

Chapter 6

Bioremediation Approaches for Degradation and Detoxification of Polycyclic Aromatic Hydrocarbons



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Abstract Waste from industry is a noteworthy risk to the earth as it contains different poisonous, mutagenic and cancer-causing substances including polycyclic aromatic hydrocarbons (PAHs). PAHs are a class of different organic compounds with two or more intertwined benzene rings in a linear, angular or cluster array. Eviction of PAHs is crucial as these are persevering toxins with ubiquitous event and adverse natural impacts. There are several remedial techniques, which are productive and financially savvy in elimination of PAHs from the affected environment. These removal approaches are not just eco-friendly; they additionally display an emerging and new strategy in mitigating the ability of PAHs to cause potential risk to living beings. Accessible physical and synthetic techniques are neither eco-accommodating nor financially viable in this way. Natural strategies such as bioremediation techniques are most appropriate for biodegradation of PAHs. Such techniques require less chemicals, less time and less contribution of energy and are cost-effective and eco-accommodating. The lethal PAH mixes can be changed into non-harmful and more straightforward ones utilizing normally occurring microorganisms like algae, bacteria and fungi in a procedure called biodegradation. This chapter mainly focuses on the enhancement in biodegradation of hazardous PAHs by using bioremedial approaches.

Keywords Polycyclic aromatic hydrocarbons · Enzymatic approach · Biodegradation · Bioremediation

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1 Introduction

PAHs are effective environmental toxicants that consist of fused aromatic rings. PAHs are originated in unrefined petroleum, coal tar and blacktop (Ukiwe et al. 2013). PAHs are included in the US Environmental Protection Agency (EPA) and European Community (EC) Contaminant Candidate List. EPA currently regulates 16 PAH compounds as priority pollutants in water and as 'aggregate PAHs' in defiled soil and sediments (Hadibarata et al. 2009). PAHs are of big concern to humans and animals as contaminant, some even recognized as cancer causing, mutagenic or teratogenic.

PAH compounds have two- to seven-membered benzene rings. They are lacking water affinity mixes with aqueous solubility declining almost linearly with increases in molecular mass (Parrish et al. 2004). Physicochemical properties and molecular weight of PAHs vary with the number of rings in the atom. Increment in subatomic weight of PAHs leads to decrease in chemical reactivity, aqueous solubility and volatility of PAHs. High-molecular-weight PAHs have high resonance energies because of the thick mists of pi-electrons surrounding the aromatic rings making them steady in the earth and recalcitrant to degradation. The recalcitrant nature of compounds may be attributed to their low water solubility and high soil sorption (Parrish et al. 2004).

PAH degradation is controlled by several physicochemical as well as biological processes, which vary their fate and transport in the surface environment. Biodegradation of hydrocarbons is achieved either by microorganism such as bacteria (Hamamura et al. 2013), fungus (Cerniglia and Sutherland 2010) or algae (Chan et al. 2006) or by enzymatic approaches. Fungi are considered as a productive competitor for effective degradation of PAHs. In any case, filamentous growth has capacity to develop on wide spectrum of substrates by secreting extracellular hydrolytic enzymes, even equipped for growing under non-ambient environment (Juhasz and Naidu 2000). Bioremediation includes either indigenous or exogenous microbial population, which is known as not proficient degraders in contaminated site (Yadav et al. 2017; Bharagava et al. 2017). Fungi have advantages over bacteria because of their fungal hyphae and potent hydrolytic enzymes, which can enter and corrupt the hydrocarbons affected environment (Venkatesagowda et al. 2012). Fungal enzymes especially oxidoreductases, laccase and peroxidases have noticeable application in removal of PAH contaminants either in fresh, marine water or terrestrial. Nevertheless, interest on growths gets an impressive consideration for bioremediation of hydrocarbon contaminated sites associated fungi for enzyme discharge (to expel hydrocarbons from nature). The persistence, toxicity and carcinogenicity of PAH molecules draw public concern to decontaminate PAH-polluted sites.

2 Sources of Polycyclic Aromatic Hydrocarbons in the Environment

For the most part, PAH contamination happens by unprocessed and processed oil, which comes from tanker accidents, refinery effluents, metropolitan and modern release from pipelines and seaward productions and waste oil from two-wheeler and four-wheeler administration stations, which causes contamination (Uzoamaka et al. 2009). Polycyclic aromatic hydrocarbon compounds are formed, what's more, discharged into the earth through both natural and anthropogenic sources. Natural sources of PAHs include their development as exudates from trees woods and rangeland fires, fungi and bacteria (Fig. 6.1). In nature, PAHs remain prevalently distributed as parts of plant oils, cuticles of insects, components of surface waxes of leaves and lipids of microorganisms. PAHs are formed naturally during thermal geologic reactions associated with fossil fuel and mineral generation.

