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## Unlocking the Potential: Microbial Strategies for Mercury Detoxification: A Review

Arindam Chakraborty<sup>\*</sup>, Arnab Ganguli, Pritilata Adhikary, Anuranjana Mukherjee

Techno India University, Kolkata -70009, West Bengal, India

#### Abstract

Mercury pollution, resulting from the combustion of fossil fuels and various industrial activities, has significant ecological consequences, affecting both aquatic and terrestrial ecosystems. The process of biosorption, where microorganisms selectively absorb and adsorb solutes onto their cell surfaces, plays a crucial role in mercury removal from the environment. Bacteria, algae, fungi, yeasts, and biofilms are the primary microbial candidates involved in biosorption. Bioaccumulation in microorganisms is another essential phenomenon for mercury detoxification. Microbes gradually accumulate mercury within their cells through uptake and storage processes, with intracellular sequestration occurring through the interactions with various cellular components. Bioprecipitation facilitates the removal of mercury ions from solution by forming insoluble metal compounds through microbial-mediated precipitation processes. Bioleaching plays a key role in transforming mercury forms in the environment by solubilizing heavy metals from solid matrices, making them available for subsequent microbial processes. Biovolatilization is a nonmetabolic process that converts toxic inorganic contaminants into less toxic organic and volatile compounds, reducing the risk of mercury accumulation in food chains and subsequent human exposure. Mercuric reductase and organomercurial lyase are enzymatic systems involved in mercury detoxification, with mercuric reductase reducing mercury ions to their elemental form and organomercurial lyase cleaving carbon-mercury bonds in organic mercury compounds. Microorganisms regulate genetic expression to optimize detoxification processes based on environmental mercury concentrations. Horizontal gene transfer enables the dissemination of mercury resistance genes among bacteria, contributing to their adaptation to diverse environments. Understanding these mechanisms offers opportunities for bioremediation strategies, harnessing microbial capabilities to address environmental pollution challenges.

Keywords: Mercury Toxicity; Bioremediation; Mercury Detoxification

## 1. Introduction to Mercury Contamination and Its Impact on Health

**1.1. Overview of mercury as a hazardous heavy metal:** Mercury, a naturally occurring element, is renowned for its hazardous properties as a heavy metal. Its prevalence in various forms across the environment poses significant risks to human health and ecosystems alike. Mercury exists in different chemical forms, with methylmercury being the most toxic and commonly encountered in the environment. The primary sources of mercury contamination

include industrial processes such as coal combustion, mining, and waste incineration, as well as natural processes like volcanic eruptions and weathering of rocks(Virtanen et al., 2007). One of the most concerning aspects of mercury pollution is its ability to bioaccumulate and biomagnify in food chains. Methylmercury, formed through microbial processes in aquatic environments, accumulates in fish and seafood. As larger predatory fish consume smaller ones, the concentration of mercury increases up the food chain, posing significant risks to humans who consume contaminated fish as part of their diet(Virtanen et al., 2007). This bioaccumulation phenomenon has led to widespread advisories on fish consumption, particularly for vulnerable populations such as pregnant women and children. The toxic effects of mercury on human health are well-documented (Chen and Dong, 2022). Exposure to even small amounts of mercury can lead to severe neurological, developmental, and reproductive disorders. Acute mercury poisoning can result in symptoms such as tremors, insomnia, memory loss, and respiratory issues. Chronic exposure, often through long-term consumption of contaminated food or water, can lead to more subtle yet equally damaging health effects, cognitive impairment, cardiovascular including compromised immune problems, and function (Fernandes Azevedo et al., 2012). Beyond its impact on human health, mercury contamination also poses significant threats to ecosystems. Aquatic organisms, particularly those in sensitive ecosystems such as wetlands and estuaries, are highly susceptible to mercury toxicity. Mercury pollution can disrupt reproductive cycles, impair growth and development, and even lead to population declines in affected species. Furthermore, mercury contamination can persist in the environment for extended periods, contributing to long-term ecological damage. A combination of international cooperation, technological improvements, and regulatory measures used in the fight against are mercury pollution.(Driscoll et al., 2013). The Minamata Convention on Mercury, adopted in 2013, aims to reduce mercury emissions and phase out the use of mercury in various industrial processes. Additionally, strategies such as improved waste management practices, adoption of cleaner technologies, and restoration of contaminated sites play crucial roles in addressing mercury pollution.

1.2. Sources of mercury pollution (natural vs. anthropogenic): Mercury pollution stems from both natural processes and human activities, with distinct sources contributing to its environmental presence. Naturally, mercury is released into the environment through geological processes such as volcanic eruptions, weathering of rocks containing mercury ores, and the degassing of mercury-rich soils and sediments(Pirrone et al., 2010). These natural sources have been contributing to atmospheric mercury levels for millennia, albeit at relatively low levels compared to anthropogenic sources.On the other hand, anthropogenic activities have dramatically increased mercury emissions in recent centuries. Industrial processes, particularly those involving the combustion of fossil fuels such as coal and oil, represent one of the largest sources of anthropogenic mercury pollution. Coal-fired power plants, in particular, release substantial amounts of mercury into the atmosphere during combustion, either as elemental mercury vapor or in the form of particulate matter. Other industrial activities such as metal smelting, cement production, and waste incineration also release mercury into the air and water(Streets et al., 2018).Furthermore, certain human practices, such as small-scale gold mining, contribute significantly to mercury pollution. Artisanal and small-scale gold mining operations often use mercury to extract gold from ore, leading to the release of large quantities of mercury into the environment. In these operations, elemental mercury is mixed with crushed ore to form an amalgam, which is then heated to vaporize the mercury, leaving behind the gold. However, much of the released mercury ends up in the environment, contaminating soil, waterways, and ecosystems.Additionally, mercury pollution can result from the improper disposal of mercury-containing products such as fluorescent light bulbs, thermometers, and batteries. When these products end up in landfills or incinerators, mercury can be released into the environment through leaching or combustion processes. Similarly, the use of mercury in various industrial processes and consumer products, although declining due to regulatory measures, continues to contribute to mercury pollution through emissions and improper waste management(Lindqvist, 1995).The distinction between natural and anthropogenic sources of mercury pollution is crucial for understanding its environmental distribution and devising effective mitigation strategies. While natural sources contribute to background levels of mercury contamination, anthropogenic activities have significantly amplified emissions, leading mercury to widespread environmental and human health impacts. Addressing mercury pollution requires comprehensive efforts to reduce emissions from industrial processes, phase out mercury-containing products, promote cleaner technologies, and implement proper waste management practices(Mohapatra et al., 2007).By targeting both natural and anthropogenic sources, it is possible to mitigate the adverse effects of mercury pollution and protect ecosystems and human health from its harmful effects.

