



# A REVIEW: HARNESSING SALT-TOLERANT PLANT GROWTH-PROMOTING RHIZOBACTERIA FOR SUSTAINABLE AGRICULTURE

Prachi Patel<sup>1</sup>, Neha Maisuria<sup>2</sup>

<sup>1</sup>PG Student, Department of Microbiology, Bhagwan Mahavir College of Basic and Applied Sciences

<sup>2</sup>Assistant Professor, Department of Microbiology Bhagwan Mahavir College of Basic and Applied Science

Article DOI: <https://doi.org/10.36713/epra16605>

DOI No: 10.36713/epra16605

## ABSTRACT

The salinity of the soil is becoming a major concern for the world's food security. It is essential to create environmentally friendly sustainable techniques that boost saline soil productivity. Certain microorganisms have a dependency on salt as well as tolerance to it. Microbes that are resistant to salt may withstand ionic and osmotic stress. From very alkaline, saline, and sodic soils, many genera of salt-tolerant plant growth-promoting rhizobacteria (ST-PGPR) have been discovered. Numerous of them are also known to help plants cope with a range of biotic and abiotic stressors. The possibility of using ST-PGPR to improve the productivity of plants under salt stress has been studied in the past several years, indicating that saline agro-ecosystems can be restored through its use. This review will focus on ST-PGPR and how they might boost saline agro-ecosystem production. Additionally, it sheds light on future research directions on the use of ST-PGPR for saline soil reclamation and PGPR-mediated mechanisms of salt tolerance in various crop plants.

**KEYWORDS:** saline soils, salt tolerant, PGPR mechanism, sustainable development, sodic soil-----

## 1. INTRODUCTION

6.73 Mha land of India is damaged by salt. Of this, 16 states receive 2.95 million hectares of salinity. A major threat to the sustainability of the crop production system is the potential more than twofold increase in salt-affected soils (SAS), which are currently distributed over 6.73 million hectares (Mha) in India by 2050, according to research conducted by the Central Soil Salinity Research Institute (CSSRI). Currently, there is scientific consensus that the human production model and energy consumption have an effect in climate change [1]. Recent research shown that 20% of raised land and 33% of irrigated land are vulnerable to high salinity, with a predicted rise of 10% each year. More than 1.257 million hectares of soil are affected by salt [2].

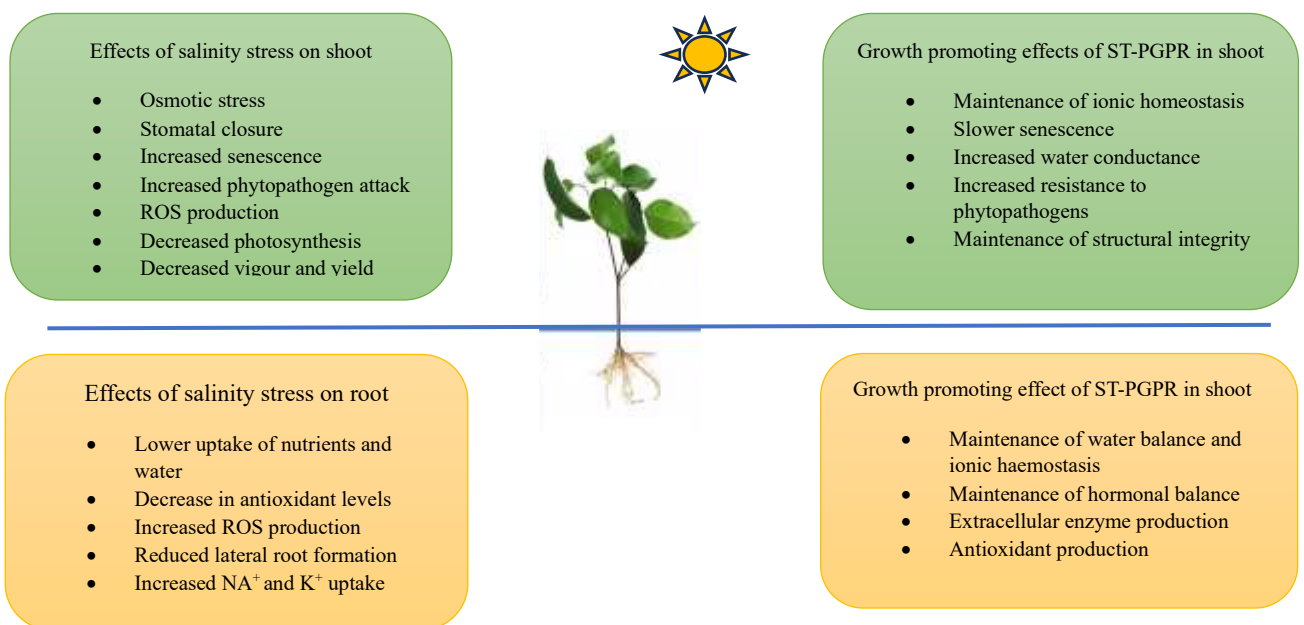
Global climate change generates adverse environmental conditions, including salinity in soils, severe temperatures, droughts, and floods, limiting plant species spread and crop production [3, 4]. Severe weather conditions negatively impact the environment, economy, and society, particularly in semi-arid regions, while arid land degradation is a consequence of the Green Revolution's excessive use of synthetic substances [5,6]. The global agricultural system's primary objective is to increase output, but negative environmental effects must be reduced through rethinking existing and future approaches and developing sustainable technology [7]. Among the methods developed by scientists are the breeding of salt-tolerant crops, the physical removal of salts from the soil's surface, and the chemical treatment of the soil [8,9]. By controlling hydric resources and their quality, sustainable land management, although gradual and expensive, can lessen the consequences of soil salinity [10,11].

Plant growth-promoting rhizobacteria (PGPRs) are a vital component of organic agriculture. Their presence in the rhizosphere is crucial for improving soil production, promoting plant development, and squelching plant diseases. The majority of promising and extensively documented genera of photosynthesis-generating bacteria (PGPR) comprise *Pseudomonas*, *Aeromonas*, *Klebsiella*, *Azoarcus*, *Enterobacter*, *Azospirillum*, *Clostridium*, *Azotobacter*, *Arthobacter*, *Rhizobium*, *Gluconacetobacter*, *Bacillus*, and *Serratia* [7].