Anthropogenic sources like fuel ignition, vehicles, spillage of petroleum products, electric fuel generation, internal ignition motors and waste incinerators are critical sources of PAHs into the environment (Arulazhagan and Vasudevan 2011). Anthropogenic wellsprings of PAHs are the real reason for natural contamination and, hence, the focus of a lot of bioremediation programmes. PAHs remain saved in the environment through generally scattered sources covering significantly the land surface area. At such sources PAHs are observed to be consumed strongly to soil particles.

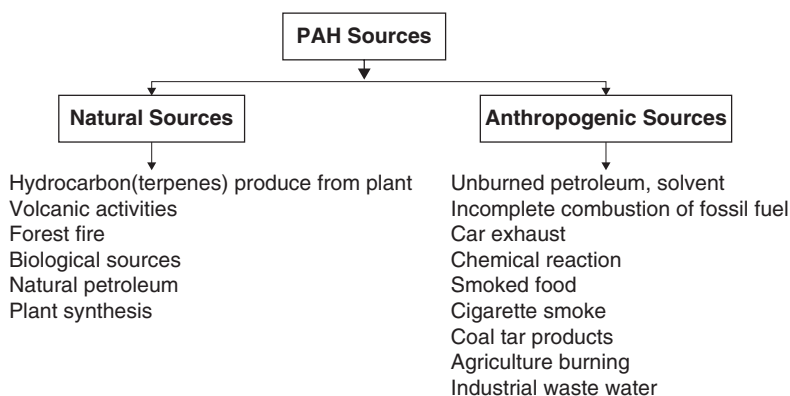


Fig. 6.1 Natural and anthropogenic sources of polycyclic aromatic hydrocarbons (PAHs)

3 Environmental Problems of PAHs

Increasing knowledge of potent adverse effects of toxicant on human health and environment has led to enhanced attention and measures for remediation and renovation of environment contaminants. Polycyclic aromatic hydrocarbons (PAHs) are omnipresent organic pollutants with serious environmental concerns. PAHs are distributed in various ecosystems and are pollutants of severe concern due to their potential toxicity, mutagenicity and carcinogenicity.

PAHs have an innate property for bioaccumulation in food chains, which makes their presence in any ecosystem alarming for human health (Morehead et al. 1986; Xue and Warshawsky 2005). Accumulation of PAHs in plants poses hazardous effect to human health because of their position in the food chain. Experimental studies have also demonstrated interference with plant carbon allocation and root symbioses by PAHs, which ultimately affect plant growth and the environment. Because of their concentration and toxicity, 16 PAHs have been enlisted as priority environmental pollutants by the US Environmental Protection Agency (US EPA).

4 Various Techniques for PAH Degradation

4.1 Chemical Degradation

The availability of PAHs in anaerobic conditions relies on certain components, which incorporate substrate interaction, pH and redox conditions (Chang et al. 2002). The most oxidation reactions in the environment are initiated by oxidants such as peroxides (H_2O_2), ozone (O_3) and hydroxyl radicals generated by photochemical processes. The degradation pathways are such that the oxidation reactions involving hydroxyl radicals or O_3 react with aromatic compounds such as PAHs at near diffusion-controlled rates by abstracting hydrogen atoms or by expansion to twofold bonds (Ukiwe et al. 2013). The reaction proceeds with complex pathways creating various intermediates. In such reactions, the last response products include a mixture of ketones, quinones, aldehydes, phenols and carboxylic acids for both oxidants (Reisen and Arey 2002). During chemical reaction PAHs are transformed into other polyaromatic hydrocarbons (they do not lose their aromatic character). Their aromaticity is conserved since considerable amounts of energy are required to change an aromatic compound into a non-aromatic compound. PAHs could be degraded through fermentative digestion system, while some studies have demonstrated that PAH degradation in anaerobic environment is much slower than in aerobic environment (Ambrosoli et al. 2005). Effects of carbon (C) and nitrogen (N) on PAH degradation have also been investigated by several authors (Quan et al. 2009). The efficiency of PAH chemical degradation is limited by their low aqueous solubility and vapour pressure (Fernando et al. 2009). However, surfactants enhance the

solubility of hydrophobic compounds (Ukiwe et al. 2013). Several reports have been focused on the significance of surfactants to expand the solubility of PAHs by decreasing the interfacial surface tension amongst PAH and the dirt/water inter-phase (Li and Chen 2009).