2. Ecological consequences of mercurv contamination: Mercury contamination has profound ecological consequences, affecting both aquatic and terrestrial ecosystems. Let's explore these impacts in detail: Aquatic Ecosystems: Bioaccumulation: Mercury enters aquatic systems primarily through industrial processes, mining, and atmospheric deposition. Once in water bodies, it transforms into methylmercury, a highly toxic form. Methylmercury accumulates in aquatic organisms, especially fish, as they ingest contaminated prey. This bioaccumulation affects entire food chains, with top predators accumulating the highest concentrations. Impaired

Reproduction and Behavior: High mercury levels in fish can lead to reproductive failure, developmental abnormalities, and altered behavior(Mohapatra et al., 2007).For instance, bald eagles and ospreys suffer reduced breeding success due to mercury exposure. Additionally, mercury disrupts the endocrine system, affecting hormone regulation in aquatic organisms(Zhu et al., 2000). Ecosystem Function: Mercury influences nutrient cycling, impacting primary production and decomposition rates. Altered nutrient dynamics affect algae, macrophytes, and zooplankton, which form the base of aquatic food webs(Dranguet et al., 2014).Consequently, mercury contamination can lead to shifts in ecosystem structure and function. Wetlands and Marshes: Wetlands play a critical role in mercury cycling. They can either methylate or demethylate mercury. Methylmercury production occurs in wetland sediments, leading anaerobic to its accumulation in aquatic organisms. Wetland loss or degradation exacerbates mercury contamination. Terrestrial Ecosystems: Soil and Forests: Mercury affects soil microbial communities, altering nutrient cycling and organic matter decomposition. Trees can take up mercury through their roots, impacting forest health and nutrient availability. Mercury-contaminated forests may contribute to downstream water pollution. Wildlife and Biodiversity: Mercury contamination affects wildlife populations. Birds, such as kingfishers, herons, and ducks, accumulate mercury from their aquatic prey. This can lead to reduced breeding success, impaired immune function, and altered migration patterns. Additionally, mercury exposure affects small mammals and reptiles(Wolfe et al., 1998).Climate Change Interaction: Climate change influences mercury cycling. Thawing permafrost releases stored mercury into aquatic systems. Increased temperatures enhance mercury methylation rates, exacerbating contamination. The interplay between climate change and mercury warrants further research(Chételat et al., 2022).

### 3. Microbial Bioremediation: A Green Approach

**3.1.** Understanding bioremediation and its advantages: Bioremediation is a powerful and eco-friendly approach for cleaning up contaminated environments. It relies on natural processes, utilizing microorganisms, fungi, green plants, or their enzymes to break down hazardous substances into less toxic or non-toxic forms (Fig1). Bioremediation minimizes damage to ecosystems by relying solely on natural processes. Unlike chemical methods, it does not introduce harmful substances into the environment. The use of microorganisms instead of chemicals makes it an environmentally sustainable approach. It is a costeffective method. It requires minimal complex tools and equipment.Traditional cleanup methods often involve expensive machinery and extensive labour, whereas bioremediation utilizes naturally occurring organisms(Mukherjee et al., 2021).Bioremediation practices can enhance soil health. Microorganisms break down contaminants, improving soil quality.Nutrient cycling and organic matter decomposition are positively influenced by microbial activity during bioremediation.Bioremediation strategies are adaptable to various types of contaminants and environmental conditions.Whether it's groundwater, soil, or air pollution, bioremediation offers flexible treatment solutions.

## **3.2. Importance of microbial strategies for environmental health**

Microbial strategies play a pivotal role in safeguarding environmental health, impacting ecosystems, human well-being, and the delicate balance of our planet. Let's explore why these tiny organisms wield such significant influence:

**3.2.1. Bioremediation**, driven by microorganisms, is a sustainable approach to mitigate pollution. These microbes break down harmful substances, such as heavy metals, pesticides, and hydrocarbons, into less toxic forms. For instance, bacteria can transform oil spills into harmless compounds, aiding ecosystem recovery(Verma et al., 2017).

**3.2.2. Detoxification** occurs when microbes alter pollutants, rendering them less harmful. They metabolize chemicals, reducing their impact on soil, water, and air quality(Devi et al., 2022).

**3.2.3. Nutrient Cycling and Soil health:** Microbes are essential for nutrient cycling. They decompose organic matter, releasing nutrients like nitrogen and phosphorus. These nutrients nourish plants, supporting terrestrial ecosystems. Soil health relies on microbial activity. Beneficial bacteria and fungi enhance soil structure, nutrient availability, and water retention. Healthy soils sustain agriculture, prevent erosion, and sequester carbon(Devi et al., 2022).

**3.2.4. Wastewater Treatment:** Microbes are unsung heroes in wastewater treatment plants. They break down organic pollutants, purifying water before it reenters natural systems. Without them, water bodies would suffer from contamination.

**3.2.5. Biogeochemical Cycling:** Microbes participate in biogeochemical cycles, including the carbon, nitrogen, and sulfur cycles. They convert organic matter, regulate greenhouse gas emissions, and maintain global climate balance.Nitrogen-fixing bacteria transform atmospheric nitrogen into forms usable by plants, enriching soil fertility.