## 2. PROBLEM OF SOIL SALINIZATION

While it is known that soil salinity is a concern on every continent in the world, a reliable map of the locations and distribution of salty soils is lacking. The majority of crops in these regions are irrigated, and to make matters worse, poor irrigation management causes secondary salinization, which threatens 20% of irrigated land globally [12]. Poor-quality groundwater is being utilized for irrigation to produce more on each hectare of arable land that

is available [13]. Almost 70% of ground water is used for agricultural globally, but when it is depleted, lower-quality water is used instead [14]. Salt-affected soil solenchak and solonetz are identified by the soil map legend and are distinguished by an accumulation of highly soluble salt. Its EC varies from 8 to 15 dS/m<sup>1</sup> and it is found in semi-arid coastal regions.[15]. Unsustainable groundwater withdrawals decrease crop quality and range, often rendering a region unusable for cultivation, and cause waterlogging, soil toxicity, and acidity [16]. salinity diminishes agricultural productivity of cereals, legumes, forages, and horticultural crops, while also affecting soil microbial ecology and physicochemical qualities. Salt stress can lead to reduced agricultural productivity, increased soil erosion, and lower economic returns [17]. Plants have physiological changes due to salinity stress, including stomatal closure, early senescence, reduced photosynthesis, and increased oxidative damage [18]. For actual use, salt-affected areas are classified as saline, alkali, sodic, saline-alkali, or saline sodic. Saline soil has an excess of soluble salt in the soil solution, which is the liquid that exists between soil particles [19]. Solonetz soils—also referred to as alkaline or sodic soils—have a pH above 8.5 and are very alkaline. They are found on 135 million hectares around the world in semi-arid temperate continental climates, including China, Argentina, Kazakhstan, Hungary, Bulgaria, Romania, and the Ukraine [15]. Plant development is hindered by salinity, which is caused by the buildup of dissolved salts in soil water, either naturally or by human activity [20].



**Figure 1. Effects of salinity stress and associated tolerance mechanisms induced by salt-tolerant plant growth-promoting rhizobacteria (ST-PGPR) in both plant roots and shoots.**

According to estimates, saline soil covers around 7 million hectares of land in India, which includes Punjab, Haryana, Uttar Pradesh, Bihar, and portions of Rajasthan [21]. Soil salinity not only hinders plant growth and development, but it also has a negative influence on microbial composition, variety, and functions [22].

### 3. IMPACT OF SALINITY ON PLANTS

Soil salinity poses a risk to plant health. Salinity alters the blooming and fruiting patterns, causing a disruption in reproductive physiology that, in turn, impacts crop yield and biomass [23]. Plant development is inhibited by salinity stress, which affects physiological and metabolic processes including water stress, ion toxicity, nutritional problems, oxidative stress, altered metabolic processes, membrane disruption, reduced cell division and expansion, and genotoxicity based on severity and duration. [24]. Salt-affected soil consists of saline soil, sodic soil, and saline-sodic soil [25]. Plant phosphorus (P) absorption is considerably reduced by soil salinity, since phosphate ions precipitate with Ca ions [26]. High salt stress (150 mm NaCl) in tomato has been demonstrated to affect flowering transition time, producing a delay in the first inflorescence as well as a reduction in shoot and root formation [27]. Deferred blooming in chickpea is directly associated with increased Na<sup>+</sup> concentrations in the laminae of fully expanded leaves [28]. Interestingly, the Salt Overly Sensitive (SOS) route is an important protective system linked with Na<sup>+</sup> ejection and maintaining particle homeostasis at the cell level [29,30]. However, salt stress has been demonstrated in multiple studies to deactivate SOS, photoperiodical, and circadian clock switch proteins associated with flowering [31,32,33]. The primary cations linked with salinity are Na<sup>+</sup>, Ca<sup>2+</sup>, and Mg<sup>2+</sup>, while the most prevalent anions are Cl<sup>-</sup>, SO<sub>4</sub><sup>2-</sup>, and HCO<sub>3</sub><sup>-</sup> [34]. Salinity stress increases signal perception

and transduction, restricting  $\text{Na}^+$  uptake and accumulation, while osmotic stress tolerance is mediated by a decrease in stomatal conductance.[35]. Accumulation of  $\text{Na}^+$  ions in plant tissues causes the generation of reactive oxygen species (ROS), including singlet oxygen ( $^1\text{O}^2$ ), hydrogen peroxide ( $\text{H}_2\text{O}_2$ ), superoxide ( $\text{O}_2^{\cdot-}$ ), and hydroxyl radical ( $\cdot\text{OH}$ ), which impede photosynthesis [36]. Compared to other ions like  $\text{Ca}^{2+}$ ,  $\text{Mg}^{2+}$ , or  $\text{K}^+$ , sodium is a highly poisonous and soluble ion that is extensively disseminated and ineffectual in causing salinization in plants. [37]. Certain elements, especially sodium, chlorine, and boron, can be hazardous to plants. Excessive sodium buildup in cell walls can swiftly promote osmotic stress and cell death [38]. Plants grow well in salty conditions, but osmotic equilibrium is essential for development because it keeps cells from drying up and dying.[39]. A high concentration of  $\text{K}^+$  is also necessary for tRNA binding to ribosomes and consequently protein synthesis. and cannot be replaced by  $\text{Na}^+$ . However, ion toxicity caused by the substitution of  $\text{K}^+$  by  $\text{Na}^+$  in metabolic processes make it unavailable for the process. [40,41]. The detrimental effects of salinity are due to stress on differentiation and the cell cycle, which results in fewer cells and limits growth. Additionally, it has a deleterious effect on all aspects of plant development, including mitosis, DNA, RNA, enzyme activity, and seed germination [42,43,44].

#### 4. MECHANISMS OF SALINITY TOLERANCE/RESISTANCE INDUCED BY PGPB

Crop production is significantly reduced by salinity's physiological impacts [1,45]. On the other hand, it has been demonstrated that applying Plant Growth Promoting Bacteria increases crop production by lessening the physiological harm brought on by high soil ion concentrations [1,46]. Elevated ion concentrations inside the cytoplasm cause an ionic imbalance, which in turn preserves the  $\text{K}^+$  osmotic equilibrium, triggers osmotic reactions, and activates genes involved in adaptive, metabolic, protective, and amino acid transport pathways. Furthermore, the synthesis of organic solutes (alanine, serine, glutamic acid, sucrose, choline, and betaine) [47,48] bolster the osmotic force inside cells to stabilize macromolecules within cells during salt stress [49]. In general, direct and indirect mechanisms make up the two subtypes of PGPB modes of action. In direct mechanism, organisms reside inside plants to have an impact on plant metabolism, indirect organisms are thought to be external [50] By functioning as hormone sinks and releasing growth factors, microbes enhance adaptability through indirect mechanics. By regulating secondary metabolites and signal susceptibility, indirect mechanisms help increase resistance to pathogen invasions and stress [51,52]. The role of halotolerant plant growth-promoting rhizobacteria (HT-PGPR) in reducing salt stress in agricultural plants has grown in the last several years.

