



Potential Role of Microbes in Insect, Human, Plant and Environment for Present and Future Perspective

Vimalanathan Arun Prasanna^{1*}, Mani Kannan², Sankarappan Anbalagan³ and Yongqi Shao⁴

¹Department of Environmental Biotechnology, Bharathidasan University, Tamil Nadu, India

²Department of Entomology, Chemistry and Nematology, Institute of Plant protection, Agricultural Research Organization, Volcani Research Centre, Israel

³Department of Zoology, Government Arts College, Tamil Nadu, India

⁴Institute of Sericulture and Apiculture, College of Animal Sciences, Zhejiang University, China

*Corresponding Author: Vimalanathan Arun Prasanna, Department of Environmental Biotechnology, Bharathidasan University, Tamil Nadu, India.

Received: February 22, 2020

Published: July 21, 2020

© All rights are reserved by Vimalanathan Arun Prasanna, et al.

Abstract

After a very long adaptation and evolution, now microbes exist anywhere in the world. They are co-evolved with insect, plant and human/animal in different conditions which has possible benefits and least microbes become pathogen when the host had decline in the level of symbionts. Recently many research dealt with identification of microbes and production of useful compounds, enzymes, peptides and even antibiotics under selective pressure. Microbes are used as a disease marker in the recent era. In this review, we highlights the significance of microbes and their uncanny ability assisting with different host and environments.

Keywords: Biotechnology, Insect Symbiosis, Plant Microbe Interaction, Cancer, Dysbiosis, Enzymes, Gut Microbes.

Introduction

Microbes are the first form of living organism and omni present on earth originate before 3 to 4 billion years. Microbial ecosystem (host) are in different environmental niche and the microbes possess potential characteristics for various process such as degrading pollutant [1] or modifying complex substance to simple compounds, cycling the ratio of carbon and nitrogen in soil for plant growth and plant defence agent. The microbial enzymes are used in different industries like textiles, paper industry, pharmaceutical industries, food industries etc. The enzyme market globally in 2020 going to reach \$7.2 billion and a CAGR of 5.8% expected during 2016 - 2023. The enzyme market globally reach a need of 6 to 7% every year [2,3]. Screening of potential microbes from different environment and utilizing it for human society in current demand

is more important. The microbes utilized from different environment for the beneficial of human society are discussed in figure 1.

Insect gut microbes

Microbes play a broad symbiotic interaction in insects from beneficial to parasitic association. The main moto of symbiosis is mostly based on sharing nutrition or defence mechanism. The insect gut bacteria in tsetse-fly was studied in the earlier times of 20th century since it carries the vector for trypanosomal diseases. The trypanosome interaction with gut microbes are much interested to study in earlier times by Wigglesworth., et al. [4]. The understanding of host-microbe interaction mechanism become more important to develop strategies to inhibit transmittable diseases. The insect gut is big reservoir for microbes and their symbiotic interaction contribute to the growth and development of their host insect [5,6].

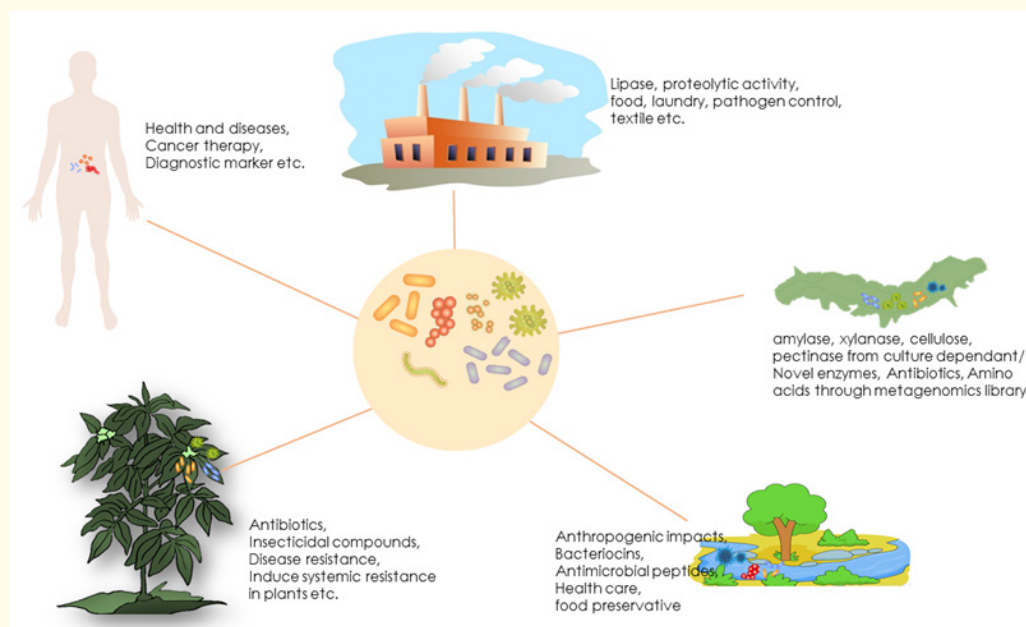


Figure 1: Microbes from environment have once adapted to different lifeforms, later become beneficial bacteria and promote growth and metabolites to their symbiotic partner.