4.2 Phytodegradation

It is characterized as the utilization of plants to expel contaminations from the earth to render them nontoxic (Table 6.1). Plants can take the toxicant up and accrue them in their tissues. It is an in situ, solar energy-regulated technique, which minimizes environmental disturbance and reduces costs (Haritash and Kaushik 2009). Researchers have indicated that various grasses and leguminous plants are potential candidates for phytodegradation of organics (Newman and Reynolds 2004; Ukiwe et al. 2013). Some tropical plants have also been reported to show effective degradation tendency due to inherent properties such as deep fibrous root system and tolerance to high hydrocarbon and low nutrient availability (Dzantor et al. 2000; Chandra et al. 2012). Many species of grass such as *Agropyron smithii*, *Bouteloua gracilis*, *Cynodon dactylon*, *Elymus canadensis*, *Festuca arundinacea*, *Festuca rubra*, *Melilotus officinalis*, etc. are known to degrade PAHs (McCutcheon and Schnoor 2003). Researchers are also investigating that grasses and legumes induce the removal of PAHs from affected soil. Plants also play an indirect role in the removal of PAHs by releasing of enzymes by roots. These enzymes are capable of transforming organic contaminants by catalysing chemical reactions in soil (Ndimele et al. 2010). Plant enzymes also act as causative agents in the transformation of contaminants mixed with sediment and soil. The identified enzyme systems included dehalogenase, nitroreductase, peroxidase and laccase (Thomson and Ndimele 2010). Rasmussen and Olsen studied the efficiency of orchard grass (*Dactylis glomerata*) towards PAH removal. The study reported that a soil/sand mixture vegetated with orchard grass exhibited high treatment efficiency with an input from the microbial catabolic degradation by plant exudates (Parish et al. 2004).

Table 6.1 Plants useful in phytodegradation of PAHs

S.No.	Isolate name	Compound	% Removal	Incubation	References
1.	<i>Festuca arundinacea</i>	Pyrene	38%	190 days	Chen et al. (2003)
2.	<i>Panicum virgatum</i>	Pyrene	38%	190 days	Chen et al. (2003)
3.	<i>E. crassipes solani</i>	Naphthalene	45%	7 days	Nesterenko et al. (2012)
4.	<i>Scirpus Grossus</i>	Petroleum hydrocarbons	81.5%	72 days	Al-Baldawi et al. (2015)

4.3 Biodegradation

Biodegradation is a reasonable technique for degradation of natural contaminations. It is the use of microorganisms to degrade or detoxify environmental pollutants (Bamforth and Singleto 2005; Saxena and Bharagava 2017). Several additional factors affecting rate of degradation incorporate pH, temperature, nearness of oxygen and supplement accessibility. The concentration of nutrients and the state of the nutrients (organic, inorganic) are important for biodegradation. The biodegradation of PAHs is financially savvy, eco-friendly (as it prevents environmental damage during transportation of contaminants). Biological degradation is an approach that presents the possibility to remove organic pollutants with the help of natural biological activity available in the substrate (Zeyauallah et al. 2009; Bharagava and Chandra 2010). The microorganisms used for biodegradation could be indigenous to the contaminated region or site (Das and Chandran 2011). The complete mineralization products of the pollutant by biodegradation process include CO₂, H₂O and cell biomass (Gratia et al. 2006). During biodegradation process optimization involves many factors such as microbial consortia capable of degrading the pollutant, bioavailability of the pollutants to microbial attack and soil type, temperature, soil pH, oxygen level of soil, electron acceptor agents and nutrient content of soil (environmental factors) contributing to microbial growth (Gratia et al. 2006; Epelde et al. 2009; Mulla et al. 2017; Bharagava and Chandra 2010). Complete degradation of PAHs to CO₂, water, microbial carbon and other inorganic compound is the ultimate goal. Haeseler et al. (2001) showed enhanced, but incomplete, degradation of PAH compounds in a field study. When remediation was complete, final toxicity was very less because the metabolites tended to be less stable and more soluble than the parent compounds, making them more available to degraders (Haeseler et al. 2001). Microbial degradation is the most suitable alternative and effective method of removal of those toxic chemicals. Microbes (including bacteria, fungi and algae) can biologically degrade PAH compounds during direct microbial metabolism of carbon energy sources or by co-metabolism while consuming another substrate (Lundstedt et al. 2006) (Table 6.2).

5 Role of Various Microorganisms in Bioremediation

The problem linked with the PAHs can be mitigating by the use of conventional approaches, which involve degradation, modification or isolation of the toxicant. These approaches involve excavation of contaminate and its incineration or containment. These technologies are expensive and in many cases transfer the pollutant from one phase to another. On the other hand, bioremediation is the tool to transform the compounds to less hazardous/nonhazardous forms with less input of chemicals, energy and time (Ward et al. 2003; Yuan et al. 2001; Chandra et al. 2011). Microorganisms are known to be their catabolic activity in biological remediation

Table 6.2 Movement and fate of organic chemicals, such as PAHs, in the environment

