a single dump site, 29.4% (5) had two dump sites, 5.88% (1) had three dump sites, and 11.76% (2) have 4 dumpsites. A total of 100% of the waste collected in Greater Mumbai and Ludhiana is dumped at the dumpsite.

16.4 Problems Caused Due to MSW Mismanagement: Waste Decomposition in Open Dumpsites and Their Impacts

Waste production across the globe and India is ever-increasing and it is impossible to control and manage liquid and gaseous by-products of waste decomposition at open dumpsites without the necessary facilities and measures.

By burning plastics and other materials openly (usually to reduce volume), toxic substances are released into the atmosphere. As a result of these toxic fumes, the chronic respiratory disease can be caused and air pollutant levels are increased, including nitrogen oxides (NO_x), sulfur oxides (SO_x), heavy metals (mercury, lead, chromium, cadmium, etc.), dioxins and furans, and particulate matter. Open dumpsites have a negative impact on health and the environment because of:

1. Unplanned siting.
2. Operations are chaotic since there are no overarching standards defining how the facility should be run and many owners and operators of these dumpsites lack the essential tools and knowledge.
3. Ineffective waste input control, either in terms of amount or composition (or both).
4. Random garbage dumping at the dump site.
5. Lack of regulation of the pollutants emitted during waste decomposition and waste burning.

16.4.1 Effect on Health

Dumping of solid waste is hazardous to human health and causes various diseases, waterborne as well as vector borne diseases. It is due to the standing of water in an area and the flow of water is stopped due to dumping near the area and can also result in flooding by obstructing storm water. This waste is dumped because

people frequently leave their trash on the roadways, where it is then dispersed by rag pickers who are looking for food. There have been reports of a greater incidence of congenital birth defects, malignancies, and respiratory illnesses in people who live or work close to landfill dumps. World Health Organization (WHO 2000, 2007) says there is insufficient data on the health effects of landfills and incinerators, including cancer, reproductive issues, and mortality. In contrast, studies from Ray et al. (2005, 2009) have confirmed this at the Okhla dump site in Delhi. Comparing workers of the same age, gender, and socioeconomic status to controls of similar age, gender, and socioeconomic status, 62% had lung infections. Workers in landfills are more likely to suffer cardiovascular disease and tissue damage due to leukocytes, platelets, and inflammation in the air (Ray et. al. 2009). These and other types of issues are maintained by a vicious cycle that includes municipal solid wastes and the pollutants emitted as a result:

- (i) Chemical poisoning through chemical inhalation
- (ii) Low birth weight
- (iii) Cancer
- (iv) Congenital malformations
- (v) Neurological disease
- (vi) Nausea and vomiting
- (vii) Mercury toxicity from eating fish with high levels of mercury.

Non-Hodgkin lymphoma and soft tissue sarcomas are the illnesses that are frequently examined in epidemiological studies near MSW incinerators. This is the case because abandoned materials, like old tyres, often contain standing water that serves as a breeding ground for fleas and mosquitoes that can spread illnesses like malaria.

16.4.2 Effect on the Environment

Open dumping is a persistent issue in our nation and around the world, as particular areas turn into common places and dump piles draw new dumping. Disposing of garbage produced by communities and industry on land, in the sea, or in low-lying areas is a highly widespread activity. Sites may be entirely or primarily made up of home or industrial garbage.

Biodegradable waste components (food and yard waste) generally degrade anaerobically at landfills and dumpsites. Additionally, methane (about 50–65%) and carbon dioxide (about 35–45%) are generated as a result of the decomposition of the waste. Several researchers have measured the rates of atmospheric gas release via a landfill cover. Methane and carbon dioxide emissions from landfill surfaces have a big impact on the greenhouse effect and global warming. Methane released into the atmosphere significantly contributes to global warming since its global warming potential is about 21 times that of carbon dioxide (Esakku et al. 2005). Extensive characterization studies of the world's methane sources and sinks have been prompted by recent

increases in atmospheric methane concentrations. This study found that atmospheric methane concentrations were rising on average between 1 and 2% per year. Methane is estimated to be responsible for around 18% of all global warming (Church and Shepherd 1989). Methane is flammable since it is lighter than air. Sparks or flames are likely to cause serious explosions in closed buildings if its concentration reaches as high as about 5–15 percent in the air (Esakku et al. 2005). The dumping and land-filling of waste are estimated to contribute between 5–15% to methane emissions into the atmosphere. Multiple environmental and health risks are produced by the open disposal of solid trash.

The management of all produced garbage must be done appropriately. In India, the location for waste disposal is chosen based on convenience rather than taking the environment into account first. Communities near upstream plants and those around the waste dumpsite may be impacted in different ways. Diversity is referred to by scientists as a technique to determine whether a location has been disturbed. The ecology expert examined the region around the dumps and a “reference site” that was distant from the dumps but had similar topography and water flow patterns. The reference location and the dump region should have had a lot of the same flora if the dump wasn’t having any effects on the soil. The local plant population did not significantly differ from that of the reference region without a dump. However, the waste areas have less plant diversity than they ought to have due to worse management and design. The plant diversity at the YK Delta landfill, which was also poorly maintained, was higher than at the reference site but was partially brought on by weedy plants like pet sites, fireweed, grass, and sourdock. Additionally, there were few of each kind of plant. As a result, just a few numbers of plants would predominate in poorly maintained wastes, and it would be more difficult to find other kinds of plants.

The impact of improper solid waste management is finding its way into water bodies as well and impacting marine and terrestrial life. About 80 percent of ocean plastic comes from land and of that 75 percent comes from poorly operating MSWM systems (Kaza et al. 2018).

16.4.3 Effect on Soil

In order for plants to develop, the soil must be “a dynamic natural body on the surface of the earth, made of mineral and organic components and living forms”. It also serves as a layer of protection and filtration that is placed over groundwater to lessen the effects of many dangerous pollutants (Venkatesan and Swaminathan 2009). The weight of MSW on land has increased due to urbanization and industrialization, which is negatively affecting the soil’s yield and biotic and abiotic qualities. In comparison to control sites, the soils at the disposal sites had higher pH, TDS, and electrical conductivity regimes. Lead (Pb), Copper (Cu), Nickel (Ni), Chromium (Cr), and Zinc (Zn) were among the heavy metal concentrations that were found to be greater at the waste dumping sites, except for Cadmium (Cd), which had a higher

value at the control site. Salinity is another major danger to soil; it results from incorrect human land use, including deforestation, construction work, and industrialization. Salinity has an impact on the physical and chemical makeup of soil, leads to soil degradation, and raises the salinity of subsurface waters (Iwai et al. 2013). Additionally, (Biro et al. 2013) shown in their research that a decrease in soil productivity may be caused by a change in soil characteristics, which may have been 25 related to a change in land use and cover. In natural ecosystems, the restoration of degraded soil is a challenging and protracted process that involves the development of a vegetative community and the resumption of microbial activity (De Souza et al. 2013). Total nitrogen, pH, and the degree of erosion are critical variables that affect soil growth. Soil richness reduced as these factors rose (Pallavicini et al. 2015). Quantifying the ecotoxicological risk of composted residuals is a topic of growing attention. The review by Kapanen and Itavaara (2001) discusses the wide range of ecotoxicity test techniques for compost that use microbes, enzymes, soil fauna, and plants. An early overview of the impacts of MSW-compost incorporation on soil biological characteristics is provided by Gallardo-Lara and Nogales (1987). These typically included rises in the populations of bacteria and fungi in modified soil.

16.5 Municipal Solid Waste Management: Treatment and Practices for Municipal Solid Waste Management

The proper management of solid waste promotes economic growth and a higher standard of living while minimising or eliminating negative effects on the environment and human health. For a municipality, managing garbage successfully involves a variety of procedures. These comprise supervision, gathering, transportation, processing, recycling, and disposal (Fig. 16.1).

