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Insights into the Chemical Interaction between Plants and Microbes and its Potential Use in Soil Remediation

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Abstract

Soil bacteria are very vital and they are frequently used in production of crop. Chemical dialogues between bacteria and plant roots result in the proliferation and biofilm formation of plant growth promoting and contaminant degrading bacteria. Plant-bacterial interactions in the rhizosphere are the determinants of plant health and soil fertility. Plant growth promoting rhizobacteria (PGPR) which is also known as plant health promoting rhizobacteria (PHPR) or nodule promoting rhizobacteria (NPR). It can benefit the host plant directly by enhancing plant growth or indirectly by producing hydrolytic enzymes and by priming plant defence. This review elaborates the effect of plant and bacterial products on the remediation of contaminated soil.

1. Introduction

Terrestrial ecosystem comprises numerous niches among which the association between plants and microbes plays an integral role [1]. Plant-microbe interactions are of various types and include mutualism, commensalism, parasitism and antagonism [2]. The most common interactions include mutualism and commensalism, where one or both participating organisms benefit from them [3].

Rhizosphere contains the root system of plants. It exhibits a flexible morphology and structure that help to respond to any change in the surrounding environment. The health and productivity of plant is influenced by its rhizosphere [4]. The root system provides anchorage, helps in nutrients uptake and facilitates many underground interactions [5]. These interactions comprise a mutualistic

relationship with plant associated bacteria, such as rhizobia, mycorrhizae, endophytes, and PGPR [6, 7, 8]. Besides these interactions with beneficial microorganisms, plant roots also interact with pathogenic microbes [9]. The roots of plants release a variety of chemicals in a large amount which help to attract beneficial bacteria and in the fight against pathogenic ones, such as *Azospirillum*, *Bacillus*, *Pseudomonas* species [10, 11]. The aim of this article is to explore the hidden mechanism of key root exudate components responsible for the accelerated rhizodegradation of organic pollutants. It also intends to indicate the responses of soil microorganisms to different root exudate components.

2. Chemical Communication between Plants and Microbes

Rhizosphere is the narrow region of soil

where the root exudates of plants drive an intense microbial activity [12, 13]. Rhizobacteria colonize the roots of plants and stimulate their growth. Inoculation and colonization of plant roots with PGPR has been used as a way to enhance the growth of plants [14]. The signalling between plants and bacteria influence the bacterial colonization of plant roots which is a complicated multistep process [15, 16]. Bacterial traits that affect the process of root colonization include motility and components of bacterial surface, such as flagella and pili [17, 18]. The growth and physiology of rhizobacteria is influenced by specie specific exudates secreted by plants. These exudates contain various biochemical components such as carbohydrates, proteins, vitamins and different organic acids [19, 6, 20]. The composition of these exudates and the ability of bacteria to catabolize these chemicals are the key determinants of primary colonizers [21, 22, 23]. *Pseudomonas* is a bacterial specie which can compete for limited carbon source and is capable of catabolizing a wide range of plant root exudates. This ability has made *pseudomonas* one of the most successful root colonizers [24, 25].

Quorum sensing is a mechanism which is utilized by bacterial communities to sense and communicate with environment [26, 27]. It is a sensing mechanism based on cell population density. Bacterial cells, particularly the gram negative bacteria, release molecules such as N-acyl homoserine lactone (AHL) which can sense quorum [28, 29]. Plants have evolved various methods to identify and respond to AHLs. This helps in the establishment of symbiotic associations between bacteria and plants [30, 31, 32]. Plants perceive surrounding AHL molecules and modify their genetic expressions accordingly, which in turn changes their protein profile and finally

adjust development based on the nature of signals [30, 33].

Various direct and indirect mechanisms are used by certain AHL producing bacterial strains to stimulate the growth of plants. Such mechanisms include nitrogen fixation, synthesis of phytohormones, siderophores, removing toxic chemicals and fighting pathogenic microbes [33, 34, 35]. Besides this, these bacterial strains also enhance the bioavailability of mineral nutrients such as iron and phosphorous through the decomposition and mineralization of organic matter [25]. Various chemicals produced by these bacteria may also help to enhance the plant community. Roots contain epidermal extensions called root hairs which directly participate in nutrient and water absorption [7, 20]. The development of primary roots, root hairs and adventitious roots is strongly induced by AHL molecules. The acyl chain length of AHL molecules may influence the thickening of root hairs. Small chains of these molecules (C6-C8) do not modify the development of root hairs [36, 32]. Long chains of these molecules (C8-C12) attenuate the development of root hairs in a dose dependant manner [37, 31]. For example, C10- AHL molecule has been reported to stimulate the lateral root hair formation on the tip region. When these quorum signal molecules are exogenously applied on root hairs, they may produce a response similar to auxins. Their role in cell division is evident by the regulation of lateral root hair formation [38].