S.No.	Degradation process	Consequence	Factors	Advantages/ disadvantages
1.	Chemical degradation	Alteration of PAHs by chemical processes such as photochemical (i.e., UV light) and oxidation-reduction reactions	High and low pH, structure of PAH, intensity and duration of sunlight, exposure to sunlight and same factors as for microbial degradation	Limited effectiveness and can be expensive
2.	Phytodegradation	Breakdown of contaminants or pesticides through metabolic processes within the plant	Molecular weight of PAHs, sunlight, enzymes	Low-cost methods for cleaning the environment
3.	Biodegradation	Degradation of PAHs by microorganisms, biodegradation and co-metabolism	Environmental factors (pH, moisture, temperature, oxygen), nutrient status, organic matter content, PAH bioavailability, microbial community present, molecular weight of PAH (LMW or HMW)	Cost-effectiveness and complete cleanup, catabolic versatility of microorganisms, less labour-intensive, relying on solar energy, have a lower carbon footprint and have a high level of public acceptance

(Bharagava et al. 2009), but changes in microbial communities are still unpredictable, and the microbial community is still termed as a ‘black box’ (Dua et al. 2002). The PAH-degrading microorganism could be algae, bacteria or fungi. It involves the breakdown of organic compounds either usually by microorganism in to less complex metabolites or through mineralization into inorganic minerals, H₂O, CO₂ (aerobic) or CH₄ (anaerobic). The extent and rate of contaminant degradation depend on many factors including pH, temperature, O₂, microbial population, degree of acclimation, convenience of nutrients, compounds chemical structure, properties of cellular transport and chemical partitioning in the growth medium (Singh and Ward 2004).

5.1 Biodegradation of PAHS by Fungi

Fungus is known to have the properties of degradation of persistent organic pollutants (Table 6.3). Distinct properties differentiate filamentous fungus from other life forms to decide why they are potent biodegraders agents. First, the mycelial growth habit gives a competitive benefit over single cells such as bacteria and yeasts, especially with respect to the colonization of insoluble substrates (Bennet et al. 2002). The isolates identified as *Deuteromycetes* belonging to the genera *Cladophialophora*, *Exophiala* and *Leptodontium* and the ascomycete *Pseudeurotium zonatum* are

Table 6.3 Fungal isolates involve in degradation of PAHs

S.No	Isolate name	Compound	% Removal	Incubation	References
1.	<i>Phomopsis liquidambari</i>	Indole	41.7%	6 days	Chen et al. (2013)
2.	<i>Fusarium verticillioides</i>	Naphthalene	87.78%	8 days	Mohamed et al. (2012)
3.	<i>Fusarium solani</i>	Anthracene, benz[a]anthracene	40% and 60%, resp.	40 days	Wu et al. (2009)
4.	<i>Aspergillus terreus</i>	Naphthalene, anthracene	98.5% and 91%, resp.	4 weeks	Mohamed et al. (2012)
5.	<i>Fusarium</i> sp.	Naphthalene	42%	7 days	Ahirwae and Dehariya (2013)

toluene-degrading fungi; they use toluene as sole carbon and energy source (Francesc et al. 2001). Clemente et al. (2001) reported that degree of degradation of PAH varies with a variation of lignolytic enzymes producing deuteromycete ligninolytic fungal isolates.

Low-molecular-weight PAHs (two to three rings) were found to be degraded most extensively by *Aspergillus* sp., *Trichocladium canadense* and *Fusarium oxysporum*. For high-molecular-weight PAHs (four to seven rings), maximum degradation has been observed by *T. canadense*, *Aspergillus* sp., *Verticillium* sp. and *Acremonium* sp. Such studies have found that fungi have a great capability to degrade a broad range of PAHs under low-oxygen conditions. As a large and novel microbial resource, endophytic fungi have been paid more attention in their ecological functions.

The effect of microbes on litter component decomposition (Osono and Hirose 2011) but extended fungal degradation to more recalcitrant carbohydrate (Russell et al. 2011).

PAHs degradation by fungi has mostly focused on white-rot fungi. Their broad-range degradation potential is one reason of PAHs, such as *Irpex lacteus* found with a degradative ability of ANT, phenanthrene, pyrene as well as fluoranthene, and their degradative mechanisms were also investigated (Cajthaml et al. 2002). The other reason is because of their efficient production of ligninolytic enzymes; e.g. *Phanerochaete chrysosporium* could degrade ANT and phenanthrene by producing lignin peroxidase (LiP) and manganese-dependent peroxidase (MnP) (Hammel 1995). *Lentinus (Panus) tigrinus* showed out the MnP transformation ability after carrying out in vivo and in vitro degradation of PAHs (Covino et al. 2010). *Cunninghamella* sp. and *Aspergillus* sp. were reported for their potential in the transformation of benzo[a]pyrene and the conjugation mechanisms during the degradation (Wu et al. 2009). *Fusarium* spp. have shown their capability to degrade high-molecular-weight organic compounds such as coal cellulose, xylan, pectin,

different hydrocarbons (Kang and Buchenauer 2000) as well as PAHs (Chulalaksananukul et al. 2006). Lignin peroxidase (LiP), manganese peroxidase (MnP) and laccase (Lac) have shown to degrade not only lignocellulose but also pollutants such as crude oil wastes, textile effluents, distillery wastewater pollutants, organochloride agrochemicals and pulp effluents which are a cause of serious environmental pollution (Mtui and Nakamura 2004; Chandra and Chowdhary 2015; Chowdhary et al. 2017a, b, 2018).