16.5.1 Waste Reduction, Reuse and Recycling

Waste reduction and product reuse are examples of trash avoidance strategies. Waste avoidance is given top priority in integrated waste management. Backyard composting can significantly reduce waste volume.

Removing items from the waste stream so that they may be utilized as raw materials to make new goods is the process of recycling (USEPA 2022). Accordingly, recycling happens in three stages: first, waste is separated, recyclables are collected, and then the recyclables are converted into raw materials. These fundamental resources are then used to create new items. It is theoretically feasible to recycle a wide range of materials, including plastics, wood, metals, glass, textiles, paper, cardboard, rubber, ceramics, and leather. Increased diversity of recyclable raw materials is often needed in regions with a broad industrial base and economy. The economy, the environment,

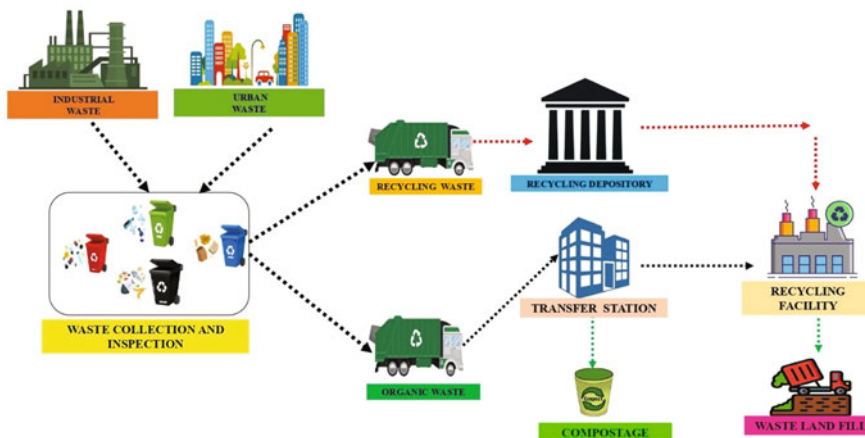


Fig. 16.1 Recycle/composting process of waste in urban area

and society are all benefited by recycling. Because recycling consumes less water, energy, and pollution than obtaining new raw materials, it lowers operating costs (Sandin and Peters 2018).

16.5.2 Treatment and Disposal

With the use of waste treatment procedures, trash may be made simpler to dispose of by being changed into a more manageable form, having its volume reduced, or having its toxicity decreased. In order to select the best treatment option, we need to know the size, shape, and composition of the waste material. Today's waste treatment techniques include using biological processes, exposing the garbage to extremely high temperatures, disposing of it on land or in landfills, and dumping it. Biological degradation can be achieved by aerobic degradation or composting.

16.5.2.1 Anaerobic Digestion

Composting and anaerobic digestion both employ biological processes to break down organic waste. Methane and carbon dioxide are produced as a result of anaerobic respiration. Anaerobic digestion is used to produce biogas, which may be used to produce power, in addition to the humus that is produced and utilised to improve soil. Nutrients like nitrogen, phosphorus, and potassium are needed for the process to run well. The pH must also be kept around 7, the alkalinity must be suitable to buffer pH variations, and temperature must be managed (Christy et al. 2014).

16.5.2.2 Composting and Its Potential in Solid Waste Management

Composting is the controlled aerobic breakdown of organic material by tiny invertebrates and microbes. Aerobic degradation generates better quality by products as compared to anaerobic composting. There are several composting methods in use right now. These include windrow composting, static pile composting, in-vessel composting, and vermicomposting. Municipal solid waste composting is influenced by a number of parameters, including waste source and type, composting design, maturation period, and composting process (Hargreaves et al. 2008). Composition and components of the materials, namely their Carbon/Nitrogen (C/N) ratio, temperature, moisture content, and air volume all affect how quickly compost forms. For the procedure to be effective, the C/N ratio is crucial. Carbon serves as the microbes' energy source, while nitrogen is necessary for the creation of certain proteins. The use of dehydrated mud is useful when the C/N ratio is too high, and the use of cellulose can be used when the ratio is too low. While the moisture content of the compost is significantly affecting the composting process, it is not the only factor.

The wetness is necessary for the microorganisms to carry out their metabolic processes. Composting is not favoured if the trash is too dry. For the purpose of getting rid of harmful germs, a high temperature is preferred. The organisms required to finish the composting process are nevertheless wiped off at temperatures over 75 °C. The procedure performs best at temperatures between 50 and 60 °C, with 60 °C being optimum. It is crucial to ensure that compost is properly aerated and that the amount of air is controlled. When oxygen levels drop, aerobes start to perish and are replaced by anaerobes. A large number of anaerobes will slow down the process, cause smells, and release methane, which is one of the most combustible substances in nature. The first large-scale composting facility, operated by Excel Industries Ltd. in Mumbai, processes 500 tonnes of MSW every day. In Vijaywada, another factory that can process 150 tonnes per day is in operation. The first large-scale composting facility, operated by Excel Industries Ltd. in Mumbai, processes 500 tonnes of MSW every day. In Vijaywada, another factory has been running with a daily capacity of 150 tonnes. End MSW compost is offered for sale for 3.50 INR (Chattopadhyay , 2009).

The MSW rules have established that only non-biodegradable inert trash and compost rejects should be disposed of in landfills; instead, biodegradable wastes should be managed via composting, vermicomposting, and other techniques. Composting is a preferred option in non-hazardous organic waste management and is part of many state SWM Plans. For instance, value added products from compost to be sent to the nearby rural farming lands (Gandhinagar Municipal Corporation 2020; HSPCB 2008; CPCB 2019) achieving circularity in solid waste management. MSW composition in India is on an average 50% compostable in nature. Paper and plastic waste increased by 280% and 1,200%, respectively, during 1996–2011, showing a rapid change in the physical structure of Indian MSW. Utilizing compostable and waste-to-energy (WtE) technologies were the best means to use compostable and combustible waste (Joshi and Ahmed 2016).

16.6 Organic Solid Waste Management as Manure

Manures has been long used as a fertilizer for agricultural systems and if well-managed in terms of storage can boost the crop production and thereby opening the realms of sustainability (Ndambi et al. 2019). Manures are known to improve soil's physical and chemical properties; increase NPK, CEC, Organic carbon, secondary nutrients like Ca, Mg; regulate pH, EC; reclaim soil by alleviating soil salinity; improve harvest index (Das et al. 2016; Wang et al. 2017; Prasad and Singh 1981; Bhatt et al. 2019; Dhar et al. 2015). As the waste enters the natural cyclic pathways via composting, not only will landfill burdens and their associated problems decrease, but also half of the nutritional needs of the plants can be met, resulting in higher crop yields (Singh and Longkumer 2018; Barrera and Hooda 2016) promoting both sustainable agriculture and sustainable development. But thorough waste segregation is the key. So, waste from one becomes a resource for another. For restoring the fertility of soil, organic fertilizer and manure use need to be encouraged. Under the scheme Paramparagat Krishi Vikas Yojana of the National Mission for Sustainable Agriculture, the Indian government promotes the use of organic manures (Press Information Bureau, Government of India, Ministry of Agriculture and Farmers Welfare 2017).

By stimulating enzymatic activities, organic amendments in conjunction with microflora increase mineralization and, therefore, soil aeration (Kang et al. 2021). The increased mineralization in soil improves soil fertility and crop productivity. Compost can be formed from various kinds of waste. Because of high organic matter and low concentrations of pollutants, MSW is used for composting, improving soil physic-chemical and biological characteristics. Bioconversion of organic waste fraction to compost is safe and straightforward answer to costly chemical treatment and fertilization for increasing productivity of salt-affected soil (Meena et al. 2016; Sundha et al. 2018).