The gut bacteria community of termites are well documented in providing glucanase, mannanase, and cellulase which the insect relies on to degrade lignocellulose based plant biomass which are used in various industrial applications [7,8]. Many novel enzymes like amylase, xylanase, cellulase, pectinase from silkworm gut bacteria and their application in food, poultry and paper industries were studied [9-12]. Esterase are used in degrading natural materials and industrial pollutions and perfume productions. Similarly tannase from the bacteria *Enterobacter cloacae* strain 41 of *Gryllotalpa krishnani* gut could be used in tannery effluent degradation, food, and pharmaceutical industrial applications [13]. Apart from this, quorum sensing compounds which are important in microbial intracellular communication in pathogenic bacteria have become a significant area in microbial research are also identified in different insects. Study in cabbage white butterfly larval (*Pieris rapae*) midgut was done to understand the quorum sensing of expressed signals in a multispecies microbial community [14]. Apart from this, various microbes are used as probiotics (lactic acid bacteria; LAB) to improve the human health through dairy, food and beverage industries which earns billions of dollars every year [15]. A

broad research must be carried out in different insect and animal gut to find more effective microbes for the benefit of human health and industry [16-18].

Revealed the world of unculturable microbes through metagenomic study

Recently, the screening of unculturable gut bacteria from several insect orders to obtain novel enzymes, antibiotics, amino acids etc. by preparing metagenomic libraries and accurate identification using advanced sequencing technologies such as genome sequencing, Next generation sequencing, Illumina HiSeq were explored. The role of insect gut microbes and their different roles on different insect species such as *Rhynchophorus ferrugineus* [19], *Acrididae* sp., *Cerambycidae* spp., *Bombyx mori* [20], *Saperda vestita*, *Dendroctonus frontalis* [21], *Hyalomma anatolicum* [22], *Blattella germanica* [23], *Spodoptera littoralis* [24], *Plutella xylostella* [25] were well studied. The Illumina MiSeq sequencing of silkworm gut bacteria revealed the presence of Proteobacteria, Firmicutes, Actinobacteria, Bacteroidetes and a unique fungus which belong to the phyla Ascomycota and Basidiomycota. The well-studied microbiota of

silkworm with previous findings of its physiology and genetics are more useful to manipulate the host microbes to improve its disease resistance, metabolism and improve economy of silk [26,27].

Impact of human gut microbes as a marker

Some of the gut microbes from human gut were studied deeply in recent times to find the correlation, their impact on health and diseases. The increased evidence on human gut bacteria involvement on cancer through dysbiosis in the human gut and these specific microbial communities are identified as a diagnostic marker in certain types of cancer [28,29] which are remarkable findings discovered by scientist to date. In this context, they reported that the loss of beneficial microbes leads to gut dysbiosis which shows a strong connection between gut microbes and health of human and animals. Dysbiosis in the healthy human affect the health and associated with diabetes, colorectal cancer (CRC), obesity. CRC case study have *Fusobacterium* association and it can be detected as fecal marker [30]. More research are need to be studied in order to understand the bacteria and human relationship to gain information related to microbe as advance marker in detecting diseases.

Significance of plant endophytic microbes

Some insect associated symbiotic microbes transmitting diseases to plants [31]. In natural, the plant endophytic microbes suppressing the pathogenic microbes through unknown mechanism because, this endophytic bacteria are evolutionarily different from soil bacteria and their precise metabolic process help in plant growth and immunity. Plant growth promoting rhizobacteria are more specific group of microbes helps in plant growth and metabolism. *B. amyloliquefaciens* isolated from *Musa* sp. and *B. subtilis* from *Gossypium* sp. are analysed as plant associated and non-plant associated strain from same species, the evolutionary trait shows the difference in strain even though they are from same species. This shows the bacterial adaptation towards plants as a unique character [32]. The antagonistic role expressed from unique bacteria against fungus are studied in recent times to find a better eco-friendly tactic instead of synthetic fungicides. *Bacillus amyloliquefaciens* an endophytic bacteria from tomato plant is treated against *Fusarium oxysporum* f. sp. *lycopersici* a pathogen to tomato shows promising results in suppressing fungus [33]. Still much research are progressing well in laboratory and under field trial against plant pathogen and pest in all over the world but agricultural industry is in an urgency for the product commercialization.