5.2 Biodegradation by Bacterial Isolates

Several reports refer to degradation ability of different bacterial isolates of environmental pollutants (Table 6.4). Many bacterial spp. are even known to nourish completely on hydrocarbons (Yakimov et al. 2007). Degradation of PAHs can occur under aerobic and anaerobic conditions, as in the case for the nitrate-reducing bacterial strains *Pseudomonas* sp. and *Brevibacillus* sp. isolated from petroleum contaminated soil (Grishchenkov et al. 2000). Several species of microorganisms have been successfully utilized in major hazardous waste cleanup processes (Levinson et al. 1994). Abd et al. (2009) reported that two- to three-ring PAHs (naphthalene, anthracene and phenanthrene) can be degraded using *Pseudomonas geniculata* and *Achromobacter xylooxidans*. Bacterial isolates capable of chrysene metabolism include *Rhodococcus* sp. strain UW1 (Walter et al. 1991) and *Sphingomonas yanokuyae* which oxidized chrysene (Boyd et al. 1999), while *Pseudomonas fluorescens* utilized chrysene and benz[a]anthracene as sole carbon sources (Caldini et al. 1995). The microorganisms capable of surviving in such a polluted environment are those that develop specific enzymatic and physiological responses that allow them to use hydrocarbon as a substrate. Kafilzadeh et al. (2011) reported ten genera as follows: *Bacillus*, *Corynebacterium*, *Staphylococcus*, *Streptococcus*, *Shigella*, *Alcaligenes*,

Table 6.4 Biodegradation of PAHs by various bacterial isolates

S.No	Isolate name	Compound	% Removal	Incubation	References
1.	<i>Pseudomonas</i> sp.	Naphthalene, fluorene	95%	4 days	Kumar et al. (2010)
2.	<i>Pseudomonas</i> sp.	Anthracene	74.8%	10 days	Kumar et al. (2010)
3.	<i>Bacillus</i> sp.	Anthracene	82.6%	72 h	Neelofur et al. (2014)
4.	<i>Mesoflavibacter zeaxanthinifaciens</i>	Benzo[a]pyrene	86%	42 days	Okai et al. (2015)
5.	<i>Mycobacterium flavescens</i>	Pyrene	89.4%	2 weeks	Dean-Ross et al. (2002)
6.	<i>Rhodococcus</i> sp.	Anthracene	53.0%	2 weeks	Ross et al. (2002)

Acinetobacter, *Escherichia*, *Klebsiella* and *Enterobacter* out of 80 bacterial strains. *Bacillus* was the best hydrocarbon-degrading genus. Bacterial strains that are able to degrade aromatic hydrocarbons have been repeatedly isolated, mainly from soil. The bacterial genera *Mycobacterium*, *Corynebacterium*, *Aeromonas*, *Rhodococcus* and *Bacillus* have been also reported for biodegradation pathways (Mrozik et al. 2003).

5.3 Biodegradation of PAHs by Algal Isolates

Algae are important microbial members in both aquatic and terrestrial ecosystems; reports are insufficient regarding their concern in hydrocarbon biodegradation (Table 6.5). (Das and Chandran 2011; Walker et al. 1975) isolated an alga, *Prototheca zopfii* which was capable of utilizing crude oil and a mixed hydrocarbon substrate and exhibited extensive degradation of n-alkanes and isoalkanes as well as aromatic hydrocarbons. Cerniglia and Gibson (1977) observed that nine cyanobacteria, five green algae, one red alga, one brown alga and two diatoms could oxidize naphthalene. Some research has demonstrated that certain fresh algae (e.g., *Chlorella vulgaris*, *Scenedesmus platydiscus*, *S. quadricauda* and *S. capricornutum*) are capable of uptaking and degrading PAHs (Wang and Zhao 2007). Warshawsky et al. (2007) found that *Selenastrum capricornutum*, a freshwater green alga, metabolizes BaP to cis-dihydrodiols using a dioxygenase enzyme system as found in heterotrophic prokaryotes. Certain algae have been reported to enhance the removal fluoranthene and pyrene when present with bacteria. Borde et al. (2003) first reported that photosynthesis enhanced degradation of toxic aromatic compounds by algal-bacterial microcosms in a one-stage treatment. *Pseudomonas migulae* and *Sphingomonas yanoikuyae* were studied for phenanthrene degradation. The green alga *Chlorella sorokiniana* was cultivated in the presence of the pollutants at different concentrations, showing increasing inhibitory effects in the order salicylate < phenol