There is often concern about heavy metal contamination in food production. Heavy metal can enter the organic materials of agricultural systems from pesticide spray, farm manures from municipal and industrial sludges which can slowly accumulate and cause severe harm to compost producers and handlers, plants and soil biota and end consumers. Through humification, although, organic compounds are made stable and pollution concentrations are reduced (Kumar and Kaushal 2015) and the manure produced can be fortified with additional treatments but strict enforcement of meeting the compost quality standards Fertiliser Control Order (FCO) (2009, 2013) and SWM Rules (2016) should be observed so that the nutrients can be cycled back to the environment for quality and environmentally sound food production in the realm of Sustainable Agriculture.

16.7 Sustainable Agriculture and Organic Farming

The health of our soil is critical to the success of our whole farming industry. Soil fertility refers to a soil's capacity to tolerate plant development while increasing output. To increase crop yields, they amended the soil with organic and inorganic fertilizers. Some soil fertility management strategies for improving agricultural productivity include the use of organic and inorganic fertilizers, legume crop rotation, and expanded germplasm. Soil fertility is critical for long-term agricultural productivity. As a result, organic farming plays a significant role, such as green manure processing. This verse influences soil physicochemically, and biologically, and about soil fertility and nutrient delivery to plants. (Saini et al. 2019). Soil fertility refers to the soil's capacity to deliver needed nutrients for biodiversity. Excessive application of costly chemical fertilizers significantly impacts the decrease of soil physicochemical and biological characteristics (NRCCA 2016).

Earles and Williams (2005) and Trivedi et al. (2016) advocated that for achieving sustainability in agriculture, soil structure, nutritional fertility and management, plant productivity, the healthy diversity and abundance of soil microbial population is an uncompromisable parameter and microbial community also serves as an essential early indicator of changing soil health even before physicochemical parameters show soil degradation.

Microbes in the soil produce polysaccharides and mucins that promote soil fertility and improve soil structure and health. A healthy plant should have a healthy rhizosphere rich with beneficial microorganisms. In contrast, optimum plant growth is challenging in soil contaminated with harmful microbes (Mendes et al. 2013). Microorganisms help to improve soil fertility by transforming available nutrients into usable forms. They consume and generate organic carbon in the Soil (Johns 2017). Organic carbon in the soil enhances soil fertility and water retention capacity. Microorganisms also help to reduce pollution in the environment (Lorenz and Lal 2014). Some of the aspects that are taken care of in Sustainable agriculture are as follows.

16.7.1 *Soil Tillage in Sustainable Agriculture*

Some of the technologies utilized to integrate amendments are tillage facilities for seeding or transplanting, fertilizer and lime as field-fixed inputs in residues and residues (Bermudez 2002). Deep ploughing promotes soil aeration and breaks up soil clumps to build healthy seed and root beds, improves water infiltration, boosts microbial activity and mineralization rate, and may smash through dense layers that hinder plant development and water movement. Tillage losses can influence the rate and severity of long-term decomposition of soil organic matter raise subsoil compaction problems, and hinder nutrient absorption and drainage (Naresh et al. 2013). Tillage losses can also cause bare pottery, hindering germination and water

infiltration. There are several advantages to low and no-tillage systems: the residue layer on the soil surface protects the soil from wind and water erosion; reduced tillage systems store more moisture in rainfed systems, which endure longer.

16.7.2 Cover Crops in Sustainable Agriculture

Legume cover crops' function in biological nitrogen determination and nutrient budgeting and *rhizobium* bacteria-infected legume cover crops can turn atmospheric nitrogen (N_2) into plant nitrate (NO_3^-) (MacKinnon et al. 2005). As a carbon source, nutrients are released into the soil solution as decaying soil microorganisms break down cover crop leftovers. Cover crops increase microbial activity and would enhance the breakdown of existing soil organic matter, allowing for establishing deep-rooted cover crops (Gougoulas et al. 2014). Temperature and humidity affect microbial activity (at low temperatures and during drought or waterlogging) and the location of cover crop residues on the soil surface. They coordinated the time of nutrient release from cover crops with agricultural productivity, crop demand, and the destiny of essential plant nutrients. Another effect of cover crops on agricultural soil physical characteristics internal (Patil and Lamnganbi 2018). Using cover crops to improve carbon and nutrient cycling results in a short-term improvement in soil physical properties. Many legumes are sensitive and can enhance nematode bug populations, as are cover crops' weed suppressive effects (Clark 2008).

16.7.3 Composts and Animal Manures in Sustainable Agriculture

Compost is used to regenerate soil and restore fertility, store carbon in the soil, and eliminate chemical inputs (fertilisers, pesticides, and fuel), all of which are environmentally friendly and lead to lower processing costs (Diacono and Montemurro 2011). The most suitable composting technology is chosen based on farm assessments (volumes of materials to be composted, matrix sort and supply locations, machinery/facilities already in place), preliminary environmental and economic sustainability assessments using Life Cycle Assessment, Life Cycle Costing, and Energy Analysis methodologies (Pergola et al. 2018). In addition, the global increase in agricultural livestock units is causing an increase in animal waste. Animal waste cannot be applied directly to soil as a soil amendment therefore biological stabilization is a better approach for use. Animal dung may be utilized as a high-value fertilizer for crops and grasslands if distributed correctly and at the right time to minimize losses and maximize nutrient delivery per crop need (Goss et al. 2013).

16.7.4 Crop Rotation in Sustainable Agriculture

This is the process whereby a crop is continuously moved from one place to a different place, principally for ecological reasons. The following are the reasons for crop rotation: Interrupting pest-host interactions, preventing the accumulation of rodents, weeds, and illnesses (Anaya 1999). Various crops (tomatoes, eggplants, peppers and potatoes) share pests and diseases, repeated farming in the same site can lead to pest population growth. Crop rotation to achieve the most fertilizer supplies and to distribute nutrient demand throughout the soil. Multi-year crop rotations also include fallow cycles and perennial cover crop rotations. Areas kept fallow and planted with permanent cover crops (e.g., perennial ryegrass) permit the soil to stay undisturbed and the aggregation processes to continue uninterrupted (Tadesse et al. 2021).

16.8 Basics of Bioelectromagnetic Energy and its Practical Application

Agriculture is open to experimentation from all fields to enhance production for fulfilling the needs of the ever-growing population in a sustainable way. One such approach is Sustainable Yogic Agriculture (SYA) which involves the utilization of basic techniques of sustainable agriculture along with immaterial energy (Brahma Kumaris 2009). In secular terms, immaterial means something that is not perceived with our five senses at first, a dimension that involves vibrational energy, consciousness, ether, sentience, intelligence, and/or electromagnetic waves or sound frequencies. Various disciplines involved in understanding these processes includes quantum mechanics, consciousness and phenomenological studies, bioelectromagnetic, chronobiology, sonochemistry, neuroscience and transpersonal psychology. The concept of “life energy” is prevalent across many cultures from a long time. It is called by various names like Qi (chi), prana, universal fluid, cosmic aether, archaicus, animal magnetism etc. Very less work is done to understand such phenomena, in the meantime, technological advances from here facilitate our daily lives in an increasing number of ways. Various electrical and electromagnetic devices that modify and improve bioenergy fields are employed for better health in energy medicine practices (Wright 2021).

At fundamental levels, atoms and molecules vibrate at various frequencies. It is not that only high vibrations have impact, forces which are at lower frequencies if resonate with something having the same frequency can have powerful impacts. Takoma Bridge incident of 1940 is an infamous example of impact of resonance. The closest we have come to understand the immaterial energy is comparing it with electrical or electromagnetic signals (Rosch 2009). Ravitz (1982) showed the behaviour of the energy to be similar to electrical activity which he studied as life (L) fields, the energy fields encompassing living beings. He reported that mobilization in galvanometer was caused due to stimulus electrical in nature, and emotional activity

evoked stimuli of the same energy. So, he concluded that “Emotions can be equated with energy” and this energy could be influenced by mental and emotional states of mind. Quantum physics suggest we are *electromagnetic* human beings. Our thoughts radiate out from us like broadcast signals like vibrations (Lipton 2012).