**3.2.6. Human Health and Immunity:** Gut microbiota influence human health. They aid digestion, synthesize vitamins, and modulate immune responses. Disruptions in the gut microbiome correlate with diseases like obesity, diabetes, and allergies.Skin and oral microbiomes also impact health. Understanding these microbial communities can lead to preventive measures and personalized medicine.

**3.2.7. Biocontrol and Agriculture:**Beneficial microbes act as biocontrol agents. They combat plant pathogens, reducing reliance on chemical pesticides. This sustainable approach promotes crop health and food security.Microbes enhance soil fertility by fixing nitrogen, solubilizing phosphates, and promoting plant growth.

**3.2.8. Genetic Resources and Biodiversity:**The vast microbial diversity harbors geneticresources. These genes encode enzymes, bioactive compounds, and metabolic pathways. Bioprospecting unlocks novel applications in medicine, industry, and biotechnology.Conserving microbial biodiversity ensures resilience against environmental changes and supports ecosystem functions.

## 3.2.9. Mercury Transformation by Microorganisms:

3.3. Biosorption-(Mechanisms and microbial candidates): Biosorption is a fascinating physicochemical process that relies on the remarkable abilities of microorganisms to interact with and sequester various solutes from aquatic solutions. Let's delve into the mechanisms behind biosorption and explore some microbial candidates involved in this essential environmental process(Tsezos, 2013). Now comes the mechanism which is followed during biosorption. FirtAbsorption and Adsorption takes place where Microbes selectively absorb and adsorb solutes onto their cell surfaces. This process occurs independently of their metabolic activities. Followed by ion exchanging through microbial biomass exchanges ions with the surrounding environment, effectively capturing pollutants. After that Surface Complexation reactions occur at the microbial surface, binding solutes through specific interactions. Due to which precipitation occurs in which some microorganisms facilitate the precipitation of metal ions, removing them from the solution following a non metalic pathway, unlike traditional metabolic processes, biosorption is nonmetabolic, making it an efficient and spontaneous phenomenon.

# **3.3.1. Microbial Candidate involved for Biosorption:**

**3.3.1.1. Bacteria:** Various bacterial species exhibit excellent biosorption capabilities. Their cell walls contain functional groups (such as carboxyl, amino, and sulfhydryl) that readily bind to metal ions.

**3.3.1.2. Algae and Fungi:** Algae and fungi possess abundant surface sites for biosorption. Algal cell walls, rich in polysaccharides, can sequester heavy metals.

**3.3.1.3. Yeasts:** Yeast cells have been studied extensively for their biosorption potential. Their surface components, both metabolically and nonmetabolically linked, can complex with metal ions.

**3.3.1.4. Biofilms:** Microbial biofilms—structured communities of bacteria—enhance biosorption due to their increased surface area and cooperative interactions.

**3.4. Bioaccumulation:** Bioaccumulation, is a fascinating phenomenon, involves the gradual accumulation of contaminants within an organism's tissues over time. In the case of mercury (Hg), microorganisms play a crucial role in sequestering this heavy metal. Let's explore how bioaccumulation occurs and the microbial mechanisms behind it(Fisher, 1995). The process of Bioaccumulation is divided into few steps they are-

**5.2.1. Uptake and Storage:** Microorganisms, particularly aquatic ones, take in mercury from their surroundings. This process involves the absorption and storage of mercury within their cells.

**5.2.2. Intracellular Sequestration:** Once inside the microbe, mercury can be sequestered intracellularly. Various cellular components, such as proteins and organelles, participate in this process.

**5.2.3.** Accumulation: As microorganisms continue their life cycles, they accumulate mercury. This gradual buildup occurs as they feed, grow, and reproduce.

5.3. **Bioprecipitation**is fascinating microbial processes that contribute significantly to transforming mercury forms in the environment(Essa et al., 2002).Bioprecipitation involves the microbialmediated precipitation of heavy metals, including mercury, from contaminated sites. In those contaminated sites, microorganisms, such as bacteria and fungi, interact with metal ions present in the environment along with it they facilitate the formation of insoluble metal compounds (precipitates) by

binding the metal ions to their cell surfaces or secreting extracellular substances. And in the case of mercury, bioprecipitation leads to the removal of mercury ions from solution. These precipitates are less mobile and less toxic, reducing the risk of mercury exposure to living organisms. Due to all these reasons bioprecipitation is used in wastewater treatment, groundwater remediation, and soil cleanup. It offers a sustainable and cost-effective approach to managing mercury contamination.

Bioleaching: These are essential microbial 5.4. strategies for transforming mercury forms, making them less harmful and aiding in environmental detoxification. Bioleaching is a process where microorganisms solubilize heavy metals from solid matrices, such as ores or sediments. There are some acidophilic microorganisms present, including bacteria and fungi, play a key role in bioleaching. They release organic acids and other metabolites that dissolve mercury from its mineral forms.Mercury often associates with iron or sulfur minerals. Bioleaching disrupts these associations, liberating mercury ions into the surrounding environment. Due to above all the reasons bioleaching enhances the availability of mercury for subsequent microbial processes, such as methylation (conversion to methylmercury). It influences the fate of mercury in aquatic ecosystems and affects its bioavailability to organisms.

**5.5. Biovolatilization** is a fascinating microbial process that involves the conversion of toxic inorganic contaminants, such as heavy metals, into less toxic organic and volatile compounds. Unlike traditional metabolic pathways, biovolatilization is nonmetabolic, making it an efficient and spontaneous phenomenon(Yin et al., 2019).

**5.5.1. Mercury Methylation:** Microorganisms, particularly bacteria and certain algae, methylate inorganic mercury (Hg^2+) to form methylmercury (Me-Hg).Methylmercury is more volatile and less toxic than its inorganic counterpart.This process occurs in aquatic environments, where mercury is present in various forms.

