

Groundwater Organic Pollution Remediation Using Microbial Techniques: A Study of Palm Oil Industrial Areas

K. R. Muthusamy¹

¹ Independent Researcher, Malaysia

Correspondence: K. R. Muthusamy, Independent Researcher, Malaysia.

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Abstract

Groundwater contamination from palm oil mill effluent (POME) poses significant environmental challenges in Malaysia. This study highlights microbial remediation as a sustainable solution, utilizing indigenous and engineered microbial communities to degrade organic pollutants through key enzymes like lipases and catechol dioxygenases. Techniques such as biostimulation, bioaugmentation, and anaerobic-aerobic systems are examined, along with optimization strategies like nutrient supplementation and pH control. Case studies demonstrate significant reductions in BOD, COD, and phenolic levels, showcasing microbial remediation's potential for sustainable groundwater management in palm oil industrial areas.

Keywords: groundwater contamination, POME, microbial remediation, biostimulation, bioaugmentation

1. Introduction

Groundwater contamination is one of the most critical environmental issues globally, particularly in regions with rapid industrial development. In Malaysia, the palm oil industry represents a key economic driver, contributing significantly to global palm oil production. However, its environmental impact cannot be overlooked, especially concerning groundwater pollution. Among the most significant threats is contamination from Palm Oil Mill Effluent (POME), a highly organic waste product generated during the extraction process. POME contains large quantities of organic compounds such as fatty acids, carbohydrates, and phenolic compounds, as well as suspended solids, which collectively pose a severe risk to groundwater quality. When POME is discharged improperly or accidentally leaks into the soil, it percolates down to aquifers, compromising the quality of groundwater—a critical resource for domestic, agricultural, and industrial use in many parts of Malaysia. The high biochemical oxygen demand (BOD) and chemical oxygen demand (COD) of POME lead to oxygen depletion in aquatic ecosystems, altering their chemical composition and causing ecological stress. Furthermore, long-term groundwater contamination threatens human health, as organic pollutants can degrade into toxic by-products over time. Conventional remediation techniques, such as physical and chemical treatments, are often expensive, labor-intensive, and unsustainable in the long term. Microbial techniques, on the other hand, offer a promising alternative by utilizing the natural metabolic processes of microorganisms to degrade and transform organic pollutants into less harmful compounds. These bioremediation methods are not only cost-effective and environmentally friendly but also scalable for large-scale applications. Given Malaysia's extensive reliance on groundwater resources and the prominence of its palm oil industry, addressing POME-related contamination is critical for sustainable water resource management. By focusing on microbial remediation strategies, this study aims to explore practical, efficient, and sustainable solutions to mitigate groundwater pollution, ensuring both environmental protection and economic viability in Malaysia's palm oil industrial zones.

2. Sources and Impacts of Organic Pollution

The palm oil industry is one of the most resource-intensive sectors in Malaysia, producing an estimated 2.7

metric tons of palm oil mill effluent (POME) per metric ton of crude palm oil. POME is a highly organic wastewater by-product generated during the sterilization, clarification, and hydro-extraction processes of palm oil production. Rich in organic matter, POME contains elevated concentrations of fatty acids, carbohydrates, lipids, proteins, and phenolic substances, along with suspended solids and traces of heavy metals. Due to its high biochemical oxygen demand (BOD) and chemical oxygen demand (COD), POME is considered one of the most challenging industrial effluents to treat effectively.

Table 1. POME Components and Impacts on Groundwater

Component	Concentration Range (mg/L)	Potential Impact on Groundwater
Fatty Acids	1000-8000	Can cause oxygen depletion, leading to anoxic conditions.
Phenolic Compounds	50-200	Toxic to aquatic life and can degrade into harmful by-products.
Carbohydrates	3000-5000	Elevated levels can promote microbial growth, leading to eutrophication.
Suspended Solids	5000-15000	Reduces water clarity, clogs filtration systems, and impacts aquatic habitats.
Nitrogen (Total Kjeldahl)	200-1000	Excessive nitrogen can lead to nitrate contamination, harmful to human health.
Phosphorus	50-100	Phosphorus promotes algal blooms, reducing water quality.