The array of signalling metabolites produced by microbe assemblages present in the soil have become an interesting and important subject for investigators. These metabolites affect the genetic expression of host plants [39]. Volatile organic compounds (VOCs) are among the very well-documented signalling molecules

produced by bacterial communities. These lipophilic compounds with low molecular weight serve as a chemical window to release information. These compounds are synthesized by various metabolic pathways [26]. Recent findings suggest a more crucial role of VOCs in microbial communication than their other volatile equivalents. For example, numerous VOCs are produced by rhizobacteria that may contain alkanes, alkenes, ketones, alcohol, terpenoids and sulphur compounds [39, 32].

3. Role of Root Exudates in Biodegradation

Recently, root exudation has gained significant attention as the process used for the biodegradation of hydrocarbons occurs in the rhizosphere. As shown in Fig. 1, root exudates act by enhancing the hydrocarbon degrader population of the rhizosphere and serve as a carbon source for microorganisms, thereby enhancing hydrocarbon degradation [11]. The concentration of root exudates was found to be negatively correlated with the concentration of hydrocarbons, that is, the higher the root exudate concentration the lower the concentration of hydrocarbons in that area. As the distance from the root increases, the biodegradation of phenanthrene also decreases [16]. Biodegradation reportedly remains 86% in the first 3mm from the roots and gradually decreases to 48% between 3-6mm. It declines to 36% between 6-9mm [26]. A number of external factors influence the chemical complexity of exudates including the size of plant, condition of soil and photosynthetic activity [40]. These factors are also species specific or genotype specific. These exudates can be differentially modified depending on the source of secretion. The potential of root exudates to cover the information about the events that mediate the communication

events in rhizosphere is evident from their strong specificity and complexity.

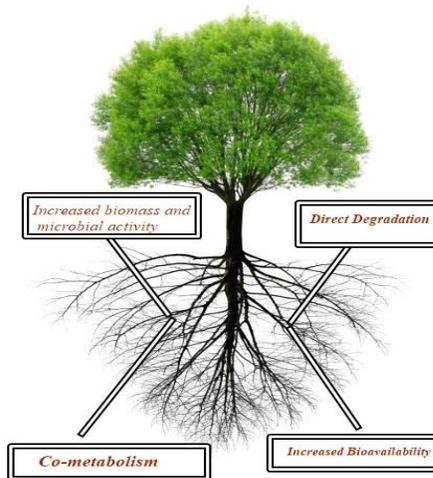


Figure 1. Improvement of petroleum hydrocarbon degradation by root exudates

4. Conclusion

Plants are environmentally safe, economically advantageous, and a robust renewable resource of *in situ* reduction of contaminants. Their root systems are generally extensive and provide a conducive environment for microbial propagation and activity, both within and outside of plant tissues. The combined use of root exudates, contaminant degrading rhizosphere and/or endophytic bacteria provides an effective approach for the remediation of contaminated soil.

References

- [1] Xu Y, Lu M. Bioremediation of crude oil-contaminated soil: comparison of different biostimulation and bioaugmentation treatments. *J Hazard Mater.* 2010;183(1): 395–401.
- [2] Ite AE, Ibok UJ. (2019). Role of plants and microbes in bioremediation of petroleum hydrocarbons contaminated