**5.5.2. Redox Reactions:** Microbes facilitate redox reactions, converting mercury ions between different oxidation states. These reactions alter the chemical speciation of mercury, affecting its mobility and toxicity. By participating in redox processes, microorganisms contribute to the transformation of mercury into less harmful forms.

**5.5.3.** Cell Membrane Transport:Once methylated, mercury can cross cell membranes more readily.This allows for its release into the environment, where it can enter the atmosphere and participate in global biogeochemical cycles.Biovolatilization

minimizes the bioavailability of mercury to organisms. Along with it reduces the risk of mercury accumulation in food chains and subsequent human exposure. By understanding this microbial process is essential for managing mercury pollution and safeguarding environmental health.

## 6. Enzymatic Systems Involved in Mercury Detoxification

6.1. Mercuric reductase, a critical enzyme in microbial metabolism, plays a pivotal role in the reduction of mercury ions. Let's delve into the significance of this enzyme and its mechanisms(Marteyn et al., 2013).Mercuric reductase catalyzes the reduction of mercuric ions (Hg<sup>2</sup>+) to their elemental form, mercury (Hg<sup>0</sup>). This transformation is essential for detoxifying mercury in the environment. Also the enzyme transfers electrons from a suitable electron donor (such as NADH or NADPH) to mercuric ions, leading to their conversion into less toxic and less mobile elemental mercury.Mercuric reductase is found in various microorganisms, including bacteria, archaea, and fungi. These organisms have evolved this enzyme to survive in mercury-contaminated habitats.Different microbial species possess distinct forms of mercuric reductase, reflecting their adaptation to specific environmental conditions.Researchers harness mercuric reductase for bioremediation purposes. By expressing this enzyme in engineered microbes, we can enhance mercury reduction in contaminated sites.Efficient reduction of mercuric ions prevents their subsequent methylation to toxic methylmercury (Me-Hg). Me-Hg bioaccumulates in food chains, posing risks to ecosystems and human health. But the mercuric reductase exhibits substrate specificity, limiting its effectiveness against other heavy metals. Further researches increasing the understanding of mercuric reductase at the molecular level opens avenues for sustainable biotechnological solutions to mercury pollution.

6.2. Organomercrudial lyase: It is a remarkable enzyme that plays a crucial role in breakdown of organic mercury compounds. Let's explore the significance of the enzyme and its mechanism(Lello et al., 2004). Organomercurial lyase catalyzes the cleavage of carbon-mercury bonds found in organic mercury compounds.After the breakdown of these compounds, the enzyme prevents their accumulation and potential harm to living organisms.Organomercurial lyase specifically targets the carbon-mercury bonds in organic molecules. It cleaves these bonds, releasing elemental mercury (Hg<sup>0</sup>) and the corresponding organic fragment. Different forms of organomercurial lyase exist, each adapted to specific organic mercury compounds. For

example, some enzymes act on methylmercury (Me-Hg), while others target ethylmercury or phenylmercury. The merits of Organomercurial lyase contributes in cycling of mercury in ecosystems. It converts organic mercury back into elemental mercury, which can then volatilize into the atmosphere. It also helps in breaking down of organic mercury, the enzyme influences its availability to organisms. This impacts the transfer of mercury through food chains.

7. Genetic regulation of mercury uptake: The genetic regulation of mercury uptake and detoxification is a dynamic interplay that allows microorganisms to thrive in diverse environments and contribute to environmental health(Foster, 1987). Microorganisms possess specific genes encoding mercury transporters(Wilson et al., 2000). These transporters facilitate the uptake of mercury ions (Hg^2+) from the surrounding environment into the cell.The expression of these transporters is tightly regulated. When mercury levels are low, the cell downregulates their production. However, in the presence of mercury, these genes are upregulated, allowing efficient uptake.

**7.1.** Mercury Detoxification Genes: It is a fascinating area of study that sheds light on how microorganisms adapt to their environment and cope with mercury contamination.

**7.1.1.** Mercuric Reductase (merA): This key enzyme is encoded by the merA gene. It catalyzes the reduction of mercuric ions  $(Hg^{2}+)$  to elemental mercury  $(Hg^{0})$ . By doing so, it detoxifies mercury within the cell.

**7.1.2. Organomercurial Lyase (merB):** The merB gene encodes organomercurial lyase. This enzyme breaks down organic mercury compounds, cleaving carbon-mercury bonds. It prevents the accumulation of toxic organic mercury.

**7.1.3. Regulatory Proteins:** Microorganisms also produce regulatory proteins (such as MerR) that sense mercury levels. These proteins bind to specific DNA sequences (mer operators) and control the expression of detoxification genes.

7.2. Mercury Resistance Operons:

**7.2.1. mer Operon:** In many bacteria, mercury resistance genes are organized into an operon called the **mer operon**. This operon includes genes for transporters, reductase, and other components.

**7.2.2.** Transcriptional Control: The mer operon is regulated at the transcriptional level. When mercury is present, the MerR protein binds to the operator region,

activating transcription. This ensures that the necessary detoxification machinery is produced.

## 7.3. Horizontal Gene Transfer:

**7.3.1. Plasmids and Transposons:** Mercury resistance genes can be transferred horizontally between bacteria. Plasmids and transposons carry these genes and allow their dissemination.

**7.3.2.** Adaptation and Evolution: Through horizontal gene transfer, microorganisms acquire mercury resistance genes from other species. Over time, this contributes to their adaptation and survival in mercury-contaminated environments.

### 8. Ecological Implications:

**8.1. Bioremediation Potential:** Understanding genetic regulation helps design strategies for enhancing microbial mercury detoxification. Engineered microbes with optimized detoxification genes can aid in bioremediation efforts.

**8.2.** Natural Variation: Different microbial species exhibit varying levels of mercury resistance due to genetic differences. Some thrive in highly contaminated sites, while others are more sensitive.