Table 6.5 Biodegradation of PAHs by various algal isolates

S.No	Isolate name	Compound	% Removal	Incubation	References
1.	<i>Prototheca zopfii</i>	Petroleum hydrocarbons	12–41%	3 days	Kirk et al. (1999)
2.	<i>Selenastrum capricornutum</i>	Benzo[a]pyrene	41%	4 days	Warshawsky et al. (2007)
3.	<i>Selenastrum capricornutum</i>	Fluoranthene	99%	7 days	Ke et al. (2010)
4.	<i>Lyngbyala gerlerimi</i>	Naphthol	36.6%	5 days	Mostafa et al. (2012)
5.	<i>Nostoc linckia</i>	Catechol	56.38%	5 days	Mostafa et al. (2012)
6.	<i>Oscillatoria rubescens</i>	β-naphthol	3.04%	7 days	Mostafa et al. (2012)

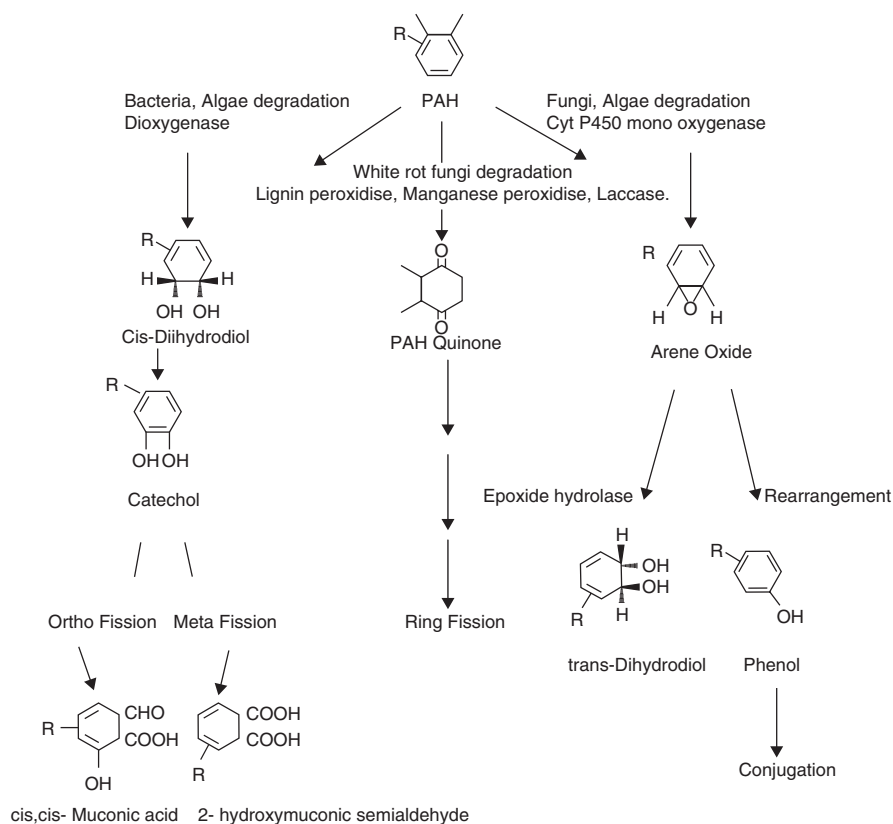


Fig. 6.2 The three main pathways for polyaromatic hydrocarbon degradation by fungi and bacteria. (Adopted by Muthuswamy et al. 2008)

<phenanthrene (Lei et al. 2007). The study of fluoranthene, pyrene and a mixture of fluoranthene and pyrene by *Chlorella vulgaris*, *Scenedesmus platydiscus*, *Scenedesmus quadricauda* and *Selenastrum capricornutum* has shown that removal is algal species-specific and toxicant-dependent. PAH removal in 7 days of treatment was 78% and 48%, respectively, by *S. capricornutum* and *C. vulgaris*. Hong et al. (2008) studied the accumulation and biodegradation of phenanthrene and fluoranthene by the algae enriched from a mangrove aquatic environment (Fig. 6.2).