The magnetic property of the bioelectromagnetic energy can be inferred by the experiment by laboratory experiments on Wheat and Chickpea at G.B. Pant University of Agriculture and Technology (Pandey et al. 2015). Seeds were exposed to a magnetic field of 100-250mT (milli Tesla) in steps of 50 mT for 1 to 4 h, in steps of 1 h for all field strengths and other batch was of Seeds exposed to metaphysical immaterial energy treatment for 1 to 4 h, in steps of 1 h by trained Rajyoga meditation practitioners. For wheat, slight increase in germination percentage in magnetic field (250 mT) as well as metaphysical energy treatments. Mean root length, shoot length and seedling vigor index was significantly higher in metaphysical energy treatment as compared to magnetic field treatment. Control showed lower root length, shoot length and seedling vigor index. Chickpea Seeds at 150, 250 mT for 1 h and metaphysical energy exposure were at par with all other treatment combinations except 200 and 250 mT for 3 to 4 h. Root length, shoot length and seedling vigor was significantly higher due to metaphysical energy exposed to seeds for 4 h compared with remaining treatment combinations.

These vibrations are subtle in nature and can be measured using electrophotonic emissions/gas discharge visualization (GDV). It is nonintrusive technique that helps capture the physiological and psychoemotional status of a person and the functional status. The sensor used is sensitive enough to catch the change in conductivity of the space in the presence of emotional people which might be due to formation areas of decreased entropy in space, or, associated with the buildup of a negative magnetic charge manifesting in the environment and identify deviations from the normal functional state at early stages and in real time. Biofield aura intensity and related characteristics are dependent on health and emotional state of mind (Kostyuk et al. 2011; Suru Krilian Photography Centre 2022).

Earlier scientists acknowledged the immaterial layer of universe and some have worked and have been working to bring to surface this phenomenon and its impact in the practical physical world. But, in twentieth century many electrotherapeutic energy devices made worthless claims and further doctors viewed anything and everything related to energy medicine as fraudulent which negatively affected the acceptance of immaterial phenomena (Rosch 2009). Otherwise, John van Neuman, mathematician and pioneer in quantum mechanics stated that “measurement chain ends only when the knowledge of the measurement is registered by an extra physical factor, that is the mind (not the brain)”. Radin et al. (2012) using his double slit experiment significantly proved that mind actively participates in manifested reality. He showed in his experiments that mental observation of light beam causes interference and collapses the wave function.

In the medical field, Alda et al. (2016) experimented with a group of 40 individuals divided into meditators and non-meditators showed that in comparison with non-meditators, expert meditators’ telomere lengths were significantly longer and their percentage of shorter telomeres was lower. Chromosomal protection, end to end

fusion and senescence of cells from degradation from nuclease is extended by degree of intactness of telomeres. Telomeres are reported to decrease in length with age and can help in early detection of diseases like hypertension, atherosclerosis, type 2 diabetes mellitus, cancer mortality, cardiovascular disease, cognitive decline and dementia.

Luders (2014) reviewed and interpreted 3 studies. Data suggested that meditation practice may slow, stall, or even reverse age-related brain degeneration. García-Campayo et al. (2018) conducted an experimental study on 34 participants divided equally into treatment and control suggested that in distinct regions of genomes that are responsible for CpG methylation have shown loss in long-term meditation practitioners. Epigenetic modulations can be impacted by mindfulness which in turn affect biological processes related to lipid or insulin metabolism. By functional analysis, at molecular level, mindfulness practices have crucial roles in signaling pathways of TNF and NF- κ B which is also linked to organizing cellular resistance to invading pathogens and enhance immunity (Bonizzi and Karin 2004).

Back in 1999, Harris et al. experimented to assess the effect of positive healing at a distance in a human clinical trial (1013 subjects). Patients that were admitted to a coronary care unit, a team of intercessors on a randomized basis prayed for patients. Severity score when compared with control group was statistically lower in the patients who were subjected to remote, intercessory prayer.

With living systems of plants in 1963, Grad established that plants are receptive to energy; healing energy improved growth, and negative energy stunted it. Backster (2003) demonstrated plants and other living organisms respond to human intention. Emoto (2004) and Radin et al. (2006) have shown impact of thoughts, intentions and words on water crystals, how positive intentions helps form well defined structure in water molecules and negative did the opposite. Oschman (2004) proposed that a photon of electromagnetic energy works the same way as a single antigen, hormone, pheromone, growth factor, neurotransmitter molecule which can provoke a series of signals at cellular levels that affect biological processes by initiating or inhibiting or even accelerating them. Sum of molecular events on the cell surface can in turn activate enzymatic activities which can catalyse and accelerate biochemical processes. Benveniste (2004) proposed that the theory of trial-error method of reception of ligand molecule and receptor can possibly be also that ligand emits an electromagnetic signal with a frequency resonating to the receptor's molecules that causes them to co-resonate and activate the same intracellular responses which can further explain the anomalies that are present despite of the quantitative structure-activity relationship theory for molecular signalling. Some processes are sensitive to these fields and some are not. Electromagnetism needs to be studied and tested more rigorously in order to establish which reactions are sensitive to these fields.

16.9 Potential of Yogic Farming and Related Research Work

Many additive techniques can be applied along with Organic farming to increase the production in fields in a sustainable way (Fig. 16.2). Adding to the fortification of quality of compost, yogic farming, an additive to organic farming is a novel yet ancient approach that harvests the potential of human mind. It refers to the agricultural practices which involve bio-organic and natural inputs along with application of the power of positive thinking and pure feelings, which are immaterial in nature (Brahma Kumaris 2009). Now, subtle, the immaterial counterpart of agroecology in advanced ecology is revealing itself and it needs to be considered in the farming practices and it is not a new concept. Ancient, traditional Indian farming practices well understood and incorporated the material and immaterial science (Wright 2021). Thought based meditation practice of Rajyoga meditation, practitioners create powerful pulses of vibrations by gently guiding the thoughts to a higher level of positive divine thoughts and direct them towards the farming setup—the seeds, water, and/or crops (Brahma Kumaris 2009). Experimental studies reveal that exposing the farming set up to thought based vibration energy helps enhance soil microbiological activity for microbes like Rhizobium, Azotobacter, Azospirillum and PSB under Organic + Yogic Farming module and crop quality in terms of higher values of germination %, vigour, higher dehydrogenase activity and low electrical conductivity and pest damage control when compared with control (Sharma et al. 2014; Pandey et al. 2015; Ndiritu et al. 2016; Yadav and Vats 2021).

This thought-based energy can be applied during the stages of bio conversion of organic waste and crop production to maximize the quality of compost and crop yield.

According to the Ministry of Agriculture and Farmers Welfare, their target is to increase the farmers' income by 2022 twofold, and for this purpose, Agriculture Minister Radha Mohan Singh pointed out that techniques additive to Organic farming, the Sustainable Yogic farming using Rajyoga meditation which works best under organic farming as research shows is being encouraged under the Parampragat Krishi Vikas Yojana (PKVY) which will also have positive societal benefits. (The Economic times 2016; Yadav 2017).

The outcome of some more research works undertaken in agriculture incorporating the bioelectromagnetic energy are tabulated and given in Table 16.1.

A preliminary research, pot scale study was conducted in urban setting of Sector 20, Gurugram with an aim to assess the possibility of using locally available organic wastes for manure production and aiding in SWM and crop production in space constrained urban areas (Fig. 16.3). The locally available waste selected was keeping in mind COVID-19 restrictions. Organic waste from household (fruit and vegetable peels) and paper waste was locally available and is well suited for composting.