**9. Microbial Communities and Hybrid Approaches** Maintaining stable consortia over time is essential. Environmental fluctuations can impact their composition and function.Researchers explore genetic modifications to optimize consortia for specific pollutants. Whilebridging the gap between lab-scale knowledge and large-scale applications remains a challenge.

Efficiency of microbial consortia: Microbial 9.1. consortia for efficient mercury remediation it represents a powerful approach for efficient mercury These diverse communities remediation. of microorganisms work synergistically to tackle mercury contamination in various environments. The significance of microbial consortia and their strategies for effective mercury detoxification are-

**9.1.1. Cooperative Efforts:**Microbial consortia consist of different species like bacteria, archaea, and fungi, each contributing unique capabilities. By working together, they enhance overall remediation efficiency.Some microbes specialize in mercury uptake, while others excel at detoxification. Their combined efforts lead to more comprehensive and sustainable remediation.

#### 9.2. Mechanisms of Mercury Detoxification:

**9.2.1. Metabolic Diversity:** Within consortia, microorganisms employ various metabolic pathways to detoxify mercury:

**9.2.1.1. Biosorption:** Some species adsorb mercury onto their cell surfaces.

**9.2.1.2. Bioprecipitation:** Others facilitate the precipitation of mercury ions, removing them from the environment.

**9.2.1.3. Bioleaching:** Certain microbes dissolve mercury from solid matrices.

**9.2.1.4. Bioaccumulation:** Some accumulate mercury within their cells.

**9.2.1.5. Biovolatilization:** A few volatilize mercury, releasing it into the atmosphere.

**9.2.1.6.** Enzymatic Systems: Enzymes like mercuric reductase and organomercurial lyase play critical roles in these processes.

**9.3.** Adaptation and Synergy:Microbial consortia adapt to specific conditions, such as pH, temperature, and oxygen availability. This adaptability allows them to thrive in diverse habitats. And cooperative interactions within consortia enhance their resilience. Biofilms, for example, provide a structured environment where microbes exchange nutrients and signals.

**10. Bioremediation Applications:** 

**10.1. In Situ and Ex Situ Remediation:** Microbial consortia can be applied directly in contaminated sites (in situ) or in controlled laboratory conditions (ex situ).

**10.2. Hybrid Approaches using advanced materials:** Combining microbial consortia with novel biosorbents (such as carbon-based materials or nanoparticles) boosts their efficiency. Hybridizing microbes with advanced materials (e.g., carbon-based, nanoparticles) It is an innovative approach that holds immense promise for mercury bioremediation. By combining the unique properties of microorganisms with those of advanced materials, researchers aim to create efficient and sustainable strategies for removing mercury from contaminated environments.

## 10.2.1. Carbon-Based Materials:

**10.2.1.1.Carbon Nanotubes (CNTs)**: These onedimensional carbon structures possess high surface area, excellent conductivity, and mechanical strength. When hybridized with microbes, CNTs enhance their ability to adsorb and sequester mercury ions(Alijani et al., 2015).

**10.2.1.2. Graphene**: A single layer of carbon atoms arranged in a two-dimensional lattice, graphene offers remarkable electrical conductivity and chemical stability. Integrating graphene with microbial systems improves their performance in mercury removal(Kabiri et al., 2016).

### 10.2.2. Metal and Metal Oxide Nanoparticles:

**10.2.2.1.Silver Nanoparticles**: Microbes can synthesize silver nanoparticles with antimicrobial properties. These hybrids exhibit dual functionality—mercury adsorption by the microbial component and antibacterial action by the nanoparticles(Karuppiah Chandran and Pambayan Ulagan, 2017).

**10.2.2.2.Iron Oxide Nanoparticles**: These magnetic nanoparticles can be functionalized and combined with microbial biofilms. They enhance mercury uptake and facilitate easy separation from the solution using external magnets(Assa et al., 2016).

#### 10.3. Synergistic Effects:

**10.3.1.** Enhanced Adsorption: The combination of microbial biomass and advanced materials results in synergistic adsorption of mercury. The materials provide additional binding sites, while microbes contribute their natural affinity for metal ions.

**10.3.2. Redox Reactions**: Nanoparticles participate in redox reactions, converting mercury ions into less toxic forms. Microbes enhance these reactions by providing reducing equivalents(Wang et al., 2017).

#### **10.4.** Applications:

**10.4.1. Wastewater Treatment**: Hybrid microbial systems can be employed in wastewater treatment plants. They efficiently remove mercury from industrial effluents, reducing its environmental impact.

**10.4.2.** In Situ Bioremediation: By introducing these hybrids into contaminated sites, we can enhance mercury detoxification. The materials stabilize the microbial consortia and improve their overall performance.

#### 11. Metagenomics and novel biosorbents:

Metagenomics, a powerful field within microbiology, has revolutionized our understanding of microbial communities by directly analyzing DNA from environmental samples. Unlike traditional culturebased methods, metagenomics allows us to explore the genetic information of uncultivable microorganisms within a specific environment. This approach provides insights into the diversity, functional potential, and adaptive strategies of microbial communities in response to mercury contamination(Allan, 2014).

## 11.1. Metagenomics for Mercury Bioremediation:

**11.1.1. Genetic Diversity**: Metagenomic studies reveal the presence of diverse microbial taxa with the potential to interact with mercury. By analyzing their genomes, we identify genes involved in mercury uptake, detoxification, and resistance(Foster, 1987).

**11.1.2.** Functional Annotation: Metagenomics helps annotate functional genes related to mercury transformation. These include mercuric reductases, organomercurial lyases, and transporters(Foster, 1987).

**11.1.3. Community Dynamics**: Understanding the dynamics of microbial consortia in contaminated sites is crucial. Metagenomics provides snapshots of community composition and gene expression over time(Desai et al., 2010).