The improper handling, storage, or disposal of POME is a major contributor to groundwater contamination in Malaysia. In many cases, POME is stored in unlined ponds or lagoons for treatment, which risks seepage into the soil and subsequent infiltration into aquifers. Moreover, accidental spills or poorly managed discharge systems exacerbate the situation, creating pathways for contaminants to enter groundwater reserves. The porous nature of the tropical soils in palm oil industrial zones facilitates the rapid migration of pollutants into aquifers, making the contamination more widespread and difficult to contain. The environmental impacts of organic pollution are extensive. Elevated BOD and COD levels deplete dissolved oxygen in aquatic ecosystems, leading to anoxic conditions that are detrimental to aquatic life. This disrupts ecological balance, affects biodiversity, and degrades natural ecosystems. Additionally, organic pollutants, such as phenolic compounds, can persist in groundwater systems, undergoing degradation into toxic by-products such as phenol derivatives or even carcinogenic substances, further compounding environmental and health risks. On a societal level, groundwater contamination threatens the safety and availability of potable water for local communities. For instance, many rural and agricultural areas in Malaysia rely heavily on groundwater for drinking, irrigation, and household use. Contaminated water sources expose populations to chronic health risks, including gastrointestinal diseases, reproductive disorders, and even cancer. Additionally, such pollution undermines agricultural productivity due to poor water quality, leading to broader socio-economic consequences. The palm oil industry's expansion in Malaysia has exacerbated the challenge of managing organic pollution, calling for sustainable, scalable, and effective remediation strategies. Addressing these challenges through advanced bioremediation techniques, such as microbial approaches, offers a promising pathway to mitigate the environmental and health impacts of groundwater contamination in palm oil industrial regions.

3. Microbial Remediation Techniques

Microbial remediation, or bioremediation, is an environmentally friendly and cost-effective method for addressing groundwater contamination caused by organic pollutants. It leverages the metabolic capacities of microorganisms to break down, transform, or neutralize harmful compounds into less toxic or inert by-products. The application of microbial remediation in palm oil industrial areas is particularly promising due to its scalability, adaptability, and compatibility with the local environment. This section delves into key microbial techniques used for groundwater remediation, including biostimulation, bioaugmentation, and combined anaerobic-aerobic processes.

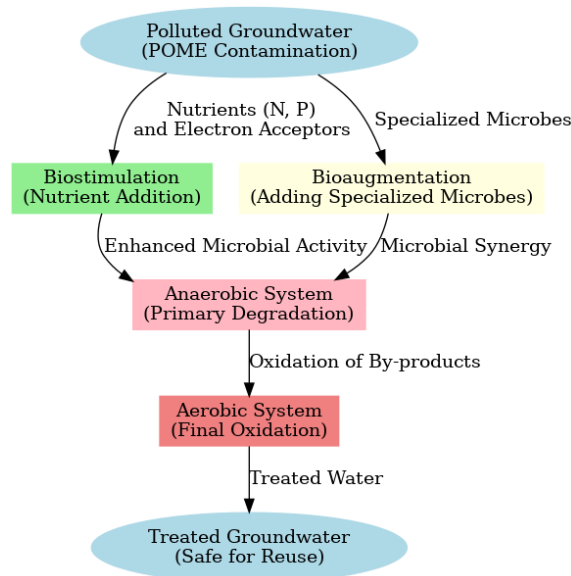


Figure 1. Microbial Remediation Process

3.1 Biostimulation

Biostimulation involves enhancing the activity of indigenous microbial populations by introducing essential nutrients, co-factors, or electron acceptors to optimize their metabolic processes. In the context of palm oil industrial areas, the organic-rich nature of POME often supports diverse microbial communities with the potential to degrade specific pollutants. However, these microbes often require external stimulation to function optimally in contaminated groundwater. For instance, adding nitrogen and phosphorus in specific ratios addresses the nutrient imbalance commonly found in POME-contaminated sites, accelerating microbial growth and pollutant degradation rates. Similarly, introducing electron acceptors such as oxygen, nitrate, or sulfate enhances microbial respiration, allowing the breakdown of recalcitrant organic compounds. In one study in Malaysia, biostimulation with nitrate led to significant reductions in chemical oxygen demand (COD) and phenolic concentrations in groundwater within a few months.

3.2 Bioaugmentation

Bioaugmentation is the introduction of specialized or genetically engineered microorganisms with specific degradative capabilities to supplement indigenous microbial communities. This technique is particularly useful when the native microbial population lacks the enzymatic pathways to degrade certain organic pollutants found in POME, such as phenolic compounds or polycyclic aromatic hydrocarbons (PAHs). For example, hydrocarbon-degrading bacteria like *Pseudomonas putida* and *Alcaligenes* have been successfully employed in groundwater remediation projects. These bacteria possess unique metabolic pathways that allow them to degrade phenolic compounds and other persistent pollutants into simpler, less harmful molecules. In palm oil industrial areas, bioaugmentation has been applied in pilot-scale studies to target specific contaminants, achieving reductions in COD and other pollutant metrics by as much as 80% in some cases.