- soils. *Int J Environ Biorem Biodegrad.* 2019;7(1): 1–19.
- [3] Fatima K, Afzal M, Imran A, Khan QM. Bacterial rhizosphere and endosphere populations associated with grasses and trees to be used for phytoremediation of crude oil contaminated soil. *Bull Environ Contam Toxicol.* 2015;94 (3): 314–320.
- [4] Saleem H, Arslan M, Rehman K, Tahseen R, Afzal M. *Phragmites australis* a helophytic grass can establish successful partnership with phenol-degrading bacteria in a floating treatment wetland. *Saudi J Biol Sci.* 2019;26 (6), 1179–1186.
- [5] Afzal M, Rehman K, Shabir G, et al. Large-scale remediation of oil-contaminated water using floating treatment wetlands. *NPJ Clean Water.* 2019;2(1): 3.
- [6] Ajuzieogu CA, Ibiene AA, Stanley HO. Laboratory study on influence of plant growth promoting rhizobacteria (PGPR) on growth response and tolerance of *Zea mays* to petroleum hydrocarbon. *Afr J Biotechnol.* 2015;14(43): 2949–2956.
- [7] Dong R, Gu L, Guo C, Xun F, Liu J. Effect of PGPR *Serratia marcescens* BC-3 and AMF *Glomus intraradices* on phytoremediation of petroleum contaminated soil. *Ecotoxicology.* 2014;23(4): 674–680.
- [8] Swamy MK, Akthar MS, Sinniah UR. *Root exudates and their molecular interactions with rhizospheric microbes plant, soil and microbes.* New York: Springer; 2016.
- [9] Compant S, Saikkonen K, Mitter B, Campisano A, Mercado-Blanco J. Soil, plants and endophytes: editorial special issue. *Plant, Soil Environ.* 2016;405(1-2): 1–11.
- [10] Sasse J, Martinoia E, Northen T. (2018). Feed your friends: do plant exudates shape the root microbiome? *Trends Plant Sci.* 2018;23(1): 25–41.
- [11] Yuan J, Raza W, Shen Q. *Root exudates dominate the colonization of pathogen and plant growth-promoting rhizobacteria root biology.* New York: Springer; 2018.
- [12] Njoku KL, Akinola MO, Oboh BO. Phytoremediation of crude oil contaminated soil using *Glycine Max* (Merril); through phytoaccumulation or rhizosphere effect. *J Biol Environ Sci.* 2016;10(30): 115–124.
- [13] Omotayo AE, Ajayi A, Amund OO. Occurrence of hydrocarbon-degrading nitrogen-fixing bacteria in the rhizosphere of *Paspalum vaginatum* Sw. *J BioSci Biotechnol.* 2017;6 (2): 119–127.
- [14] Podile A, Kishore G. *Plant growth-promoting rhizobacteria: plant-associated bacteria.* New York: Springer; 2006.
- [15] Burghal AA, Al-Mudaffarand NA, Mahdi KH. *Ex situ* bioremediation of soil contaminated with crude oil by use of actinomycetes consortia for process bioaugmentation. *Eur J Experimental Biolo.* 2015;5,: 24–30.
- [16] dos Santos JJ, Maranhão LT. Rhizospheric microorganisms as a solution for the recovery of soils contaminated by petroleum: a review. *J Environ Manage.* 2018;210: 104–113.
- [17] Cheng L, Zhou Q, Yu B. Responses and roles of roots, microbes, and degrading genes in rhizosphere during phytoremediation of petroleum