The bioelectromagnetic treatment given is the energy from thought-based meditation. The experimental set up consisted treatments on the basis of distance—2 m, 4 m, 6 m and Control. Three types of compost compositions—Paper waste compost,

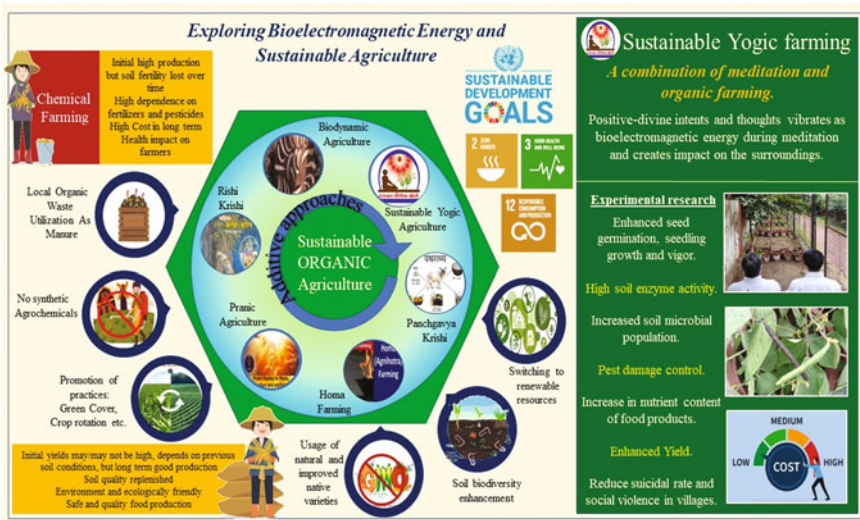


Fig. 16.2 The additive approaches that are being explored under sustainable organic agriculture and sustainable yogic farming incorporating bioelectromagnetic energy to enhance the agro-production in a sustainable way

Kitchen waste compost and Mix waste compost was produced. Keeping the similar environmental conditions, the similar treatment of average 30 min thought-based meditation energy was given by meditation practitioner. The sampling of compost soil was done at 30, 60, 75 days. Highlight of study was that porosity %, organic carbon and organic matter percentage is higher among the compost exposed to thought based energy supporting the use of SYA in organic manure production. The compost was further applied as manure for crop production of Moong beans. The treatment with organic + yogic showed higher germination percentage when compared to inorganic chemical treatment and only organic treatment.

16.9.1 Conclusion

It is one of the basic principles in ecology that waste that we generate will not disappear because cyclic pathways in the natural environment rearrange, cycle and recycle all materials. This circular pattern has been disrupted when we talk about the mismanaged systems of waste management and it is not a problem to ignore anymore. Bioconversion of locally available organic waste to manures for sustainable SWM will reduce a huge burden from landfills. This will aid in nutrient management of agricultural soil and crop production and their quality enhancement by incorporating Sustainable Yogic Agriculture (SYA) approach seems to be a holistic solution harnessing the combined potential of human mind and natural system for agricultural sustainability enabling the prosperity of ecology, economy and society.

Table 16.1 Research works conducted in agriculture using bioelectromagnetic energy

Treatment	Parameter studied	Plant studied	Outcome	References
Pranic energy application	Growth, yield and nutritional content	Finger millet	Plant growth and production parameters were higher as compared to control-plant height, productive tillers count, number of panicles, fingers count and grain yield were 26%, 35%, 54%, 13% and 44% respectively higher statistically (at $p < 0.05$) Protein content in straw and grain was 4.38% and 6.13% compared to control than control with 3.5 and 4.75% Nitrogen and zinc content was also higher in millet grain was higher in treatment	Poornima et al. (2021)
Yogic farming through Brahma Kumaris raja yoga meditation	Crop growth performance	Wheat and groundnut	Seeds exposed to meditative energy showed enhanced germination, seedling growth and vigor. The quality of groundnut and wheat improved, along with increased soil microbial population. Organic + meditation gave comparable results to chemical inputs	Pandey et al. (2015)

(continued)

Table 16.1 (continued)

Treatment	Parameter studied	Plant studied	Outcome	References
Yogic farming (Raja Yoga Meditation)	Germination and seedling vigor + enzymatic activity	Chickpea	Significantly increased root and shoot length, seedling vigor index and dry weight, enzymatic activity of dehydrogenase over control. Electrical conductivity of seed leachate of treated seed significantly decreased over control	Sharma et al. (2014)
Pranic treatment	Plant growth parameters	Cluster beans	For pranic group, significant increase in vegetative growth and higher flowering and yield	Prasad and Jois (2021)
Meditation listening to shamanic double drumming	Germination and growth of plants	Wheat and pea	The seed germination rate exposed to treated water with stimulating intent was about 8.5% higher in treatments at significant p values. Inhibiting intent experiment with wheat showed 4.2% lower germinate rate compared to control seeds. Mass of seedlings was lower in the inhibitory intent	Haid and Huprikar (2001)
Buddhist meditation	Growth of plant due to intentionally treated water and seeds	<i>Arabidopsis thaliana</i>	Seeds grown with intentionally treated water had decreased length of hypocotyl and increase in anthocyanin and chlorophyll suggestive of enhanced photomorphogenic growth	Shiah et al. (2021)

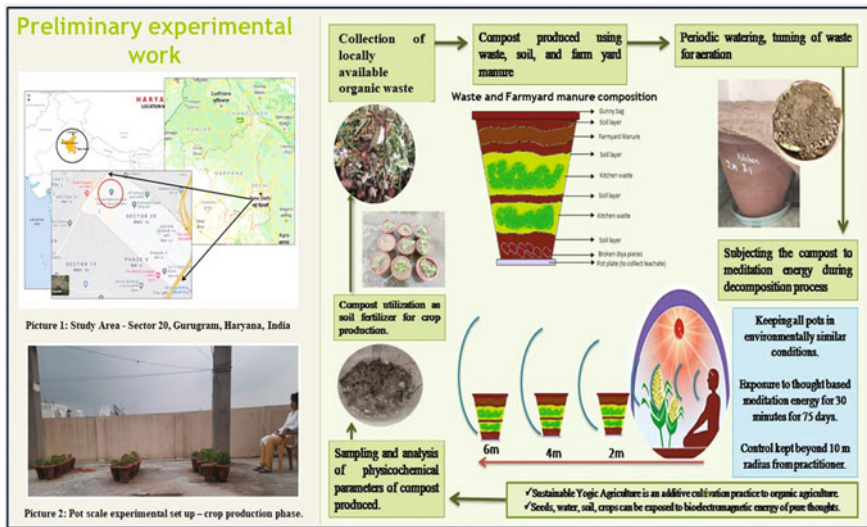


Fig. 16.3 Pot scale study conducted in Sector 20, gurugram showing study area and experimental set up

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Chapter 17

Production of Sustainable Building Products Using Micronized Biomass Silica from Rice Husk



S. Thirumalini, T. Shanmuga Priya, Mirijam Vrabec,
and Gayathri Chandran

Abstract Micronized biomass silica (MBS) is an agricultural waste obtained by burning and grinding rice husk in a controlled atmosphere to nano scale. The present chapter focuses primarily on the performance assessment of MBS as an ingredient in production of Fly ash brick and paver blocks. The research was carried in two stages. Casting and compressive strength tests of mortar mixtures with variable MBS content in the range of 0, 5, 10, 15, and 20% by weight of cement were conducted in the first step. The optimum percentage of MBS substitution in mortar was found to be 15% and the microstructural properties of hydrated cement mortar mixes using SEM and EDAX shows that the addition of MBS to the hydrated cement mortar mixes resulted in the formation of CSH gels and CaCO_3 crystals, which may be credited to improved strength. In second stage, fly ash bricks and paver blocks were casted by replacing fly ash and cement respectively with varying percentage of MBS (0, 10, 20, 30, 40 & 50%). From the compressive strength test it was found out that optimum replacement of MBS was 15% for fly ash bricks and paver blocks. Efflorescence was not observed in fly ash bricks at all proportions of MBS. When compared to control specimens, MBS blended fly ash brick and paver block specimens showed a significant resistance to water penetration. The present study indicates the feasibility of utilizing MBS as a raw material in production of sustainable building materials and products.