## 11.2. Novel Biosorbents and Metagenomics:

**11.2.1 Biosorbents**: These materials have high affinity for binding pollutants. By combining metagenomics with biosorbents, we can discover novel microbial genes encoding biosorption proteins(Vijayaraghavan and Yun, 2008).

**11.2.2** Functional Screening: Metagenomics allows us to screen for genes involved in metal uptake, detoxification, and biosorption. These genes can be expressed in suitable hosts for further study.

## 12. Case Studies and Field-Scale Applications

Successful microbial remediation projects for mercury bioremediation have demonstrated the effectiveness of harnessing microorganisms to mitigate mercury contamination in various environments. These projects leverage the natural abilities of microbes to transform and sequester mercury, offering sustainable and ecofriendly solutions(Löffler and Edwards, 2006). Here are some notable examples:

## 12.1. Industrial Wastewater Treatment:

**12.1.1. Case Study&Approach**: In industrial settings, mercury-contaminated wastewater poses a significant challenge. Successful projects have employed microbial consortia or genetically engineered bacteria to efficiently remove mercury from effluents. These systems combine microbial activity with advanced materials (such as carbon-based

nanomaterials or metal nanoparticles) to enhance mercury adsorption and reduction.

## 12.2. Mining Sites and Soil Remediation:

**12.2.1. Case Study&Approach:** Abandoned mining sites often harbor high mercury levels due to historical mining activities. Microbial remediation projects have revitalized these areas by promoting mercury uptake and transformation.Microbes, including naturally occurring bacteria and fungi, are introduced to contaminated soils. They facilitate mercury detoxification through processes like volatilization, biosorption, and reduction.

### 12.3. Aquatic Ecosystem Restoration:

**12.3.1.** Case Study&Approach: Mercury pollution in aquatic environments affects both aquatic life and human health. Successful projects have focused on restoring contaminated lakes, rivers, and wetlands.Microbial biofilms, combined with carbonbased materials or nanoparticles, have been deployed to reduce mercury levels. These systems enhance mercury removal and minimize its impact on aquatic ecosystems.

## 12.4. Phytoremediation with Microbial Assistance:

**12.4.1.** Case Study & Approach: Phytoremediation involves using plants to extract pollutants from soil. Combining plants with mercury-resistant microbes improves efficiency. Microbes enhance mercury uptake by plants, facilitating its transfer from soil to plant tissues. This combined approach accelerates mercury removal.

#### 12.5. Synergy of Natural Microbial Communities:

**12.5.1.** Case Study&Approach: Natural microbial communities in contaminated environments adapt to mercury stress over time. Successful projects have harnessed these communities for bioremediation. By understanding the genetic diversity and functional potential of these communities through metagenomics, researchers identify key players and optimize their performance.

## 13. Bridging the gap between lab-scale research and real-world implementation

Bridging the gap between lab-scale research and realworld implementation for mercury bioremediation is a critical challenge in environmental science and engineering. While promising lab-scale studies demonstrate the efficacy of microbial approaches for mercury removal, translating these findings into practical applications remains complex.

13.1. Complexity of Real-World Environments:

#### 13.3. Biocompatibility and Safety:

**13.3.1. Materials Compatibility**: Integrating advanced materials (such as carbon-based nanomaterials or nanoparticles) with microbes requires



Fig 1: Different strategies for bioremediation using Microbes and their products

**13.1.1.Lab-Scale Simplicity**: In controlled laboratory settings, researchers can manipulate variables and optimize conditions. However, real-world environments—such as contaminated soils, water bodies, or industrial sites—are multifaceted. Factors like pH, temperature, organic matter, and competing ions significantly impact microbial performance.

**13.1.2.** Adaptation and Resilience: Microbes must adapt to diverse conditions, including fluctuations in mercury concentrations and coexisting pollutants. Bridging the gap requires understanding how microbial communities respond to these complexities.

## 13.2. Long-Term Stability and Sustainability:

**13.2.1. Temporal Dynamics**: Lab experiments often run for shorter durations, while real-world remediation projects extend over years. Ensuring the stability and effectiveness of microbial consortia or genetically engineered strains throughout this extended period is essential.

**13.2.2. Maintenance and Monitoring:** Implementing bioremediation at scale demands continuous monitoring, maintenance, and adaptive management. Researchers must design systems that remain effective over time.

biocompatibility. Materials should not hinder microbial growth or function.

**13.3.2. Human and Ecosystem Safety**: Real-world applications involve exposure to humans and ecosystems. Ensuring that microbial interventions do not inadvertently harm non-target organisms is crucial.

#### 13.4. Scaling Up and Cost-Effectiveness:

**13.4.1. Economic Viability**: Lab-scale research often overlooks economic constraints. Real-world implementation involves costs related to materials, infrastructure, monitoring, and maintenance. Strategies must be economically feasible.

**13.4.2.** Scalability: Moving from small-scale experiments to large-scale field applications requires thoughtful design. Factors like reactor design, mass transfer, and logistics play a significant role.

## **13.5.** Interdisciplinary Collaboration and Policy Integration:

**13.5.1. Team Effort**: Bridging the gap necessitates collaboration among microbiologists, materials scientists, engineers, and policymakers. Interdisciplinary teams can address technical, economic, and regulatory aspects.

**13.5.2. Policy Alignment**: Real-world implementation occurs within regulatory frameworks. Researchers must engage policymakers early to align their strategies with existing guidelines and regulations.

### 14. Challenges and future prospects

Mercury bioremediation faces several challenges and holds promising future prospects.

### 14.1. Challenges:

**Complex Environmental Matrix**: Mercury contamination occurs in diverse environments—soil, water, and air. Each matrix presents unique challenges for bioremediation.

Species-SpecificResponses:Differentmicroorganismsresponddifferentlytomercury.Understandingtheirspecificmechanismsandoptimizingtheirperformanceisessential.

**Long-Term Stability**: Ensuring the stability and effectiveness of microbial consortia or genetically engineered strains over extended periods remains a challenge.