- hydrocarbons contaminated soil. *Int J Phytorem.* 2019;21(12):1–9.
- [18] Kulkarni M, Chaudhari A. Microbial remediation of nitro-aromatic compounds: an overview. *J Environ Manage.* 2007;85(2), 496–512.
- [19] Adam M. Biodegradation of marine crude oil pollution using a salt-tolerant bacterial consortium isolated from Bohai Bay, China. *Mar Pollut Bull.* 2016;105(1): 43–50.
- [20] Hou J, Liu W, Wang B, Wang Q, Luo Y, Franks AE. PGPR enhanced phytoremediation of petroleum contaminated soil and rhizosphere microbial community response. *Chemosphere.* 2015;138: 592–598.
- [21] Alarcón A, Díaz MG, Cuevas LVH, ... et al. Impact of crude oil on functional groups of culturable bacteria and colonization of symbiotic microorganisms in the *clitoria-brachiaria* rhizosphere grown in mesocosms. *Acta Biol Colomb.* 2019;24(2): 343–353.
- [22] Lueders T. The ecology of anaerobic degraders of BTEX hydrocarbons in aquifers. *FEMS Microbiol Ecol.* 2017;93(1): fiw220.
- [23] Mommer L, Kirkegaard J, van Ruijven J. Root-root interactions: towards a rhizosphere framework. *Trends Plant Sci.* 2016;21(3): 209–217.
- [24] Flury P, Vesga P, Dominguez A-F, ... et al. Persistence of root-colonizing *Pseudomonas protegens* in herbivorous insects throughout different developmental stages and dispersal to new host plants. *ISME J.* 2019;13(4): 860.
- [25] Mavrodi DV, Yang M, Mavrodi OV, Wen S. (2017). Management of soilborne plant pathogens with beneficial root-colonizing *pseudomonas*. *Adv PGPR Res.* 2017;147: 2.
- [26] Dharmasiri R, Nilmini A, Undugoda L, Nugara N, Udayanga D, Manage P. Phenanthrene degradation ability of phyllosphere bacteria inhabiting the urban areas in Sri Lanka. In: *Proceedings of the 6th International Conference on Multidisciplinary Approaches (iCMA), 2019.* Sri Lanka: Sri Jayewardenepura; 2019. Available at SSRN 3497475.
- [27] Mangwani N, Kumari S, Das S. Effect of synthetic N-acylhomoserine lactones on cell–cell interactions in marine *Pseudomonas* and biofilm mediated degradation of polycyclic aromatic hydrocarbons. *Chem Eng J.* 2016;302: 172–186.
- [28] Al-Kharusi S, Abed RM, Dobretsov S. Changes in respiration activities and bacterial communities in a bioaugmented oil-polluted soil in response to the addition of acyl homoserine lactones. *Int BiodeteriorBiodegrad.* 2016;107: 165–173.
- [29] Kumari S, Mangwani N, Das S. Synergistic effect of quorum sensing genes in biofilm development and PAHs degradation by a marine bacterium. *Bioengineered.* 2016;7(3): 205–211.
- [30] Deryabin D, Galadzhieva A, Kosyan D, Duskaev G. Plant-derived inhibitors of AHL-mediated quorum sensing in bacteria: modes of action. *Int J Mol Sci.* 2019;20(22): 5588.
- [31] Schikora A, Schenk ST, Hartmann A. Beneficial effects of bacteria-plant communication based on quorum sensing molecules of the N-acyl

- homoserine lactone group. *Plant Mol Biol.* 2016;90(6): 605–612.
- [32] Zaytseva YV, Sidorov A, Marakaev O, Khmel I. Plant-microbial interactions involving quorum sensing regulation. *Microbiology.* 2019;88(5): 523–533.
- [33] Gualpa J, Lopez G, Nieves S, ... et al. *Azospirillum brasilense* Az39, a model rhizobacterium with AHL quorum-quenching capacity. *J Appl Microbiol.* 2019;126(6): 1850–1860.
- [34] Hayat S, Faraz A, Faizan M. (2017). Root exudates: composition and impact on plant-microbe interaction. In: Ahmand I, Hussain FM (eds). *Biofilms plant soil health.* Hoboken: Wiley; 2017.
- [35] Singh H, Jaiswal V, Singh S, Tiwari S, Singh B, Katiyar D. Antagonistic compounds producing plant growth promoting rhizobacteria: a tool for management of plant disease. *J Adv Microbiol.* 2017;3(4): 1–12.
- [36] Rankl S, Gunse B, Sieper T, Schmid C, Poschenrieder C, Schröder P. Microbial homoserine lactones (AHLs) are effectors of root morphological changes in barley. *Plant Sci.* 2016;253: 130–140.
- [37] Hartmann A, Rothballer M. Role of quorum sensing signals of rhizobacteria for plant growth promotion. In: Mehnaz S (ed). *Rhizotrophs: plant growth promotion to bioremediation.* New York: Springer; 2017.
- [38] Fester T, Giebler J, Wick LY, Schlosser D, Kästner M. Plant-microbe interactions as drivers of ecosystem functions relevant for the biodegradation of organic contaminants. *Curr Opin Biotechnol.* 2014;27: 168–175.
- [39] Yap CK, Peng SHT. Cleaning up of contaminated soils by using microbial remediation: a review and challenges to the weaknesses. *Am J Biomed Sci Res.* 2019;2(3). doi: [10.34297/AJBSR.2019.02.000589](https://doi.org/10.34297/AJBSR.2019.02.000589)
- [40] Mhlongo MI, Piater LA, Madala NE, Labuschagne N, Dubery IA. The chemistry of plant-microbe interactions in the rhizosphere and the potential for metabolomics to reveal signaling related to defense priming and induced systemic resistance. *Front Plant Sci.* 2018;9: 112. doi: [10.3389/fpls.2018.00112](https://doi.org/10.3389/fpls.2018.00112)