6 Enzymatic Degradation of PAHs

Major enzymes useful in PAH degradation belong to oxygenase, dehydrogenase and lignolytic categories. Lignolytic enzymes secreted by majority of fungal species are lignin peroxidase, manganese peroxidase and laccase, which are

extracellular in nature and are secreted for catabolism of substrate food material (Chandra and Chowdhary 2015). Spent mushroom compost (SMC) is often used as an inoculum source which enhances the rate of PAH degradation. The SMC are high in laccase and Mn-dependent peroxidase, whereas the production of ligninase is reported to be low in SMC (Haritash and Kaushik 2009). In which most of enzymes are active at many temperature and have optimum activity at mesophilic temperatures and it reduces with very high and very low temperatures. Some of the enzymes are reported to be active even at extremes of temperatures (Haritash and Kaushik 2009). Enzymes also show substrate specificity, but ligninolytic enzymes are non-specific in nature, acting on phenolic and non-phenolic organic compounds via the generation of cation radicals after one e^- oxidation (Lau et al. 2003).

6.1 Lignin Degrading Enzymes

Oxidoreductive enzymes play a key role in transformation and degradation of polymeric substances (Table 6.6). The less degraded or oxidized items can without much of a stretch be taken up by microbial cells where they are totally mineralized. A class of oxidoreductive enzymes include lignin-degrading compounds (LDEs) which have practical application in bioremediation of polluted environment (Husain 2006). LDEs belong to two classes, viz., the heme-containing peroxidases and the copper-containing laccases. A progression of redox responses are started by the laccases. LDEs degrade the lignin or lignin-derived pollutants. The LDEs oxidize the aromatic compounds until the aromatic ring structure is cleaved, which is followed by further debasement with different compounds.

Table 6.6 Biological functions of ligninolytic enzymes

Enzymes	Applications	References
Lignin peroxidase	Biodegradation of lignin	Martínez et al. (2005)
	Degradation of azo	Stolz (2001)
	Mineralization of environmental contaminants	Harms et al. (2011)
	Degradation of pharmaceuticals and their metabolites	Marco-Urrea et al. (2009)
Manganese peroxidase	Degradation of lignin	Martínez et al. (2005)
	PAH degradation	Baborová et al. (2006)
	Synthetic dyes, DDT, PCB, TNT	Hernández et al. (2008)
	Textile dye degradation and bleaching	Kalyani et al. (2008)
Laccase	Spore resistance	Lu et al. (2012)
	Rhizomorph formation	Ranocha et al. (2002)
	Pathogenesis	Langfelder et al. (2003)
	Fruit bodies formation	Nagai et al. (2003)
	Pigment synthesis	Eisenman et al. (2007)

Enzymes involved in the degradation of PAHs are oxygenase, dehydrogenase and ligninolytic enzymes. Fungal ligninolytic enzymes are lignin peroxidase, laccase and manganese peroxidase. They are extracellular and catalyse radical formation by oxidation to destabilize bonds in a molecule (Hofrichter et al. 1999).

Peroxidases perform heme-containing degradation with other enzymes. Peroxidases are heme-containing enzymes that comprise manganese-dependant peroxidase (MnP), lignin peroxidase (LiP) and versatile peroxidase (VP). They oxidize lignin subunits using extracellular hydrogen peroxide generated by unrelated oxidases as co-substrate. Most mineralization activity of the lignin polymers to CO₂ and H₂O in terrestrial ecosystem is performed by fungal species. These fungi produce a wide range of lignin-degrading enzymes (LDEs), which in turn act on lignin and lignin-analogous compounds. PAHs are primarily degraded using extracellular oxidative enzymes, although use of laccases and peroxidases in PAH bioremediation is currently being studied (Harms et al. 2011). The white-rot fungi (WRF) belonging to the basidiomycetes produce various isoforms of extracellular ligninolytic enzymes, laccases (Lac) and different peroxidases, including lignin peroxidase (LiP), manganese peroxidase (MnP) and versatile peroxidase (VP), the latter sharing LiP and MnP catalytic properties (Martínez 2002). The natural substrate of these enzymes (lignocellulose) is degraded in the environment by the WRF, along with various xenobiotic compounds, including dyes (Wesenberg et al. 2003). Some WRF produce all the three lignin-modifying enzymes, while others produce only one or two of them. Lignin-modifying enzymes are produced by WRF during their secondary metabolism since lignin oxidation provides no net energy to the fungus (Eggert et al. 1996).

6.1.1 Lignin Peroxidases (EC 1.11.1.14)

LiPs (EC 1.11.1.14) are an extracellular hemeprotein. They are related to the family of oxidoreductases (Higuchi 2004). LiP has high redox potential and low optimum pH (Piontek et al. 2001) and is capable of oxidizing a variety of reducing substrates including polymeric substrates (Oyadomari et al. 2003). Due to their high redox potentials and enlarged substrate range, LiPs have more potential for application in several industrial processes (Erden et al. 2009). Enzymatic activity of LiP is H₂O₂ dependent; here H₂O₂ gets reduced to H₂O by picking up an electron from LiP (which itself gets oxidized). The oxidized LiP then returns to its native reduced state by picking up an e⁻ from veratryl alcohol and oxidizing into veratryl aldehyde. Veratryl aldehyde then gets reduced back to veratryl alcohol by picking up an electron from lignin or equivalent structures such as xenobiotic compounds.