3.3 Combined Anaerobic and Aerobic Processes

POME contaminants often contain both easily degradable compounds and more complex, recalcitrant substances that require different metabolic pathways for effective remediation. Combined anaerobic and aerobic processes offer a comprehensive solution by utilizing the complementary strengths of both systems. In anaerobic conditions, microorganisms such as methanogens and sulfate-reducing bacteria degrade high-molecular-weight organic compounds into simpler intermediates like methane, acetate, and carbon dioxide. These intermediates are further oxidized under aerobic conditions, where aerobic bacteria such as *Bacillus* and *Pseudomonas* complete the degradation process. Integrated systems, such as anaerobic bioreactors followed by aerobic treatment ponds or biofilters, have been widely implemented in Malaysia's palm oil industrial zones. These systems provide a two-stage remediation approach that enhances pollutant breakdown efficiency. For example, anaerobic processes can reduce COD by 60-70%, while subsequent aerobic processes can achieve an additional 20-30% reduction, resulting in groundwater that meets environmental standards.

3.4 Application in Palm Oil Industrial Areas

Field studies in palm oil industrial zones in Malaysia have demonstrated the effectiveness of these microbial remediation techniques. For instance, biostimulation with controlled nutrient inputs reduced COD levels by over 75% in contaminated aquifers within six months. Similarly, pilot-scale bioaugmentation projects using hydrocarbon-degrading bacteria have shown significant reductions in phenolic compound concentrations.

In conclusion, microbial remediation techniques—when tailored to the specific characteristics of POME contamination—hold great promise for addressing groundwater organic pollution in palm oil industrial areas. By combining biostimulation, bioaugmentation, and integrated anaerobic-aerobic systems, these approaches offer sustainable and efficient solutions to mitigate the environmental impact of industrial activities.

4. Functional Roles of Microbial Communities

Microbial communities play a pivotal role in the degradation and transformation of organic pollutants in groundwater contaminated by palm oil mill effluent (POME). These communities consist of diverse microorganisms, including bacteria, fungi, and archaea, each contributing specific metabolic pathways and enzymatic functions to the bioremediation process. Understanding the functional roles of these microbial communities is critical for developing targeted and effective remediation strategies. Advanced molecular tools such as metagenomics, transcriptomics, and proteomics provide insights into the composition, dynamics, and metabolic capabilities of these microbial populations, allowing for the optimization of microbial remediation processes.

4.1 Composition and Diversity of Microbial Communities

Microbial communities in POME-contaminated environments are often enriched with microorganisms capable of degrading organic pollutants. Bacterial genera such as *Pseudomonas*, *Bacillus*, *Clostridium*, and *Alcaligenes* are commonly found in such environments. These microorganisms possess specific enzymes that catalyze the breakdown of phenolic compounds, hydrocarbons, and other organic pollutants in POME. Additionally, methanogenic archaea, such as *Methanobacterium* and *Methanosarcina*, play a vital role in anaerobic degradation by converting organic matter into methane and carbon dioxide.

Fungi, such as *Aspergillus* and *Penicillium*, also contribute to pollutant degradation through their production of extracellular enzymes like lignin peroxidase and manganese peroxidase. These enzymes are particularly effective in breaking down complex, high-molecular-weight compounds that are resistant to bacterial degradation. The synergy among these microbial groups enables the effective transformation of a wide range of pollutants in POME-contaminated groundwater.

Table 2. Key Enzymes, Target Pollutants, And Degradation Pathways

Key Enzyme	Target Pollutant	Degradation Pathway
Lipase	Triglycerides and Lipids	Lipids → Glycerol + Free Fatty Acids
Catechol 2,3-Dioxygenase	Phenolic Compounds	Phenols → Catechol → Central Metabolism (TCA Cycle)
Amylase	Carbohydrates	Starch → Maltose → Glucose
Alcohol Dehydrogenase	Alcohols	Alcohol → Aldehyde → Acetate
Cellulase	Cellulose	Cellulose → Cellobiose → Glucose

4.2 Key Enzymes and Metabolic Pathways

Microbial degradation of POME involves a variety of enzymes and metabolic pathways. Key enzymes include:

- 1) Lipases: Break down triglycerides and other lipids into glycerol and free fatty acids, which are further metabolized through beta-oxidation.
- 2) Catechol 2,3-Dioxygenase: Plays a critical role in the degradation of phenolic compounds, converting them into intermediates that enter the tricarboxylic acid (TCA) cycle.
- 3) Amylases and Cellulases: Degrade carbohydrates in POME, breaking down starches and cellulose into simpler sugars that can be metabolized by microorganisms.
- 4) Alcohol and Aldehyde Dehydrogenases: Involved in the oxidation of alcohols and aldehydes derived from POME into less harmful compounds.