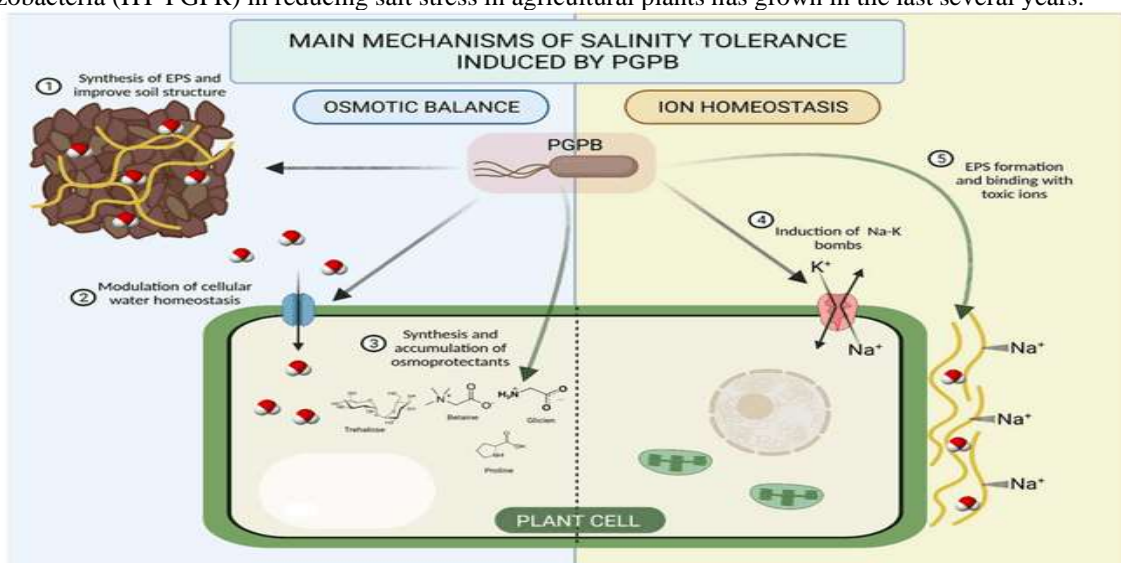


Figure 2. Main mechanisms of salinity stress tolerance induced by Plant Growth Promoting Bacteria (110).

By directly producing a variety of advantageous metabolites, including exopolysaccharides, siderophores, volatile organic compounds (VOCs), compatible osmolytes, and phytohormones, or indirectly by controlling the expression of stress-related genes and thwarting the effects of phytopathogens, ST-PGPR increase the productivity of the saline-agroecosystem (Figure 3) [53,54].

#### 4. DIRECT MECHANISMS

##### *Indole-3-Acetic Acid production*

Plant growth regulators, sometimes referred to as phytostimulators, are substances that naturally possess the capacity to control the synthesis of certain growth regulator enzymes and have a significant impact on many aspects of plant growth, including the morphological, physiological, and biochemical processes of the plant [55]. Indole acetic acid, which is necessary for cell division and elongation in plants under salt stress, is produced by



the ST-PGPR. *Azotobacter*, *Arthrobacter*, *Azospirillum*, *Pseudomonas*, *Stenotrophomonas*, and *Rahnella* are a few of those most renowned ST-PGPR generating IAA under salt stress [56,57,58]. Assessing the mode of action of various phytopathogens has shown that while excessive concentrations of IAA can be hazardous, at appropriate levels they may have positive benefits [1,59]. With its help, we may be able to identify the PGPR strain utilizing bacteria that release IAA [60]. One of the most significant plant hormones, IAA controls a wide range of processes related to plant growth and development across the course of the cell cycle, including root initiation, apical dominance, blooming, fruit ripening, and senescence, as well as cell division, elongation, and differentiation. [61] Further, by boosting the availability of water and nutrients to plants, salt-tolerant Rhizobacteria increases the production of IAA in plants, reducing the growth of tap roots, encouraging the elongation of root hairs, and lengthening lateral roots.[62]. The role of IAA in resistance to increased salt stress was demonstrated by Numan et al. (2018), who observed abundant IAA generation with osmotolerant PGPR *Azospirillum brasilense* NH at high salt concentration in durum wheat. It is well known that maize growth is enhanced by desiccation-tolerant *Micrococcus luteus* when IAA is produced [63,64]. Since the synthesis of phytohormones is a critical trait of PGPR that may be exploited, more research must be done to determine how to employ PGPR to mitigate the negative impacts of salinity [65].

### ***Symbiotic Nitrogen fixation***

For plants to develop and be productive, nitrogen (N) is the most important nutrient but developing plants cannot use it. By employing a sophisticated enzyme system called nitrogenase involving a process known as biological N<sub>2</sub> fixation (BNF), which transforms atmospheric N<sub>2</sub> into forms that plants can use [66]. Symbiotic bacteria found in legume nodules are believed to be the most crucial element in the biological fixation of atmospheric nitrogen.[67]. Numerous bacterial species have been shown to be connected to the plant's rhizosphere, which further promotes the growth of the plant. It encompasses the following, *Erwinia*, *Azospirillum*, *Flavobacterium*, *Bacillus*, *Arthrobacter*, *Rhizobium*, *Acinetobacter*, *Burkholderia*, *Pseudomonas*, *Enterobacter*, and *Serratia* [68,69]. Around the world, biological nitrogen fixation generates 180 X 10<sup>6</sup> metric tons annually, of which symbiotic relationships generate 80% and free-living or associative systems the remaining portion.[70]. In two years, a rice rhizosphere strain of *A. radiobacter* greatly boosted yields of barley and winter wheat (5–30%) and contributed 23–32% to total nitrogen absorption. [111].

### ***Non-symbiotic nitrogen-fixation***

It has been shown that when soil contains mineral nitrogen, free-living nitrogen-fixing bacteria consume the nitrogen instead of fixing it [71,72]. It is essential to both agriculture and the fixing of nitrogen. An energy-oriented fixation of nitrogen in a form that plants might easily use has been the main obstacle obscuring its effect. By putting them closer to the roots, the constraint might be readily overcome. Numerous bacteria are linked to non-symbiotic nitrogen fixation, including but not limited to *Azoarcus* sp., *Herbaspirillum* sp., and *Azotobacter* sp., [73,74]. Soil bacteria can convert mineral nitrogen into gases like NO, N<sub>2</sub>O, and N<sub>2</sub>, which harms agroecosystems. Nitrogen metabolism genes in soil bacteria are influenced by nitrogen availability, potentially aiding nitrogen accumulation.[75]. Noori et al.'s 2019 study also examined the impact of effective isolates on plant development under salt stress. The isolates included rhizobial and non-rhizobial bacteria that are tolerant of salinity and drought from surface-sterilized alfalfa root nodules cultivated in salty soils. They simultaneously injected *K. cowanii* A37, *S. meliloti* ARh29, and *Klebsiella* sp. A36 into the alfalfa plant. According to the findings, nitrogen could be supplied to plants by *Klebsiella* sp. A36 and *Kcowanii* A37 and increased plant growth indices in the absence of rhizobial bacteria [112].

### ***Phosphate solubilization***

Phosphorus is the second most essential element in plant nutrition, behind nitrogen. The pH, compaction, aeration, moisture, temperature, texture, and organic matter of soils, crop residues, the size of plant root systems, root exudate secretions, and soil bacteria that are present all affect how much phosphorus is accessible to plants [76]. As rescue mechanisms, soil microbes contribute to soil acidification and consequent solubilization of inorganic phosphates. Among the most common are phosphorus-solubilizing bacteria from the genera *Bradyrhizobium*, *Cladosporium*, *Azotobacter*, *Bacillus*, *Pseudomonas*, and *Enterobacter* [77]. Phosphoric fertilizers are widely used in agricultural areas to compensate for the soil's *Pseudomonas* deficiency. Plants absorb less phosphatic fertilizer, and what is left over is rapidly converted by the soil into insoluble complexes [78]. Not only bacteria but fungi from the following groups are also taken into account: *Achrothcium*, *Alternaria*, *Arthrobotrys*, *Aspergillus*, *Cephalosporium*, *Chaetomium*, *Cladosporium*, *Cunninghamella*, *Curvularia*, *Fusarium*, *Glomus*, *Helminthosporium*, *Micromonospora*, *Phenomiocenspora*, *Phenomycylum*, *Populospora*, *Pythium*, *Rhizoctonia*, *Rhizopus*, *Saccharomyces*, *Schizosaccharomyces*, *Schwanniomyces*, *Sclerotium*, *Torula*, *Trichoderma*, and *Yarrowia*. Of all the microorganisms that can solubilize phosphate, bacteria make up as much as 50% of the total, whilst fungi make up as little as 0.5% [79].