6.1.2 Manganese Peroxidases (EC 1.11.1.13)

MnP (EC 1.11.1.13) belong to the family of oxidoreductases (Higuchi. 2004). Studies have shown that MnP is distributed in almost all white-rot fungi (Hofrichter 2002). Manganese peroxidases (MnP) seem to be distributed amongst white-rot

fungi than LiP (Hammel and Cullen. 2008). MnP oxidizes Mn^{2+} to highly reactive Mn^{3+} , which oxidizes phenolic structures to phenoxyl radicals (Hofrichter 1999). The Mn^{3+} forms complex with chelating organic acids resulting in products such as oxalates or malates (Makela et al. 2002). The redox potential of the Mn peroxidase system is lower than that of lignin peroxidase, and it has shown capacity for preferable oxidation of phenolic substrates. On the other hand, studies indicate that contrary to LiP, MnP may oxidize Mn (II) without H_2O_2 with decomposition of acids and concomitant production of peroxy radicals that may affect lignin structure. Versatile peroxidase (VP) enzymes produced by *Pleurotus* spp. are also able to oxidize phenolic compounds and dyes efficiently that are substrates of generic peroxidases and related peroxidases or the well-known horseradish peroxidase (HRP). VP (EC 1.11.1.16) oxidizes Mn^{2+} , similar to MnP, and have a high redox potential aromatic as LiP enzymes. Due to these qualities, interest in VP has increased during the last years (Martínez et al. 2009).

6.1.3 Laccases (EC 1.10.3.2)

Laccases (EC 1.10.3.2) belong to a multicopper oxidase family (Alcalde 2007), present in bacteria, e.g., *Azospirillum lipoferum*, *Actinomyces* like *Streptomyces*, fungi, plants and insects (Baldrian 2006; Chandra and Chowdhary 2015). This enzyme had been reported more than a hundred years ago (Desai and Nityanand 2011), but the significance and broad studies over the role of this enzyme in wood degradation have been conducted in the last few decades. However, many laccases were reported from fungi, and most biotechnologically useful laccases also originated from fungus (Kalmis et al. 2008).

7 Challenges

In spite of considerable progress made in the study of the biodegradation of PAHs over the past few decades, removal of petroleum hydrocarbons in the environment is a daunting problem of the real world. Advancement in various approaches such as genomics, proteomics and metabolomic study has contributed immensely in understanding the PAH-degrading microorganisms and the biochemistry involved in the degradation pathways; however, challenges are posed by various aspects of PAH bioremediation which are either unknown or insufficient information is available regarding them.

Little or no information is available related to genes, enzymes and molecular mechanism of PAH degradation in high-salt environments or low-oxygen and anaerobic environments. Scarce data and research are there on the transmembrane trafficking of PAHs and their metabolites; no transporter molecule/protein has been characterized till date with specific role in the transport of PAHs into microorganisms.

Thorough understanding of genetic regulation of the pathways involved in PAH degradation by different bacteria and fungi has been used for efficient biotransformation or metabolism of PAH pollutants in recent past; a deeper understanding of the microorganism-mediated mechanisms of PAH catalysis will enable strategizing novel methods to enhance the bioremediation of PAHs in the environment.

The use of genetically modified organism in bioremediation represents a research frontier with broad implications to improve the degradation of hazardous wastes under laboratory conditions. The potential benefits of using genetically modified microorganisms are significant. Combining genetic engineering tools such as gene conversion, gene duplication and mutation, enzyme overexpression and novel strains can be produced with desirable properties for effective bioremediation applications. Ecological and environmental concerns and regulatory constraints pose major obstacles and challenges for testing genetically modified organism in the field. These problems must be solved before a genetically modified organism can provide an effective, safer and more efficient method than the present alternatives for removal process at very low-cost and eco-friendly way.

8 Conclusion

The present status of work done on biodegradation of biologically toxic PAHs using different microorganism has been reviewed. The environmental toxicity and persistence of PAHs have resulted in several laboratory-based experiments to change these substances into less unsafe/nondangerous substances with the use of microorganisms in the process called as biodegradation. Removal of PAHs from the affected environment is a tough job. Therefore, it is very essential to understand the mechanism of several degradation processes. Degradation of PAHs remains affected by numerous factors, which need to be addressed and explored. Biological approaches appear be the most efficient, cost-effective and eco-friendly method to mitigate/remove PAHs from affected area.

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