These enzymes work in tandem within complex metabolic pathways to ensure the complete breakdown of pollutants. For example, phenolic compounds are first hydroxylated by monooxygenases and dioxygenases, producing intermediates such as catechol, which is then cleaved by catechol dioxygenase enzymes for further oxidation in central metabolic pathways.

4.3 Role of Molecular Tools in Microbial Analysis

Advanced molecular tools such as metagenomics, transcriptomics, and proteomics have revolutionized the study of microbial communities in contaminated environments. These tools allow researchers to:

Identify Microbial Diversity: Metagenomics provides a comprehensive overview of microbial diversity, identifying species and functional genes present in contaminated groundwater. **Analyze Gene Expression:** Transcriptomics reveals the active metabolic pathways by analyzing gene expression profiles, helping to identify which microorganisms and enzymes are actively involved in pollutant degradation under specific conditions. **Characterize Protein Functions:** Proteomics helps in understanding the functional roles of enzymes and proteins in pollutant degradation, providing insights into the mechanisms of microbial activity.

For example, metagenomic studies in Malaysia's palm oil industrial areas have identified genes encoding enzymes such as lipases, catechol dioxygenases, and alcohol dehydrogenases, which are critical for breaking down POME pollutants. Transcriptomic analysis has further revealed that indigenous microbial communities upregulate genes related to fatty acid degradation and phenol metabolism when exposed to POME contaminants. These findings help in selecting or engineering microbial strains with enhanced degradative capabilities.

4.4 Case Studies

Several studies conducted in Malaysian palm oil industrial areas highlight the functional roles of microbial communities in POME remediation:

- 1) **Selangor Study on Indigenous Microbial Communities:** A study conducted in Selangor identified a microbial community dominated by *Pseudomonas* and *Bacillus* species, which were highly effective in degrading fatty acids and phenolic compounds. By stimulating these communities with nitrate as an electron acceptor, researchers observed a 70% reduction in COD levels within three months.
- 2) **Bioaugmentation in Johor:** In Johor, researchers introduced hydrocarbon-degrading bacteria such as *Alcaligenes eutrophus* into POME-contaminated groundwater. This bioaugmentation approach resulted in a significant decrease in phenol concentrations and a reduction in BOD levels by over 80%.
- 3) **Metagenomic Analysis in Sabah:** A metagenomic study in Sabah palm oil plantations revealed the presence of unique microbial genes encoding lignin-degrading enzymes. This study demonstrated the potential of fungal communities, such as *Aspergillus*, to degrade complex organic polymers in POME, enhancing the overall remediation process.

4.5 Implications for Remediation Strategies

The functional roles of microbial communities provide a foundation for designing effective remediation strategies. By combining insights from molecular studies with field-scale applications, practitioners can tailor bioremediation approaches to specific site conditions. For example, microbial consortia with complementary enzymatic activities can be deployed to degrade diverse pollutants in POME. Furthermore, optimizing environmental conditions such as pH, temperature, and nutrient availability can enhance microbial activity, leading to faster and more efficient pollutant removal.

In conclusion, microbial communities are the cornerstone of bioremediation efforts in POME-contaminated groundwater. Their diverse metabolic capabilities, coupled with advanced molecular tools, enable the effective degradation of complex organic pollutants. Understanding and leveraging these functional roles is critical for developing sustainable, cost-effective, and scalable remediation strategies in Malaysia's palm oil industrial areas.

5. Optimization Strategies for Remediation

Microbial remediation techniques hold immense potential for addressing groundwater contamination in palm oil industrial areas. However, their efficiency is often contingent upon the environmental conditions under which the microbial communities operate. Effective optimization strategies are therefore critical to maximizing the metabolic activity of microorganisms and ensuring the rapid and complete degradation of pollutants. This section delves deeper into four key strategies for optimizing microbial remediation processes: biostimulation, environmental control, integration with constructed wetlands and biofilters, and real-time monitoring and adaptive management.