**Biocompatibility**: Integrating advanced materials with microbes requires compatibility to avoid hindering microbial growth or function.

Field-Scale Implementation: Bridging the gap between lab-scale success and large-scale field applications is critical

### 14.2. Future Prospects:

**Omics Techniques**: Advances in metagenomics and other Omics techniques allow us to explore microbial diversity and functional potential. These tools guide the design of effective bioremediation strategies.

**Biotechnological Innovations**: Continued research into microbial genetics and biotechnological applications will yield novel approaches for mercury removal.

**One Health Approach**: Considering the impact of bioremediation on ecosystems, human health, and the environment in a holistic manner is crucial.

**Interdisciplinary Collaboration**: Collaborations between microbiologists, materials scientists, and environmental engineers will drive innovative solutions.

**Metagenomics and Systems Biology**: Leveraging metagenomics and systems biology will enhance our understanding of microbial communities and their functional potential. These approaches guide the design of effective bioremediation strategies.

**Community Engagement**: Involving local communities and stakeholders ensures acceptance, participation, and long-term success of bioremediation projects.

**Education and Outreach**: Bridging the gap also involves educating practitioners, policymakers, and the public about the benefits and limitations of microbial approaches.

**Synergistic Approaches**: Integrating metagenomics with novel biosorbents enhances our understanding of microbial interactions and functional networks. It guides the design of tailored bioremediation strategies.

**Biotechnological Innovations**: Metagenomics opens avenues for sustainable biotechnological solutions. By harnessing microbial diversity, we can develop efficient and eco-friendly methods for mercury removal.

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#### References

- Alijani, H., Shariatinia, Z., Aroujalian Mashhadi, A., 2015. Water assisted synthesis of MWCNTs over natural magnetic rock: An effective magnetic adsorbent with enhanced mercury(II) adsorption property. Chem. Eng. J. 281, 468–481. https://doi.org/10.1016/j.cej.2015.07.025
- [2] Allan, E., 2014. Metagenomics: Unrestricted access to microbial communities. Virulence 5, 397–398. https://doi.org/10.4161/viru.28057
- [3] Assa, F., Jafarizadeh-Malmiri, H., Ajamein, H., Anarjan, N., Vaghari, H., Sayyar, Z., Berenjian, A., 2016. A biotechnological perspective on the application of iron oxide nanoparticles. Nano Res. 9, 2203–2225. https://doi.org/10.1007/s12274-016-1131-9
- [4] Chételat, J., McKinney, M.A., Amyot, M., Dastoor, A., Douglas, T.A., Heimbürger-Boavida, L.-E., Kirk, J., Kahilainen, K.K., Outridge, P.M., Pelletier, N., Skov, H., St. Pierre, K., Vuorenmaa, J., Wang, F., 2022. Climate change and mercury in the Arctic: Abiotic interactions. Sci. Total Environ. 824, 153715. https://doi.org/10.1016/j.scitotenv.2022.153715
- [5] Desai, C., Pathak, H., Madamwar, D., 2010. Advances in molecular and "-omics" technologies to gauge microbial communities and bioremediation at xenobiotic/anthropogen

contaminated sites. Bioresour. Technol. 101, 1558–1569. https://doi.org/10.1016/j.biortech.2009.10.080

- [6] Devi, R., Behera, B., Raza, M.B., Mangal, V., Altaf, M.A., Kumar, R., Kumar, A., Tiwari, R.K., Lal, M.K., Singh, B., 2022. An Insight into Microbes Mediated Heavy Metal Detoxification in Plants: a Review. J. Soil Sci. Plant Nutr. 22, 914–936. https://doi.org/10.1007/s42729-021-00702-x
- [7] Dranguet, P., Flück, R., Regier, N., Cosio, C., Le Faucheur, S., Slaveykova, V.I., 2014. Towards Mechanistic Understanding of Mercury Availability and Toxicity to Aquatic Primary Producers. CHIMIA 68, 799. https://doi.org/10.2533/chimia.2014.799
- [8] Driscoll, C.T., Mason, R.P., Chan, H.M., Jacob, D.J., Pirrone, N., 2013. Mercury as a Global Pollutant: Sources, Pathways, and Effects. Environ. Sci. Technol. 47, 4967–4983. https://doi.org/10.1021/es305071v
- [9] Essa, A.M.M., Macaskie, L.E., Brown, N.L., 2002. Mechanisms of mercury bioremediation. Biochem. Soc. Trans. 30, 672–674. https://doi.org/10.1042/bst0300672
- [10] Fernandes Azevedo, B., Barros Furieri, L., Peçanha, F.M., Wiggers, G.A., Frizera Vassallo, P., Ronacher Simões, M., Fiorim, J., Rossi De Batista, P., Fioresi, M., Rossoni, L., Stefanon, I., Alonso, M.J., Salaices, M., Valentim Vassallo, D., 2012. Toxic Effects of Mercury on the Cardiovascular and Central Nervous Systems. J. Biomed. Biotechnol. 2012, 1–11. https://doi.org/10.1155/2012/949048
- [11] Fisher, S.W., 1995. Mechanisms of Bioaccumulation in Aquatic Systems, in: Ware, G.W. (Ed.), Reviews of Environmental Contamination and Toxicology, Reviews of Environmental Contamination and Toxicology. Springer New York, New York, NY, pp. 87–117. https://doi.org/10.1007/978-1-4612-4252-9\_4
- Foster, T.J., 1987. The Genetics and Biochemistry of Mercury Resistance. CRC Crit. Rev. Microbiol. 15, 117–140. https://doi.org/10.3109/10408418709104455
- [13] Kabiri, S., Tran, D.N.H., Cole, M.A., Losic, D., 2016. Functionalized three-dimensional (3D) graphene composite for high efficiency removal of mercury. Environ. Sci. Water Res. Technol. 2, 390–402. https://doi.org/10.1039/C5EW00254K
- [14] Karuppiah Chandran, P., Pambayan Ulagan, M., 2017. ANTIMICROBIAL AND ANTICANCER ACTIVITY OF SILVER NANOPARTICLES FROM EDIBLE MUSHROOM: A REVIEW. Asian J. Pharm. Clin. Res. 10, 37. https://doi.org/10.22159/ajpcr.2017.v10i3.16027
- [15] Lello, P.D., Lafrance-Vanasse, J., Omichinski, J.G., 2004. The Organomercurial Lyase MerB, in: Messerschmidt, A., Huber, R., Poulas, T., Wieghardt, K., Cygler, M., Bode, W. (Eds.), Handbook of Metalloproteins. Wiley. https://doi.org/10.1002/0470028637.met278
- [16] Lindqvist, O., 1995. Environmental impact of mercury and other heavy metals. J. Power Sources 57, 3–7. https://doi.org/10.1016/0378-7753(95)02229-5
- [17] Löffler, F.E., Edwards, E.A., 2006. Harnessing microbial activities for environmental cleanup. Curr. Opin. Biotechnol. 17, 274–284. https://doi.org/10.1016/j.copbio.2006.05.001
- [18] Marteyn, B., Sakr, S., Farci, S., Bedhomme, M., Chardonnet, S., Decottignies, P., Lemaire, S.D., Cassier-Chauvat, C., Chauvat, F., 2013. The Synechocystis PCC6803 MerA-Like Enzyme Operates in the Reduction of Both Mercury and Uranium under the Control of the Glutaredoxin 1 Enzyme. J. Bacteriol. 195, 4138–4145. https://doi.org/10.1128/JB.00272-13
- [19] Mohapatra, S.P., Nikolova, I., Mitchell, A., 2007. Managing mercury in the great lakes: An analytical review of abatement