### ***Exo-polysaccharides (EPS) production***

One common trait shared by several rhizosphere bacteria is the production of surface polysaccharides, or EPS. Even though the quantity and makeup of EPS might fluctuate throughout ST-PGPR strains, unfavourable circumstances lead to the formation of large amounts of EPS [80,81,82]. The synthesis of exopolysaccharides (EPS) also gives bacteria a means of preventing plants from absorbing harmful ions by enclosing the root system in a physical barrier, these chemicals mitigate the consequences of the ion toxicity phase [11,83,84]. The generation of EPS promotes the growth of biofilm, which in turn promotes soil aggregation and moisture retention [85] Stress affects EPS production. In *E. coli*, LonS, an ATP-dependent enzyme that denatures proteins under stress, regulates ResA, a positive transcriptional regulator. [86]. Additionally, the production of exopolysaccharides may improve the odds of bacterial survival in desiccation or nutrient-deprived environments, as well as aid in nitrogen fixation by limiting excessive oxygen tension. These polysaccharides may also be implicated in cell aggregation [87].

### ***Siderophore production***

Approximately 140 enzymes require iron as a cofactor, necessary for cytochrome and ribonucleotide reductase. It occurs in the forms of oxyhydroxides, insoluble hydroxides, and ferric Fe<sup>3+</sup> under circumstances of plentiful O<sub>2</sub>, none of which are accessible to microorganisms or plants [88]. Low molecular weight molecules called siderophores bind iron (Fe<sup>+++</sup>) and move it across the cell membrane. The iron siderophore complex enters cells by bacterial absorption, facilitating the growth of microorganisms. The majority of the Fe<sup>+3</sup> in the rhizosphere is bound by siderophores produced by soil bacteria, avoiding fungal diseases.[89]. The capacity of rhizobacteria to use siderophores generated by various genera of rhizobacteria is often variable; some are skilled at using siderophores produced by the same species (homologous siderophores), while others may be able to use those produced by other rhizobacteria (heterologous siderophores) [90]. Winkelman and Dreschel (1997) recognised five kinds of siderophores, formerly known as sideramines and sideromycins, which are iron chelators classified into hydroxamates and techolates.[91]. Therefore, excessive soil concentrations of heavy metals cause stressors on plants, which are lessened by bacterial siderophores [88]. In 2007, Crowley and Kraemer discovered that oat plants had a siderophore-mediated iron transport system. They deduced that siderophores generated by rhizosphere bacteria supply iron to oat, which possesses mechanisms for using Fe-siderophore complexes in situations where iron availability is restricted [114].

### ***HCN production***

It is during the early stationary growth phase that hydrogen cyanide is generated [92].The volatile chemical hydrogen cyanide (HCN) is released by *Pseudomonas*. Along with their protective function for the host plant, they have antibacterial actions [94]. In solution, cyanide takes the form of free cyanide, which consists of the non-dissociated HCN and the cyanide anion (CN<sup>-</sup>). Cyanide, a phytotoxic chemical that may disrupt key metabolic enzymes, is thought to be a common characteristic of harmful rhizobacterial isolates [95]. A broad-spectrum antibacterial substance called HCN is used in the biological management of root infections [96]. Glycine is used to create this element together with CO<sub>2</sub> [97]. According to Qurashi and Sabri (2012a), chickpea growth, soil structure stability, and aggregation under salt were all enhanced by EPS-producing ST-PGPR *Halomonas variabilis* (HT1) and *P. rifietoensis* (RT4) [93].

### ***Ammonia production***

The generation of ammonia is mostly seen in leguminous rhizobacteria and is associated with nitrogen fixing. A variety of rhizobacteria that promote plant development interact with C3 and C4 plants (such as cotton, rice, wheat, maize, sugarcane, Jatropha, and so on) to greatly enhance their vegetative growth and grain output [98]. A variety of aerobic heterotrophic bacteria called Azospirillum species proliferate widely in the rhizosphere of gramineous plants, fixing nitrogen in microaerobic circumstances. <sup>15</sup>N tracer methods revealed that through biological nitrogen fixation, *Azospirillum lipoferum* and *Azospirillum brasilense* provided 7–12% of the N in wheat plants [98,99]. In greenhouse studies (Muthukumarasamy et al. 1999) are inoculation with Azospirillum also significantly enhanced the N content of sugarcane leaves, a reflection of Azospirillum's ammonia production [113].

## **8. INDIRECT MECHANISMS**

Rhizobacteria's primary indirect method of promoting plant development is via serving as biocontrol agents [100]. In general, the main biocontrol mechanisms in PGPR are nutritional competition, niche exclusion, induced systemic resistance, and the synthesis of antifungal metabolites [101]. It has been documented that a wide variety of rhizobacteria create antifungal metabolites, including phenazines, pyrrolnitrin, 2,4-diacetylphloroglucinol, pyoluteorin, viscosinamide, and tensin [102]. On the other hand, secondary metabolites and their sensitivity to signals generated by the microbe are relayed by an indirect process. It involves, for instance, the development of tolerance to various stress circumstances as well as resistance to a variety of pathogen attacks [103]. Additionally,



ethylene and jasmonate signaling occur inside the plant during ISR, and these hormones activate the host plant's defensive mechanisms against a range of plant diseases [105]. In order to improve soil richness, this method uses a variety of tools, such as the production of hydrolytic proteins, anti-microbials, and more, to reduce the need for agrochemicals (pesticides and manures). It also addresses a wide range of misuses of plant development that elevate rhizobacteria [104].

## 9. FUTURE PROSPECTS

Salt tolerance is increased by symbiotic bacteria in the rhizosphere and roots of plants; however, it is uncertain if halophyte rhizobia can be successfully introduced to non-halophyte plants over the long term.[106]. On the one hand, a more varied group of microorganisms might offer more advantages to plants [107]. However, certain PGPB strains may concurrently display a number of actions that promote plant development. For instance, it has been reported in a number of articles that bacteria that fix nitrogen are capable of phosphate solubilization among other things. On the other hand, it is unknown how actively a single bacterial strain may display many characteristics that are advantageous for plant development at the same time [108]. The research looks at metabolites and genes related to salt tolerance, investigates salt-stressed plant growth strategies utilizing -OMICs techniques, and investigates the effect of PGPR on plant epigenome alteration.[109]. In recent year, an growing amount of research studies has confirmed the clear relevance and importance of bacterial consortia by focusing on their application and having a beneficial impact on plant growth and development.

## 10. CONCLUSION

The increasing population and demand for food have led to increased agricultural production, but saline stress is a major obstacle. PGPRs provide a reliable solution by accelerating plant development, seed germination, and protection against environmental stressors. Salt-tolerant rhizobacteria can reduce salt stress while plants grow. Further research is needed to confirm the effectiveness of PGPR formulations on different plant species and genotypes. PGPRs can improve agricultural productivity by enhancing soil fertility and plant nutrition through nutrient acquisition.