5.1 Biostimulation with Nutrients and Electron Acceptors

Biostimulation is a vital strategy to enhance the growth and activity of indigenous microbial populations by addressing their metabolic needs. Microorganisms require a balanced supply of essential nutrients such as nitrogen (N) and phosphorus (P), which are often limited in contaminated groundwater. The optimal nitrogen-to-phosphorus (N:P) ratio for biostimulation is generally 10:1 to 100:1, depending on the specific pollutants present.

Nutrient Supplementation: Adding nitrogen (in the form of ammonium or nitrate) and phosphorus (as phosphate)

can stimulate microbial growth and reproduction. For example, studies in Malaysia have shown that controlled nutrient addition led to a 50-70% increase in the degradation rates of fatty acids and phenolic compounds in palm oil mill effluent (POME)-contaminated groundwater. **Electron Acceptors:** Microbial respiration is driven by electron transfer, which requires electron acceptors such as oxygen (aerobic conditions), nitrate, sulfate, or iron (anaerobic conditions). The selection of an electron acceptor depends on the pollutants present and the natural conditions of the groundwater. For instance, nitrate biostimulation has been shown to improve the degradation of hydrocarbons and phenols in anaerobic conditions common in Malaysian aquifers.

A careful balance of nutrient and electron acceptor levels is essential, as excessive supplementation can lead to secondary pollution or microbial inhibition.

5.2 Environmental Control of pH and Temperature

Environmental factors such as pH and temperature significantly influence microbial metabolism and enzymatic activity. The majority of microorganisms involved in POME degradation thrive under neutral to slightly acidic conditions (pH 6.0–7.5) and temperatures between 30°C and 40°C, conditions commonly found in tropical environments like Malaysia.

pH Control: Adjusting pH with buffering agents such as lime (calcium carbonate) or sulfur compounds can optimize enzymatic activity. For example, neutralizing acidic groundwater environments caused by POME decomposition helps maintain optimal microbial performance, preventing the inhibition of key enzymes like lipases and dioxygenases. **Temperature Regulation:** Maintaining ambient temperatures within the optimal range ensures higher reaction rates in microbial processes. In colder seasons or shaded areas, the installation of passive solar heating systems or thermal covers over treatment lagoons can sustain favorable temperatures.

Environmental controls not only enhance microbial degradation efficiency but also prevent the proliferation of competing or pathogenic microbes.

5.3 Integration with Constructed Wetlands and Biofilters

Constructed wetlands and biofilters provide an innovative hybrid solution by combining physical filtration with microbial degradation. These systems mimic natural processes to enhance pollutant removal while offering long-term, low-maintenance solutions for groundwater remediation.

Constructed Wetlands: Wetlands act as biological reactors where microbial communities colonize plant roots and soil. Oxygen diffused from the roots into the rhizosphere supports aerobic degradation, while anaerobic zones in the soil facilitate the breakdown of recalcitrant pollutants. Plants such as *Typha* and *Phragmites* are commonly used due to their high adaptability and nutrient uptake efficiency. **Biofilters:** Engineered biofilters consist of porous media such as activated carbon, sand, or gravel that support microbial biofilms. These biofilms actively degrade pollutants as groundwater passes through the system. In Malaysia, pilot projects using biofilters in palm oil industrial areas have shown significant reductions in COD (up to 85%) and phenolic compounds.

The integration of microbial techniques with these engineered systems ensures that physical filtration and biological degradation occur simultaneously, resulting in more efficient pollutant removal.

5.4 Real-Time Monitoring and Adaptive Management

The complexity of POME-contaminated sites demands a dynamic approach to remediation, supported by real-time monitoring and adaptive management systems. Advances in monitoring technologies, such as biosensors and remote sensing, enable continuous assessment of groundwater quality and microbial activity.

Biosensors: Biosensors use biological components (e.g., enzymes or microbial cells) to detect specific contaminants, such as phenols or hydrocarbons, in real time. These sensors provide immediate feedback on pollutant levels, enabling operators to adjust remediation strategies promptly. **Machine Learning Models:** Machine learning algorithms can analyze large datasets from groundwater monitoring systems to predict contamination trends and optimize remediation protocols. For example, data on microbial gene expression, nutrient levels, and pollutant concentrations can inform decisions on nutrient supplementation or temperature adjustments.