policies. J. Environ. Manage. 83, 80–92. https://doi.org/10.1016/j.jenvman.2006.01.015

- [20] Mukherjee, S., Narula, R., Bhattacharjee, Shreya, Dutta, D., Bose, I., Mahakud, J., Paul, Sharanya, Bhattacharjee, Sourish, Paul, Sonali, 2021. BIOREMEDIATION: THE ECO-FRIENDLY SOLUTION TO THE HAZARDOUS PROBLEM OF ENVIRONMENTAL POLLUTION. J. Environ. Eng. Landsc. Manag. 29, 477–483. https://doi.org/10.3846/jeelm.2021.14439
- [21] Pirrone, N., Cinnirella, S., Feng, X., Finkelman, R.B., Friedli, H.R., Leaner, J., Mason, R., Mukherjee, A.B., Stracher, G.B., Streets, D.G., Telmer, K., 2010. Global mercury emissions to the atmosphere from anthropogenic and natural sources. Atmospheric Chem. Phys. 10, 5951–5964. https://doi.org/10.5194/acp-10-5951-2010
- [22] Streets, D.G., Lu, Z., Levin, L., Ter Schure, A.F.H., Sunderland, E.M., 2018. Historical releases of mercury to air, land, and water from coal combustion. Sci. Total Environ. 615, 131–140. https://doi.org/10.1016/j.scitotenv.2017.09.207
- [23] Tsezos, M., 2013. Biosorption: A Mechanistic Approach, in: Schippers, A., Glombitza, F., Sand, W. (Eds.), Geobiotechnology I, Advances in Biochemical Engineering/Biotechnology. Springer Berlin Heidelberg, Berlin, Heidelberg, pp. 173–209. https://doi.org/10.1007/10\_2013\_250
- [24] Verma, S.K., Trivedi, P., Gupta, A.K., Verma, R.K., 2017. Aromatic Plant–Microbe Associations: A Sustainable Approach for Remediation of Polluted Soils, in: Singh, J.S., Seneviratne, G. (Eds.), Agro-Environmental Sustainability. Springer International Publishing, Cham, pp. 85–103. https://doi.org/10.1007/978-3-319-49727-3\_5
- [25] Vijayaraghavan, K., Yun, Y.-S., 2008. Bacterial biosorbents and biosorption. Biotechnol. Adv. 26, 266–291. https://doi.org/10.1016/j.biotechadv.2008.02.002
- [26] Virtanen, J.K., Rissanen, T.H., Voutilainen, S., Tuomainen, T.-P., 2007. Mercury as a risk factor for cardiovascular diseases. J. Nutr. Biochem. 18, 75–85. https://doi.org/10.1016/j.jnutbio.2006.05.001
- [27] Wang, X., Zhang, D., Pan, X., Lee, D.-J., Al-Misned, F.A., Mortuza, M.G., Gadd, G.M., 2017. Aerobic and anaerobic biosynthesis of nano-selenium for remediation of mercury contaminated soil. Chemosphere 170, 266–273. https://doi.org/10.1016/j.chemosphere.2016.12.020
- [28] Wilson, J.R., Leang, C., Morby, A.P., Hobman, J.L., Brown, N.L., 2000. MerF is a mercury transport protein: different structures but a common mechanism for mercuric ion transporters? FEBS Lett. 472, 78–82. https://doi.org/10.1016/S0014-5793(00)01430-7
- [29] Wolfe, M.F., Schwarzbach, S., Sulaiman, R.A., 1998. Effects of mercury on wildlife: A comprehensive review. Environ. Toxicol. Chem. 17, 146–160. https://doi.org/10.1002/etc.5620170203
- [30] Yin, K., Wang, Q., Lv, M., Chen, L., 2019. Microorganism remediation strategies towards heavy metals. Chem. Eng. J. 360, 1553–1563. https://doi.org/10.1016/j.cej.2018.10.226
- [31] Zhu, X., Kusaka, Y., Sato, K., Zhang, Q., 2000. The endocrine disruptive effects of mercury. Environ. Health Prev. Med. 4, 174–183. https://doi.org/10.1007/BF02931255