## REFERENCES

1. Ilangumaran, G.; Smith, D.L. *Plant growth promoting rhizobacteria in amelioration of salinity stress: A systems biology perspective*. *Front. Plant Sci.* 2017, 8, 1768.
2. Abdelraheem, A.; Esmaili, N.; O'Connell, M.; Zhang, J. *Progress and perspective on drought and salt stress tolerance in cotton*. *Ind. Crops Prod.* **2019**, 130, 118–129.
3. Mukhopadhyay, R.; Sarkar, B.; Jat, H.S.; Sharma, P.C.; Bolan, N.S. *Soil salinity under climate change: Challenges for sustainable agriculture and food security*. *J. Environ. Manag.* 2020, 280, 111736.
4. Corwin, D.L. *Climate change impacts on soil salinity in agricultural areas*. *Eur. J. Soil Sci.* 2021, 72, 842–862.
5. Pingali, P.L. *Green revolution: Impacts, limits, and the path ahead*. *Proc. Natl. Acad. Sci. USA* **2012**, 109, 12302–12308.
6. Yang, X.; Fang, S. *Practices, perceptions, and implications of fertilizer use in East-Central China*. *Ambio* **2015**, 44, 647–652.
7. Etesami, H.; Maheshwari, D.K. *Use of plant growth promoting rhizobacteria (PGPRs) with multiple plant growth promoting traits in stress agriculture: Action mechanisms and prospects*. *Ecotoxicol. Environ. Saf.* **2018**, 156, 225–246.
8. Safdarian, M., H. Askari, V. Shariati and G. Nematzadeh (2019). *Transcriptional responses of wheat roots inoculated with *Arthrobacter nitroguajacolicus* to salt stress*. *Scientific reports*, 9
9. Safeena, M. and M. Zakeel (2019). *Strategies for the Management of Soil-Borne Pathogens and Crop Production Under Saline Environment, Plant Health Under Biotic Stress*. Springer, 213-229.
10. Shrivastava, P.; Kumar, R. *Soil salinity: A serious environmental issue and plant growth promoting bacteria as one of the tools for its alleviation*. *Saudi J. Biol. Sci.* 2015, 22, 123–131.
11. Etesami, H.; Adl, S.M. *Can interaction between silicon and non-rhizobial bacteria benefit in improving nodulation and nitrogen fixation in salinity-stressed legumes? A review*. *Rhizosphere* 2020, 15, 100229.
12. Glick, B.R., 2007. *Promotion of plant growth by bacterial ACC deaminase*. *Crit. Rev. Plant Sci.* 26, 227–242.
13. Majeed, A. and Z. Muhammad (2019). *Salinity: A Major Agricultural Problem – Causes, Impacts on Crop Productivity and Management Strategies, Plant Abiotic Stress Tolerance*. Springer, 83-99.
14. FAO and ITPS (2015). *Status of the World's Soil Resources (SWSR) – Main Report*. Rome: Food and Agriculture Organization of the United Nations and Intergovernmental Technical Panel on Soils.
15. IUSS Working Group WRB (2015). *World Reference Base for Soil Resources 2014, update 2015 International Soil Classification System for Naming Soils and Creating Legends for Soil Maps*. *World Soil Resources Reports No. 106*, Rome: FAO.
16. Abod, É., É. Laslo, S. Szentes, S. Lányi and G. Mara (2019). *Plant Growth-Promoting Bacteria: Strategies to Improve Wheat Growth and Development Under Sustainable Agriculture, Plant Growth Promoting Rhizobacteria for Agricultural Sustainability*. Springer, 1-17.
17. Hu Y, Schmidhalter U. 2002. *Limitation of salt stress to plant growth*. In Hock B, Elstner C F (eds.) *Plant Toxicology*. Marcel Dekker Inc., New York. pp. 91–224.