Keywords Micronized biomass silica · Cement mortar · Fly ash bricks · Paver blocks

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17.1 Introduction

Concrete is recognized amongst the most important components incorporated in construction sectors. Due to the various mechanical and physical properties it possesses, it is extensively used in construction activities. It is true that its application and usage are touching skies but simultaneously its ill effects on the environment are also touching skies. Amongst the major issues related to concrete structures are carbon dioxide emissions due to large amount of cement usage and cracks-especially micro cracks (Worrell et al. 2001; Coronelli and Gambarova 2004). More than 7% of greenhouse effects are due to carbon dioxide emissions from the extensive cement content in concrete (Turner and Collins 2013).

Global warming is speeding every day and it is consuming every single unit of the atmosphere. It has been caused by extreme increase in greenhouse gas emission, which has now created a blanket of pollution around the planet. Air pollutants are dangerously settling in atmosphere. The more they increase, more is the threat created for each individual's health. Construction industry is one of the main causes of this pollution. Construction companies are the source of many wastes which are untreatable. Additionally, the population is growing and the economy is developing quickly, which has increased the production of industrial and agricultural waste products such fly ash, GGBS, rice husk ash, and silica fume, whose disposal is becoming more difficult. Several research projects are currently being conducted on the use of waste in concrete (Corinaldesi et al. 2002; Mehta 2002; Naik and Moriconi 2005).

Rice is the main source of food in India. This causes a continuous demand requiring production at bulk. While processing the produce for the market, we get a byproduct called rice husk, which is the skin of rice grain. This husk is disposed of without considering the fact that it can be used for our own benefits. An agricultural waste product derived from the processing of paddy is rice husk. Of the 590 million tonnes of paddy produced worldwide, it makes about 20% (Ewais et al. 2014). The rice husk is utilised in India to make partition boards, feed animals, and other things that are considered trash (Gidde and Jivani 2007). Amorphous silica, which is utilized as a mineral additive in concrete, is found in high concentrations in rice husk ash (RHA), which is produced by controlled burning (De Sensale et al. 2008; Dellaet al. 2002). More CSH gel is created when this amorphous silica interacts with hydration products. As a result, concrete's mechanical and durable qualities are improved (James and Subba Rao 1986). It is known that adding RHA to concrete increases strength, decreases permeability, and provides corrosion resistance (Saraswathy and Song 2007). Additionally, numerous studies demonstrate that recycled aggregate concrete with rice husk has superior mechanical qualities to concrete without RHA (Tangchirapat et al. 2008; Akinkulolere 2013).

The performance of regular concrete with RHA has been the subject of numerous studies; however, research on concrete containing micronized biomass silica (MBS) is few. The MBS is made by carefully burning rice husk at a controlled temperature (500–600 C) in a rotary furnace and then grinding it for a short period of time in a jar

mill to minimize the particle size (i.e. micronized). The increased silica concentration and finer particles are the primary distinction between RHA and MBS. In order to produce high performance concrete, RHA and silica fume have been employed in place of cement (Zhang and Malhotra 1996; Appa Rao 2003). Secondary calcium silicate hydrates (CSH) were produced as a result of MBS's pozzolanic reaction with cement hydrates, and MBS's characteristics, which have the ability to lessen the intensity of Ca(OH)_2 , also improved.

Research carried by Suraya Hani et al. (2015) involved cement replacement with MBS at percentages ranging from 0 to 20%. It was noted that up to 12% of MBS, concrete showed higher strength and lower water permeability when compared with control concrete. Another work by Prameetthaa et al. (2015) incorporated replacement of cement with MBS at 0, 4, 8 and 12% in mortar. An increase in CSH gel formation along with a potential reduction in the intensity of Ca(OH)_2 was observed. Optimum results were obtained at 8% MBS on grounds off mechanical and durability properties. The research work carried by Ambily et al. (2015) involved the use of total replacement of ground granulated blast furnace slag against ordinary Portland cement and MBS was replaced at 0, 10, 20, and 30% by volume. Various mechanical and physical properties were analyzed and it was found that 10% replacement of MBS showed a considerable higher compressive strength. The current study has been undertaken to determine the mechanical and durability characteristics of cement mortar with partial replacement of cement and various percentages of MBS, as well as to assess its sustainability, a study has been undertaken.

17.1.1 Objectives

The preeminent goal of this research was to determine the ideal amount of MBS to use in the cement-mortar paste and to assess the partial substitution of cement with MBS in cement-mortar. The project's goals are outlined below in brief: To study the characteristics of Micronized Biomass Silica (MBS) as an effective replacement of cement in cement paste and mortar.

- To study characteristics of Micronized Biomass Silica as an effective substitute of cement in cement paste and mortar.
- To perform preliminary test on cement replaced with various proportions of MBS to analyze various properties such as its specific gravity, setting time and flowability.
- To design mix proportions with varying contents of MBS as a replacement of cement.
- To study and analyze strength and durability of the rendered mix designs and to compare with the control mix (Traditional cement mortar).
- To perform a cost-benefit analysis to deduce the economic feasibility of using MBS as a cementitious material.

- To undertake an environmental impact assessment to understand the impacts caused by replacing MBS in cement mortar.
- To derive an optimum replacement percentage of MBS to obtain a sustainable cement mortar.

17.2 Materials and Methods

OPC of 53 Grade was used in accordance with IS 12269:2007 and the specific gravity of cement was found to be 3.15. MBS used in this study was obtained from the Astrra Chemicals, a chemical manufacturing company in Chennai. The specific gravity of MBS used is 2.2. MBS has a pozzolanic activity index of 85.05% with a surface area of 20.83 m²/g. The chemical composition of cement and MBS are presented in Table 17.1 (Fig. 17.1).

A low calcium Class F fly ash (confirming with ASTM C311) was used and its chemical compositions are given in Table 17.1. Locally accessible natural sand with a maximum grain size of 4.75 mm that complied with Zone II was used. According to IS383-2016, the bulk density of fine aggregate was found to be 1675 kg/m³ with water absorption of 1.8%, and the specific gravity was 2.65 and fineness modulus

Fig. 17.1 Micronized biomass silica



Table 17.1 Chemical composition (%) of cement and MBS

Compound	Cement	MBS	Flyash
SiO ₂	15.55	95.25	48
Al ₂ O ₃	8.40	0.27	24.3
Fe ₂ O ₃	5.87	0.33	15.6
SO ₃	3.12	0.21	0.4
CaO	62.44	0.81	3.2
Na ₂ O	0.16	—	0.8
K ₂ O	1.27	2.26	—

was 2.36. For the preparation of the mortar, fresh, drinkable water that was devoid of organic and acidic materials was employed.

17.2.1 Mixture Proportions for Mortar Mixes

The effects on different percentages of MBS (partial replacement by weight of cement) were evaluated, expecting developments in the strength, durability and microstructural properties. A total of 5 batches of mortar samples were cast. Ordinary portland cement, river sand, MBS and Potable water was used. Mix design was done as per the procedure mentioned in ASTM C311. The mix proportion in kg/m^3 is presented in Table 17.2 (Fig. 17.2).

To maintain a consistent mixing temperature, all the ingredients were held at ambient temperature ($27 \text{ }^\circ\text{C} \pm 2 \text{ }^\circ\text{C}$). All the components were mixed using a paddle mortar mixer. The process of combining the ingredients took around 7 minutes. After combining all the ingredients, the mortar was layered three times in the mould and properly crushed using a vibrating table. The casted cubes are shown in Fig. 17.2.