Adaptive management involves the continuous refinement of remediation strategies based on monitoring data. This ensures that remediation efforts remain effective despite changing environmental conditions or pollutant profiles. Optimization strategies are essential for achieving effective microbial remediation of POME-contaminated groundwater in palm oil industrial areas. By addressing nutrient limitations through biostimulation, controlling environmental factors such as pH and temperature, integrating biological processes with engineered systems like constructed wetlands and biofilters, and leveraging real-time monitoring and adaptive management, the efficiency of microbial techniques can be maximized. These strategies not only enhance the degradation of pollutants but also contribute to the sustainability and scalability of remediation efforts in Malaysia's palm oil sector. When implemented holistically, these approaches hold great potential for

restoring groundwater quality, protecting ecosystems, and promoting sustainable industrial practices.

6. Conclusion

The contamination of groundwater by organic pollutants, particularly from palm oil mill effluent (POME), presents a pressing challenge in Malaysia's palm oil industrial areas. With POME containing high levels of organic compounds such as fatty acids, phenols, and carbohydrates, its infiltration into groundwater has serious environmental, ecological, and public health consequences. Traditional remediation techniques, while effective to an extent, are often cost-prohibitive and environmentally unsustainable. This necessitates the adoption of microbial remediation as a viable alternative. Microbial remediation techniques harness the natural metabolic pathways of microorganisms to degrade or transform harmful organic pollutants into benign or less toxic compounds. Indigenous microbial communities, supplemented with biostimulation and bioaugmentation approaches, have shown remarkable efficiency in breaking down complex pollutants. The use of anaerobic and aerobic processes in combination further ensures comprehensive degradation, especially for recalcitrant pollutants. Critical to the success of microbial remediation is the application of optimization strategies. Biostimulation with targeted nutrients and electron acceptors enhances microbial activity, while environmental controls such as pH and temperature adjustments create favorable conditions for pollutant degradation. The integration of microbial techniques with constructed wetlands and biofilters provides a sustainable and scalable hybrid solution. Real-time monitoring and adaptive management allow for dynamic and efficient remediation, even in complex and changing environments. Case studies in Malaysia have demonstrated the efficacy of these approaches, achieving significant reductions in BOD, COD, and phenolic compounds in groundwater. As the palm oil industry continues to grow, the implementation of tailored microbial remediation strategies can play a pivotal role in safeguarding groundwater resources, protecting ecosystems, and ensuring the long-term sustainability of Malaysia's natural and economic systems. By leveraging advanced microbial technologies and site-specific interventions, Malaysia can establish itself as a leader in sustainable industrial practices while addressing one of its most critical environmental challenges.

References

- Ahmad, A. L., Ismail, S., & Bhatia, S., (2003). Water recycling from palm oil mill effluent (POME) using membrane technology. *Desalination*, 157(1-3), 87-95.
- Chan, Y. J., Chong, M. F., Law, C. L., & Hassell, D. G., (2009). A review on anaerobic-aerobic treatment of industrial and municipal wastewater. *Chemical Engineering Journal*, 155(1-2), 1-18.
- Choi, W., & Lee, S. Y., (2018). Microbial degradation of phenolic compounds under anaerobic conditions. *Applied Microbiology and Biotechnology*, 102(24), 10725-10735.
- Das, N., & Chandran, P., (2011). Microbial degradation of petroleum hydrocarbon contaminants: An overview. *Biotechnology Research International*, 2011, 1-13.
- Hasanudin, U., Brebes, R., & Setyawan, R., (2015). Anaerobic-aerobic treatment for palm oil mill effluent (POME) to meet the standard quality of effluent. *Procedia Environmental Sciences*, 23, 309-316.
- Rabus, R., Hansen, T. A., & Widdel, F., (2013). Dissimilatory sulfate- and sulfur-reducing prokaryotes. *The Prokaryotes*, 309-404.
- Sánchez, C., (2009). Lignocellulosic residues: Biodegradation and bioconversion by fungi. *Biotechnology Advances*, 27(2), 185-194.
- Wu, T. Y., Mohammad, A. W., Jahim, J. M., & Anuar, N., (2010). Pollution control technologies for the treatment of palm oil mill effluent (POME) through end-of-pipe processes. *Journal of Environmental Management*, 91(7), 1467-1490.
- Yacob, S., Hassan, M. A., Shirai, Y., Wakisaka, M., & Subash, S., (2006). Baseline study of methane emission from anaerobic ponds of palm oil mill effluent treatment. *Science of the Total Environment*, 366(1), 187-196.

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