18. Mahawar, L., and Shekhawat, G. S. (2019). EsHO 1 mediated mitigation of NaCl induced oxidative stress and correlation between ROS, antioxidants and H<sub>2</sub>O<sub>2</sub> in seedlings of *Eruca sativa*: underutilized oil yielding crop of arid region. *Physiol. Mol. Biol. Plants* 25, 895–904. doi: 10.1007/s12298-019-00663-7.
19. Etesami, H. (2018). Using Halotolerant Bacteria for Farming in Salt Affected Soils. *Acta Scientific Agriculture* **Volume 2**.
20. Powrie, W. (2018). *Soil mechanics: concepts and applications*. CRC Press.
21. Patel, B.B., Patel, Bharat.B., Dave, R.S., 2011. Studies on infiltration of saline-alkali soils of several parts of Mehsana and Patan districts of north Gujarat. *J. Appl. Technol. Environ. Sanitation* 1 (1), 87–92.
22. Shrivastava P, Kumar R. 2015. Soil salinity: A serious environmental issue and plant growth promoting bacteria as one of the tools for its alleviation. *Saudi J Biol Sci.* **22**: 123–131.
23. Promila, K., and Kumar, S. (1982). Effect of salinity on flowering and yield characters in pigeonpea. *Ind. J. Plant Physiol.* 25, 252–257.
24. Carillo, P.; Annunziata, M.G.; Pontecorvo, G.; Fuggi, A.; Woodrow, P. Salinity stress and salt tolerance. In *Abiotic Stress in Plants – Mechanisms and Adaptations*; Shanker, A., Venkateswarlu, B., Eds.; IntechOpen: London, UK, 2011.
25. Mukhopadhyay, R.; Sarkar, B.; Jat, H.S.; Sharma, P.C.; Bolan, N.S. Soil salinity under climate change: Challenges for sustainable agriculture and food security. *J. Environ. Manag.* **2020**, 280, 111736.
26. Bacilio, M., Rodriguez, H., Moreno, M., Hernandez, Juan-Pablo, Bashan, Y., 2004. Mitigation of salt stress in wheat seedlings by a *gfp*-tagged *Azospirillum lipoferum*. *Biol. Fertility Soils* 40, 188–193.
27. Ghanem, M. E., van Elteren, J., Albacete, A., Quinet, M., Martínez-Andújar, C., Kinet, J. M., et al. (2009). Impact of salinity on early reproductive physiology of tomato (*Solanum lycopersicum*) in relation to a heterogeneous distribution of toxic ions in flower organs. *Funct. Plant. Biol.* 36, 125–136.
28. Panwar, M., Tewari, R., & Nayyar, H. (2016). Native halo-tolerant plant growth Promoting rhizobacteria *Enterococcus* and *Pantoea* sp. Improve seed yield of Mungbean (*Vigna radiata* L.) under soil salinity by reducing sodium uptake and Stress injury. *Physiology and Molecular Biology of Plants*, 22(4), 445–459.
29. Zhu J. K., Liu J. P., Xiong L. M. (1998). Genetic analysis of salt tolerance in *Arabidopsis*: Evidence for a critical role of potassium nutrition. *Plant Cell* 10 (7), 1181–1191. 10.2307/3870720.
30. Zhu J. K. (2003). Regulation of ion homeostasis under salt stress. *Curr. Opin. Plant Biol.* 6 (5), 441–445. 10.1016/S1369-5266(03)00085-2.
31. Kim, J., Geng, R., Gallenstein, R. A., and Somers, D. E. (2013). The F-box protein ZEITLUPE controls stability and nucleocytoplasmic partitioning of GIGANTEA. *Development* 140, 4060–4069. doi: 10.1242/dev.096651.
32. Park, H. J., Kim, W. Y., and Yun, D. J. (2013). A role for GIGANTEA: keeping the balance between flowering and salinity stress tolerance. *Plant Signal. Behav.* 8:e24820. doi: 10.4161/psb.24820.
33. Ryu, J. Y., Lee, H.-J., Seo, P. J., Jung, J.-H., Ahn, J. H., and Park, C.-M. (2014). The *Arabidopsis* floral repressor BFT delays flowering by competing with FT for FD binding under high salinity. *Mol. Plant* 7, 377–387. doi: 10.1093/mp/ss114.
34. Safdar, H.; Amin, A.; Shafiq, Y.; Ali, A.; Yasin, R.; Shoukat, A.; Hussan, M.U.; Sarwar, M.I. A review: Impact of salinity on plant growth. *Nat. Sci.* **2019**, 17, 34–40.
35. Blaylock, A.D., 1994. Soil salinity, salt tolerance and growth potential of horticultural B3 landscape plants. Co-operative Extension Service, University of Wyoming, Department of Plant, Soil and Insect Sciences, College of Agriculture, Laramie, Wyoming.
36. Ahmad M, Zahir Z A, Asghar H N, Asghar M. 2011. Inducing salt tolerance in mung bean through coinoculation with rhizobia and plant growth-promoting rhizobacteria containing 1-aminocyclopropane-1-carboxylate deaminase. *Can J Microbiol.* **57**: 578–589.
37. Maathuis, F.J.M. Sodium in plants: Perception, signalling, and regulation of sodium fluxes. *J. Exp. Bot.* **2014**, 65, 849–858.
38. Munns, R., 2002. Comparative physiology of salt and water stress. *Plant Cell Environ.* 25, 239–250.
39. Ashraf, M., 2004. Some important physiological selection criteria for salt tolerance in plants. *Flora* 199, 361–376.
40. Zhu, J.K., 2002. Salt and drought stress signal transduction in plants. *Annu. Rev. Plant Biol.* 53, 247–273.
41. Chinnusamy, V., Zhu, J., Zhu, Jian-Kang, 2006. Gene regulation during cold acclimation in plants. *Physiol. Plant.* 126 (1), 52–61.
42. Seckin, B., Sekmen, A.H., Turkan, I., 2009. An enhancing effect of exogenous mannitol on the antioxidant enzyme activities in roots of wheat under salt stress. *J. Plant Growth Regul.* 28, 12–20.
43. Tabur, S., Demir, K., 2010. Role of some growth regulators on cytogenetic activity of barley under salt stress. *Plant Growth Regul.* 60, 99–104.
44. Javid, M.G., Sorooshzadeh, A., Moradi, F., Sanavy Seyed, A.M.M., Allahdadi, I., 2011. The role of phytohormones in alleviating salt stress in crop plants. *AJCS* 5 (6), 726–734.
45. Munns, R.; Tester, M. Mechanisms of Salinity Tolerance. *Annu. Rev. Plant Biol.* **2008**, 59, 651–681.
46. Dodd, I.C.; Pérez-Alfocea, F. Microbial amelioration of crop salinity stress. *J. Exp. Bot.* **2012**, 63, 3415–3428.
47. Shabala S, Pottosin I. 2014. Regulation of potassium transport in plants under hostile conditions: Implications for abiotic and biotic stress tolerance. *Physiol Plant.* **151**: 257–279.
48. Miller K J, Wood J M. 1996. Osmoadaptation by rhizosphere bacteria. *Ann Rev Microbiol.* **50**: 101–136.
49. Lippert K, Galinski E A. 1992. Enzyme stabilization by ectoine-type compatible solutes: Protection against heating, freezing and drying. *Appl Microbiol Biotechnol.* **37**: 61–65.
50. Vessey JK. Plant growth promoting rhizobacteria as biofertilizers. *Plant Soil.* 2003; 255: 571–586.
51. Ramos-Solano B, Barriuso J, Gutiérrez-Mañero FJ. Physiological and molecular mechanisms of plant growth promoting rhizobacteria (PGPR). In: Ahmad I, Pichtel J, Hayat S, eds. *Plant-bacteria interactions: strategies and techniques to promote*