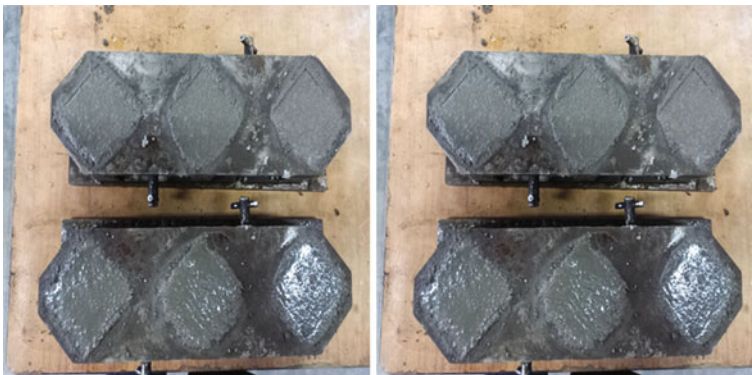


Fig. 17.2 Cubes for compressive strength test

Table 17.2 Mix proportion

Mix ID	Percentage replacement		Quantity of materials (grams)			
	Cement	MBS	Cement	MBS	Sand	Water
M1	100	0	500	0	1375	242
M2	95	5	450	50	1375	242
M3	90	10	400	100	1375	242
M4	85	15	350	150	1375	242
M5	80	20	300	100	1375	242

Tap water was used for curing these specimens for 28 days at room temperature ($27\text{ }^{\circ}\text{C} \pm 2\text{ }^{\circ}\text{C}$). The specimens were immersed in water throughout the curing process.

17.2.2 Mixture Proportions for Flyash Bricks

The various raw materials that constitute the fly-ash bricks, along with their proportions used in the industry are: Fly-ash—50%, Lime—15%, Cement—10%, Crusher dust—25% and 150–200 ml of water is used per kilogram of the raw mixture.

Fly-ash bricks were cast with varying proportions of MBS replacing flyash (0, 10, 20, 30, 40 & 50%). The raw materials were mixed and blended in a miller drum. A fly-ash brick weight about 3.3 kgs. Therefore, a batch of 20 kg of this mixture was taken for 6 bricks (each mix) to get samples with desired proportion of MBS (Fig. 17.3).



Fig. 17.3 Flyash bricks preparation **a** Mixing of raw materials **b** Bricks emerging from hydraulic press **c** Bricks after curing



Fig. 17.4 Paver blocks preparation

17.2.3 Mixture Proportions for Paver Blocks

Paver blocks of M-35 grade were used in this study. The various raw materials that constitute the fly-ash bricks, along with the mix proportions used in the industry are:

Cement: Fine aggregates: Coarse aggregates—1:2.5:3.25.

Water-cement ratio (w/c) - 0.4

Samples of M-35 grade paver blocks were cast with varying proportions of MBS replacing Cement (0, 10, 20, 30, 40 and 50%). The raw materials were mixed and blended in a mixing drum. A paver block weight about 5.4 kgs. Therefore, a batch of 33.75 kg of this mixture was taken for 6 blocks (each mix) to get samples with desired proportion of MBS. Fig. 17.4 shows the preparation steps of paver blocks.

17.2.4 Testing Procedures

In order to evaluate the compressive strength in accordance with Codal requirements, mortar specimens of $50 \times 50 \times 50$ mm, fly ash bricks, and paver blocks were employed. There were six samples cast for each combination. The strength was assessed after 7 and 28 days of curing. With a continuous load rate of 2.5 kN applied to the cubes, the test was carried out using an AIMIL compression testing machine. The produced raster scan picture for SEM can target resolutions greater than 1 nm. The sample underwent a compressive strength test and was then ground into smaller pieces before being evaluated for SEM examination. The samples were then cleaned and dried in an oven at a temperature of about 96 °C. The samples were made into a powder and put through a sieve with a mesh size of 75 microns. Additionally, the uncoated sieved samples were maintained safely over a glass plate that had been coated in Canada Balsam. Following that, the samples are examined using an LEICA microscope and visualised using QWIN software.

Fig. 17.5 Water absorption test



According to the steps outlined in IS3495 (Part 2), the fly ash bricks and paver blocks were evaluated for total water absorption as shown in Fig. 17.5. To achieve basically constant mass, specimens were dried in a vented oven at a temperature of 105 to 115 °C. After the samples had reached room temperature, its weight (M_1) was determined. The specimens were fully dry before being submerged in sterile water for 24 h at a temperature of 27 °C. The specimens were taken out, and any water residue was cleaned with a moist cloth. Three minutes after being removed from the water, the specimens were weighed, and their weight (M_2) was recorded. Following is the formula for water absorption, expressed as a percentage of mass, following a 24-hour immersion in cold water:

$$\text{Water absorption} = [(M_2 - M_1) / M_1] * 100$$

17.3 Result and Discussions

17.3.1 Compressive Strength Results—Mortar Samples

Compressive strength values of MBS incorporated mortar samples shown in Fig. 17.6. Compressive strength values showed an increase from 5 to 15% replacement of MBS. Decrease in strength was observed with 20% replacement of cement with MBS.

From Fig. 17.6, the compressive strength with 5% MBS showed an increase of 0.56% at 28th day. With 10% MBS addition an increase of 1.09% was observed with respect to reference sample. Compressive strength grows with 15% percentage of MBS addition, after which it rapidly falls. A percentage of 1.45% was observed for this percentage replacement. With 20% addition a percentage decrease of 1.57%

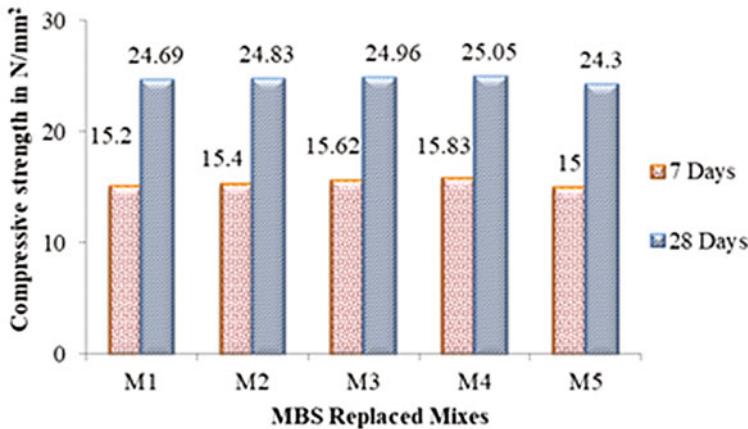


Fig. 17.6 Compressive strength results—mortar samples

was observed. The pozzolanic reaction of MBS is what causes the gain in strength. The MBS contains a lot of silica and it transforms calcium hydroxide into C-S-H molecules when combined which are in charge of developing strength. The constrictive after a certain point, strength decreases as less water is due to the MBS's high water absorption capacity, available for the reaction ability.

17.3.2 Compressive Strength Results—Fly Ash Bricks

Average strength values of MBS incorporated fly ash bricks at 21 days are shown in Fig. 17.7. Compressive strength values showed an increase from 0 (FAB1) to 40% (FAB5) replacement of MBS. With 10, 20, 30, 40% MBS addition a percentage increase of 0.2%, 0.48%, 1.4%, and 2.7% respectively. Decrease in strength was observed with 50% replacement incorporation of MBS. The percentage decrease was around 1.6%. Optimum strength value was observed at 40% of MBS in fly ash brick mix. Up until 40% replacement, the compressive strength grows as the percentage of MBS does, after which it rapidly falls.

17.3.3 Compressive Strength Results—Paver Blocks

Samples of paver blocks had their compressive strength tested after seven and twenty-eight days. For each mix, three samples were cast and evaluated over the course of seven and twenty-eight days. The average of these readings was then taken and is shown in Tables 17.3 and 17.4 below.