- plant growth, 2008: 41-54 growth promoting rhizobacteria (PGPR). In: Ahmad I, Pichtel J, Hayat S, eds. *Plant-bacteria interactions: strategies and techniques to promote plant growth*, 2008: 41-54.
52. Bhat, M. A., Kumar, V., Bhat, M. A., Wani, I. A., Dar, F. L., Farooq, I., et al. (2020). Mechanistic insights of the interaction of plant growth-promoting rhizobacteria (PGPR) with plant roots toward enhancing plant productivity by alleviating salinity stress. *Front. Microbiol.* 11, 1952. doi: 10.3389/fmicb.2020.01952.
53. Prasad, M., Ramakrishnan, S., Chaudhary, M., Choudhary, M., and Jat, L. K. (2019). "Plant growth promoting rhizobacteria (PGPR) for sustainable agriculture: perspectives and challenges" in *PGPR Amelioration in Sustainable Agriculture*, eds A. K. Singh, A. Kumar and P. K. Singh (Woodhead Publishing), 129–157. doi: 10.1016/B978-0-12-815879-1.00007-0.
54. Lugtenberg BJ, Chin A-Woeng TF, Bloemberg GV. *Microbe-plant interactions: principles and mechanisms*. *Antonie Van Leeuwenhoek*. 2002; 81: 373-383.
55. Somers E, Vanderleyden J, Srinivasan M. Rhizosphere bacterial signalling: a love parade beneath our feet. *Crit Rev Microbiol*. 2004; 30: 205-240.
56. Egamberdieva, D., Kamilova, F., Validov, S., Gafurova, L., Kucharova, Z., and Lugtenberg, B. (2008). High incidence of plant growth-stimulating bacteria associated with the rhizosphere of wheat grown on salinated soil in Uzbekistan. *Environ. Microbiol.* 10, 1–9. doi: 10.1111/j.1462-2920.2007.01424.x.
57. Piccoli, P., Travaglia, C., Cohen, A., Sosa, L., Cornejo, P., Masuelli, R., et al. (2011). An endophytic bacterium isolated from roots of the halophyte *Prosopis strombulifera* produces ABA, IAA, gibberellins A 1 and A 3 and jasmonic acid in chemically-defined culture medium. *Plant Growth Regul.* 64, 207–210. doi: 10.1007/s10725-010-9536-z.
58. Abd\_Allah, E. F., Alqarawi, A. A., Hashem, A., Radhakrishnan, R., Al-Huqail, A. A., Al-Otibi, F. A., et al. (2017). Endophytic bacterium *Bacillus subtilis* (BERA71) improves salt tolerance in chickpea plants by regulating the plant defense mechanisms. *J. Plant Interact.* 3, 37–44. doi: 10.1080/17429145.2017.1414321.
59. Korasick, D.A.; Enders, T.A.; Strader, L.C. Auxin biosynthesis and storage forms. *J. Exp. Bot.* **2013**, 64, 2541–2555.
60. Khalid A, Arshad M, Zahir ZA. Screening plant growth-promoting rhizobacteria for improving growth and yield of wheat. *J Appl Microbiol.* 2004; 96(Suppl 3): 473-480.
61. Arshad M, Frankenberger WT. Microbial production of plant hormones. *Plant Soil.* 1991; 133(Suppl1): 1-8.
62. Nawaz, A., Shahbaz, M., Asadullah., Imran, A., Marghoob, M. U., Intiaz, M., et al. (2020). Potential of salt tolerant PGPR in growth and yield augmentation of wheat (*Triticum aestivum* L.) under saline conditions. *Front. Microbiol.* 11, 2019. doi: 10.3389/fmicb.2020.02019.
63. Grover, M., Bodhankar, S., Sharma, A., Sharma, P., Singh, J., and Nain, L. (2021). PGPR mediated alterations in root traits: way toward sustainable crop production. *Front. Sustain. Food Syst.* 4, 618230. doi: 10.3389/fsufs.2020.618230.
64. Numan M, Bashir S, Khan Y, Mumtaz R, Shinwari Z K, Khan A L, Khan A, Al-Harrasi A. 2018. Plant growth promoting bacteria as an alternative strategy for salt tolerance in plants. *Microbiol Res.* **209**: 21–32.
65. Forni C, Duca D, Glick B R. 2017. Mechanisms of plant response to salt and drought stress and their alteration by rhizobacteria. *Plant Soil.* **410**: 335–356.
66. Kim, J., Rees, D.C., 1994. Nitrogenase and biological nitrogen fixation. *Biochemistry* 33, 389–397.
67. Roughley, R.J.; Gault, R.R.; Gemell, L.G.; Andrews, J.A.; Brockwell, J.; Dunn, B.W.; Griffiths, G.W.; Hartley, E.J.; Hebb, D.M.; Peoples, M.B.; et al. Autecology of *Bradyrhizobium japonicum* in Soybean-Rice Rotations. *Plant Soil* 1995, 176, 7–14.
68. Tilak KVBR, Ranganayaki N, Pal KK, De R, Saxena AK, Nautiyal CS, Mittal S, et al. Diversity of plant growth and soil health supporting bacteria. *Curr Sci.* 2005; 89: 136-150.
69. Egamberdiyeva D. Plant-growth-promoting rhizobacteria isolated from a Calcisol in a semi-arid region of Uzbekistan: biochemical characterization and effectiveness. *J Plant Nutr Soil Sci.* 168; 2005:94-99.
70. Raymond, J., Siefert, J.L., Staples, C.R., Blankenship, R.E., 2004. The natural history of nitrogen fixation. *Mol. Biol. Evol.* 21, 541–554.
71. Ludden, P.W. Reversible ADP-Ribosylation as a Mechanism of Enzyme Regulation in Prokaryotes. *Mol. Cell. Biochem.* **1994**, 138,123–129.
72. Barron, A.R.; Wurzbarger, N.; Bellenger, J.P.; Wright, S.J.; Kraepiel, A.M.L.; Hedin, L.O. Molybdenum Limitation of Asymbiotic Nitrogen Fixation in Tropical Forest Soils. *Nat. Geosci.* **2009**, 2, 42–45.
73. Vessey JK. Plant growth promoting rhizobacteria as biofertilizers. *Plant Soil.* 2003; 255(Suppl 2): 571-586.
74. Barriuso J, Solano BR. Ecology, genetic diversity and screening strategies of plant growth promoting rhizobacteria (PGPR). *J Plant Nutr.* 2008: 1-17.
75. Robson, R.L.; Postgate, J.R. Oxygen and Hydrogen in Biological Nitrogen Fixation. *Annu. Rev. Microbiol.* **1980**, 34, 183–207.
76. Gopalakrishnan, S., Sathya, A., Vijayabharathi, R., Varshney, R.K., Laxmipathi Gowda, C.L., Krishnamurthy, L. Plant growth promoting rhizobia: challenges and opportunities. *Biotech.*, 2015; 5: 355–377.
77. Chabot R, Beauchamp C J, Kloepper J W, Antoun H. 1998. Effect of phosphorus on root colonization and growth promotion of maize by bioluminescent mutants of phosphate-solubilizing *Rhizobium leguminosarum* biovar *phaseoli*. *Soil Biol Biochem.* **30**: 1615–1618.
78. McKenzie, R.H., Roberts, T.L., 1990. Soil and fertilizers phosphorus update. In: *Proceedings of Alberta Soil Science Workshop Proceedings*, Feb. 20–22, Edmonton, Alberta, pp. 84–104.
79. Kalayu, G. Phosphate Solubilizing Microorganisms: Promising Approach as Biofertilizers. *Int. J. Agron.* **2019**, 2019, 4917256.
80. Bomfeti C. A., Florentino A. L., Guimarães A. P., Cardoso P. (2011). Exopolysaccharides produced by the symbiotic nitrogen-fixing bacteria of leguminosae. *Rev Bras Ciênc Solo* 35 657–671. 10.1590/s0100-06832011000300001.





81. Tewari S., Arora N. K. (2014a). Multifunctional exopolysaccharides from *Pseudomonas aeruginosa* PF23 involved in plant growth stimulation, biocontrol and stress amelioration in sunflower under saline conditions. *Curr. Microbiol.* 69 484–494. 10.1007/s00284-014-0612-x.
82. Khan N., Bano A. (2019). Exopolysaccharide producing rhizobacteria and their impact on growth and drought tolerance of wheat grown under rainfed conditions. *PLoS One* 14:e0222302. 10.1371/journal.pone.0222302.
83. Kim, Y.; Khan, A.L.; Waqas, M.; Lee, I.; Lee, I. Silicon Regulates Antioxidant Activities of Crop Plants under Abiotic-Induced Oxidative Stress: A Review. *Front. Plant Sci.* **2017**, 8, 510.
84. Vaishnav, A.; Varma, A.; Tuteja, N.; Choudhary, D.K. PGPR-mediated amelioration of crops under salt stress. In *Plant-Microbe Interaction: An Approach to Sustainable Agriculture*; Choudhary, D.K., Varma, A., Tuteja, N., Eds.; Springer: Singapore, 2016; pp. 205–226, ISBN 978-981-10-2854-0.
85. Isfahani, F.M., A. Tahmourespour, M. Hoodaji, M. Ataabadi and A. Mohammadi (2019). Influence of Exopolysaccharide-Producing Bacteria and SiO<sub>2</sub> Nanoparticles on Proline Content and Antioxidant Enzyme Activities of Tomato Seedlings (*Solanum lycopersicum* L.) under Salinity Stress. *Polish Journal of Environmental Studies*, **28**.
86. Stewart, B.F., et al. (1997) Clinical Factors Associated with Calcific Aortic Valve Disease. *Cardiovascular Health Study. Journal of the American College of Cardiology*, 29, 630-634.
87. Gilck BR, Patten CL, Holguin G, and Penrose DM (1999). Biochemical and genetic mechanisms used by plant growth promoting bacteria. *Imperial college press, London*, pp.187-189.
88. Rajkumar M, Ae N, PrasadMN V, Freitas H. 2010. Potential of siderophoreproducing bacteria for improving heavy metal phytoextraction. *Trends Biotechnol.* **28**: 142–149.
89. Kloepper JW, Leong J, Teintze M, Schroth MN (1980). Enhanced plant growth by plant growth promoting rhizobacteria. *Nature* 286:885–886.
90. Khan, M.S., Zaidi, A., Wani, P.A., Oves, M., 2009. Role of plant growth promoting rhizobacteria in the remediation of metal contaminated soils. *Environ. Chem. Lett.* 7, 1–19.
91. Winkelman, G, Dreschel H (1997). *Microbial Siderophores*. In: Rehm HJ, Reed G (Eds.) *Biotechnology*, VCH Publishers, Weinheim, pp. 199-245
92. Vining LC (1990). Functions of secondary metabolites. *Ann Rev Microbiol* 44: 395-427.
93. Qurashi, A. W., and Sabri, A. N. (2012b). Biofilm formation in moderately
94. halophilic bacteria is influenced by varying salinity levels. *J. Basic Microbiol.* 52, 566–572. doi: 10.1002/jobm.201100253.
95. Voisard, C.; Keel, C.; Haas, D.; Defago, G. Cyanide production by *Pseudomonas fluorescens* helps suppress black root rot of tobacco under gnotobiotic conditions. *The EMBO Journal* 8 (1989) 351-358.
96. Bakker AWW, Schippers B (1987). Microbial cyanide production in the rhizosphere in relation to potato yield reduction and *Pseudomonas* spp.-mediated plant growth-stimulation. *Soil Biol Biochem* 19: 451– 457.
97. Ramette, A.; Moënne-Loccoz, Y.; Défago, G. Prevalence of fluorescent pseudomonads producing antifungal phloroglucinols and/or hydrogen cyanide in soils naturally suppressive or conducive to tobacco black root rot. *FEMS microbiology ecology* 44 (1) (2003) 35-43.
98. Mackie, A.E.; Wheatley, R.E. Effects and incidence of volatile organic compound interactions between soil bacterial and fungal isolates. *Soil Biology and Biochemistry* 31 (3) (1999) 375-385.
99. Kennedy IR, Choudhury AIMA, KecSkcs ML (2004) Non-Symbiotic bacterial diazotrophs in crop-farming systems: can their potential for plant growth promotion be better exploited? *Soil Boil Biochem* 36(8):1229-1244.
100. Malik KA, Mirza MS, Hassan U, Mehnaz S, Rasul G, Haurat J, Bauy R, Normanel P (2002). The role of plant associated beneficial bacteria in rice-wheat Cropping System. In: Kennedy IR, Chaudhry ATMA (Eds.) *Biofertilisers in action*. Rural Industries Research and Development Corporation, Canberra, pp. 73–83
101. Glick, B.R., 2012. *Plant Growth-Promoting Bacteria: Mechanisms and Applications*. Hindawi Publishing Corporation, Scientifica.
102. Lugtenberg, B., Kamilova, F., 2009. Plant-growth-promoting rhizobacteria. *Annu. Rev. Microbiol.* 63, 541–556.
103. Bhattacharyya, P.N., Jha, D.K., 2012. Plant growth-promoting rhizobacteria (PGPR): emergence in agriculture. *World J. Microbiol. Biotechnol.* 28, 1327– 1350
104. Aeron A, Kumar S, Pandey P, Maheshwari DK. Emerging role of plant growth promoting rhizobacteria in agrobiolgy. In: Maheshwari DK, ed. *Bacteria in agrobiolgy: crop ecosystems*. 2011: 1-36.
105. Tariq, M., Ali, Q., Khan, A., Khan, G. A., Rashid, B., Rahi, M. S., ... & Husnain, T. (2014). Yield potential study of *Capsicum annuum* L. under the application of PGPR. *Advancements in Life Sciences*, 1(4), 202-207.
106. Glick, B.R., 2012. *Plant Growth-Promoting Bacteria: Mechanisms and Applications*. Hindawi Publishing Corporation, Scientifica.
107. Kearl, J.; McNary, C.; Lowman, J.S.; Mei, C.; Aanderud, Z.T.; Smith, S.T.; West, J.; Colton, E.; Hamson, M.; Nielsen, B.L. Salt-Tolerant Halophyte Rhizosphere Bacteria Stimulate Growth of Alfalfa in Salty Soil. *Front. Microbiol.* **2019**, 10, 1849.
108. Hu, J.; Wei, Z.; Friman, V.-P.; Gu, S.; Wang, X.; Eisenhauer, N.; Yang, T.; Ma, J.; Shen, Q.; Xu, Y.; et al. Probiotic Diversity Enhances Rhizosphere Microbiome Function and Plant Disease Suppression. *mBio* **2016**, 7, e01790-16.
109. Kadmiri, I.M.; Chaouqui, L.; Azaroual, S.E.; Sijilmassi, B.; Yaakoubi, K.; Wahby, I. Phosphate-Solubilizing and Auxin-Producing Rhizobacteria Promote Plant Growth under Saline Conditions. *Arab. J. Sci. Eng.* **2018**, 43, 3403–3415.
110. De Palma, M.; Salzano, M.; Villano, C.; Aversano, R.; Lorito, M.; Ruocco, M.; Docimo, T.; Piccinelli, A.L.; D’Agostino, N.; Tucci, M. Transcriptome reprogramming, epigenetic modifications and alternative splicing orchestrate the tomato root response to the beneficial fungus *Trichoderma harzianum*. *Hortic. Res.* **2019**, 6, 5.



111. Kumar, V.; Raghuvanshi, N.; Pandey, A.K.; Kumar, A.; Thoday-Kennedy, E.; Kant, S. Role of Halotolerant Plant Growth-Promoting Rhizobacteria in Mitigating Salinity Stress: Recent Advances and Possibilities. *Agriculture* 2023, 13, 168.
112. Bairamov LE, Vinogradova LV, Zavalin AA (2001). Nitrogen nutrition and productivity of barley as conditioned by the application of associative diazotrophs. *Asp Appl Biol* 63:135–139
113. Noori, F., H. Etesami, H.N. Zarini, N.A. Khoshkholgh-Sima, G.H. Salekdeh and F. Alishahi (2018). Mining alfalfa (*Medicago sativa* L.) nodules for salinity tolerant nonrhizobial bacteria to improve growth of alfalfa under salinity stress. *Ecotoxicology and environmental safety*, **162**: 129-138.
114. Mathiyazhagan S, Kavitha K, Nakkeerans S, Chandrasekar MK, Renukadevi P, Krishnamoorthy AS, Fernando WGD (2004). PGPR mediated management of stem blight of *Phyllanthus amarus* (Schum and Thonn) caused by *Corynespora cassiicola* (Berk and Curt) wei. *Arch Phytopathol Plant Prot* 37:183–199.
115. Crowley, D.E., Kraemer, S.M., 2007. Function of siderophores in the plant rhizosphere. In: Pinton, R. et al. (Eds.), *The Rhizosphere, Biochemistry and Organic Substances at the Soil-Plant Interface*. CRC Press, pp. 73–109.