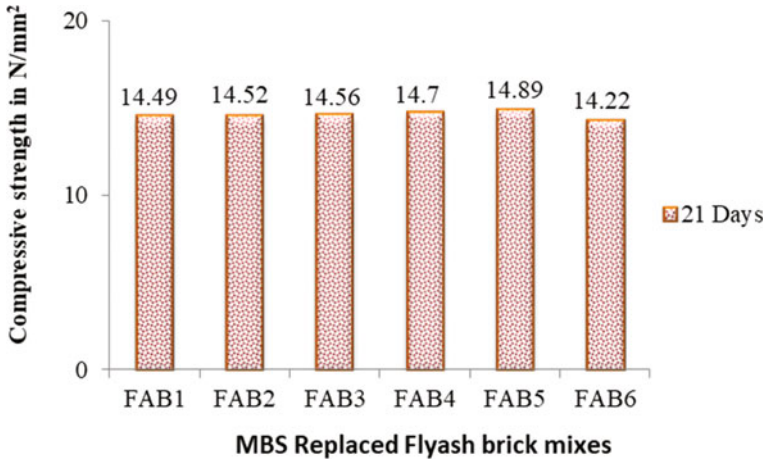


Fig. 17.7 Compressive strength results—fly ash bricks

Table 17.3 Compressive strength for paver blocks at 7 days

eMix Id	Sectional area (mm ²)	Average load at failure (kN)	Average strength (N/mm ²)
PVB1	30,000	643.2	21.44
PVB2	30,000	673.5	22.45
PVB3	30,000	675	22.5
PVB4	30,000	678	22.6
PVB5	30,000	696	23.2
PVB6	30,000	538.89	21.3

Table 17.4 Compressive strength for paver blocks at 28 days

Mix Id	Sectional area (mm ²)	Average load at failure (kN)	Average strength (N/mm ²)
PVB1	30,000	1051.5	35.05
PVB2	30,000	1125	37.5
PVB3	30,000	1128.3	37.61
PVB4	30,000	1129.5	37.65
PVB5	30,000	1148.1	38.27
PVB6	30,000	1050	35

Average strength values of MBS incorporated paver brick at 7 and 28 days is shown in Fig. 17.8. For both curing age the compressive strength values showed an increase with 0% (PAB1) to 40% (PAB4) replacement of MBS. With 10% MBS the percentage increase in strength for 7 and 28th day was 4.7% and 6.9% respectively. For 20%

replacement it was 4.9% and 7.3% respectively. Similarly for 30% replacement it was 5.4% and 7.4% respectively. Maximum hike in strength was observed at 40% replacement. A dip in strength was observed with 50% replacement incorporation of MBS for both curing age. Optimum strength value was observed at 40% of MBS in paver brick mix. Up-to 40% substitution, the compressive strength grows as the percentage of MBS does, after which it rapidly falls till 50% replacement.

17.3.4 Water Absorption Results—Fly Ash Bricks

The results of all the mixtures water absorption tests were displayed in Table 17.5. It was shown that when MBS was substituted for fly ash brick at various percentages, as percentage replacement increased water absorption decreased. A percentage decrease of 1.29%, 20.1, 33.7%, 24.02%, 25.3% was observed for 10, 20, 30, 40 and 50% MBS addition respectively. Lowest water absorption percentage was observed in 30% MBS in fly ash brick. The optimum water absorption percentage was 40%. The existence of finer MBS particles and its pozzolanic feature, which led to a discontinuous pore structure, can be a cause of this positive result.

17.3.5 Water Absorption—Paver Block

Table 17.6 displays the mixes' water absorption capabilities. When MBS is added, which has PVB4's mix having the lowest water absorption, the water absorption decreases? 20% MBS in paver blocks had the lowest recorded water absorption percentage. The recommended replacement rate was 20%.

Table 17.5 Water absorption results—fly ash bricks

Mix ID	Dry weight (kg)	Wet weight (kg)	% Water absorbed
FAB1	3.303	3.813	15.4
FAB2	2.817	3.246	15.2
FAB3	2.970	3.335	12.3
FAB4	3.213	3.540	10.2
FAB5	2.825	3.156	11.7
FAB6	2.713	3.026	11.5

Table 17.6 Water absorption results—paver blocks

Mix ID	Dry weight (kg)	Wet weight (kg)	% Water absorbed
PVB1	5.327	5.521	3.64
PVB2	4.280	4.425	3.38
PVB3	4.627	4.764	2.96
PVB4	4.962	5.062	2.01
PVB5	5.035	5.231	4.90
PVB6	5.125	5.426	5.87

17.3.6 Micro-structural Analysis—Mortar Mix

The conventional and 15% replacement of cement by MBS replacement mix were studied through micro-structural analysis and compared with CC to evaluate the texture and effect of MBS in cement mortar. Fig. 17.8a shows the microstructural arrangement of CC, being a uniformly distributed arrangement of materials in absence of any major cracks visually observable. Unreacted particles are visible on the surface of the surface which shows incomplete pozzolanic reaction. Moreover the presence of small amount of Si is reflected in the EDS analysis.

Figure 17.8b shows the micro-structural matrix for MBS addition mortar mix which shows finished pozzolanic reactions. The SEM image shows a dense matrix due to enhancement in the formation of CSH/CASH gels which is also confirmed by the EDS analysis which shows large amounts of Si and Ca governing the high strength of the mix. Lastly MBS is accounted for higher amounts of Si and Al, thus promoting a high strength mix with desirable properties.

17.4 Conclusions

The below mentioned outcomes were achieved from the above described results

- Cement replacement with MBS in mortar increased compressive strength by up to 15%.
- The optimum replacement of MBS was found to be 20% for fly ash bricks and paver blocks.
- Micro-structural analysis of mortar mixes showed the effect of pozzolanic nature of MBS in increasing the overall performance.
- The MBS addition products can be used for buildings as a sustainable and economical way for disposal of rice husk waste.
- MBS mortar can be applied other primarily places where cement mortar is used leads to paving a way for a cleaner and greener construction.

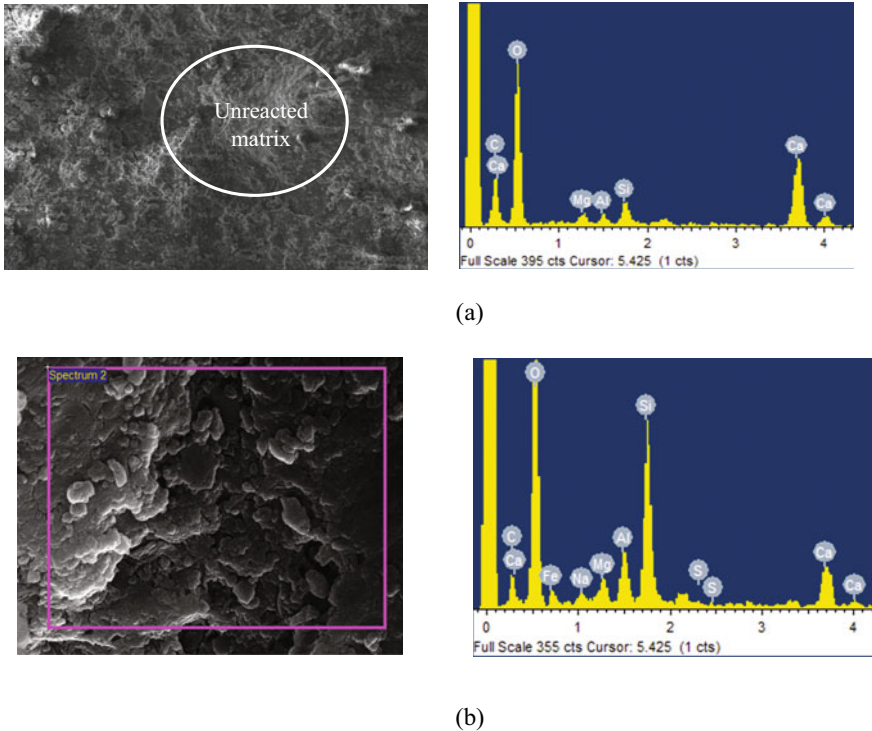


Fig. 17.8 SEM & EDS **a** CC, **b** MBS mix

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