Microbes role in diverse environment

Microbes holds various strategies to overcome environmental stress which in produce cyst and spores, alters cellular membranes, autonomous enzymes for damage resistance [34]. More

interestingly, the bacteria from river soil responds differently to anthropogenic impacts and provides an alternative indicator of stream health [35]. The soil microbes (*Pseudomonas aeruginosa*) producing bacteriocins are natural anti-microbial peptides and have possible application as food preservation and healthcare [36]. The microbes produce different enzyme under selected pressure in environment, these microbes are more efficient in producing enzymes and a key for pharma and food industries. A recent study in Brazil shows a promising approach in finding a potential lipolytic bacteria from a biological treatment system from a hotel. *Acinetobacter baylyi*, a popular genus is identified for lipase enzyme production [2]. Similarly, *Acinetobacter calcoaceticus* [37], *Acinetobacter radioresistens* strain CMC-2 [38], *Acinetobacter* sp. SY-01 [39], *Acinetobacter radioresistens* CMC-1 [40], and *Acinetobacter johnsonii* [41] are previously reported as an effective group of microbes in large scale lipase enzyme production. In recent research in proteolytic enzyme search, a *Thermococcus kodakarensis* homologous to *archaemetzincins* family shows proteolytic activity and apart from this many multiple applications like inhibiting various pathogenic microbes, can be utilized in food industry to increase shelf life of food by controlling food borne pathogens, can utilize as a additive in laundry detergent [42-44]. With this broad aspect of microbes and their role in different environment are more interesting to study. Research in Insect gut, Human digestive track, Plant-Bacteria interaction and microbes from different environment have a bright future of research scope for young scientist with a constructive prospect.

Conclusion

The microbes from different environment with a vast diversity holds both beneficial and harmful pathogens. In this many species are beneficial to humans in various aspects like bioremediation, biotechnology applications, environmental biomarkers, symbiosis with human-insect-plants. In this insect gut bacteria are targets for biotechnological applications, as they produce many metabolites which are use full in industries. Insect parasitic symbiosis is also more interested direction to be studied more for pest control aspects. Recent studies shows plant symbiotic bacteria shares metabolites, promotes plant growth and antagonistic behaviour against fungus infection in plant. Still there is no clear picture of this mutualism with microbes- plants, the major hindrance is identifying the specific endophytic microbes which are not easily be cultivable *in vitro*. The microbes as bio marker is also a remarkable approach to identify the disease as early as possible. Though microbes are capable of producing novel metabolites, enzymes, bioactive molecules their present need to society is abundant. A broad research in microbial ability using modern genomics could help us to understand more for future direction.

Bibliography

1. Jayaraman Suriya., *et al.* "Identification of a Novel Gene Through the Metagenomic Approach to Degrade the Targeted Pollutant". *Microbial Biodegradation of Xenobiotic Compounds* (2019): 204.
2. Furini Graciane., *et al.* "Production of lipolytic enzymes by bacteria isolated from biological effluent treatment systems". *Anais da Academia Brasileira de Ciências* 90.3 (2018): 2955-2965.
3. Gurung Neelam., *et al.* "A broader view: microbial enzymes and their relevance in industries, medicine, and beyond". *BioMed research international* 2013 (2013).
4. Wigglesworth VB. "Digestion in the tsetse-fly: a study of structure and function". *Parasitology* 21.3 (1929): 288-321.
5. Broderick Nichole A., *et al.* "Contributions of gut bacteria to *Bacillus thuringiensis*-induced mortality vary across a range of Lepidoptera". *BMC Biology* 7.1 (2009): 11.
6. Kannan Mani., *et al.* "An efficient method for extraction of genomic DNA from insect gut Bacteria-Culture dependent". *Current Research in Microbiology and Biotechnology* 3.1 (2015): 550-556.
7. Otagiri Masato., *et al.* "Heterologous expression and characterization of a glycoside hydrolase family 45 endo- β -1, 4-glucanase from a symbiotic protist of the lower termite, *Reticulitermes speratus*". *Applied Biochemistry and Biotechnology* 169.6 (2013): 1910-1918.
8. Tsukagoshi Hikaru., *et al.* "The GH26 β -mannanase RsMan26H from a symbiotic protist of the termite *Reticulitermes speratus* is an endo-processive mannohydrolase: heterologous expression and characterization". *Biochemical and Biophysical Research Communications* 452.3 (2014): 520-525.
9. Prem Anand A., *et al.* "Isolation and characterization of bacteria from the gut of *Bombyx mori* that degrade cellulose, xylan, pectin and starch and their impact on digestion". *Journal of Insect Science* 10.1 (2010): 107.
10. ArunPrasanna Vimalanathan., *et al.* "Characterization of amylase producing *Bacillus megaterium* from the gut microbiota of silkworm *Bombyx mori*". *Journal of Chemistry and Environment* (2014).
11. Gangwar Avdresh Kumar., *et al.* "Applicability of microbial xylanases in paper pulp bleaching: a review". *BioResources* 9.2 (2014): 3733-3754.
12. Kannan M., *et al.* "Insect gut as a bioresource for potential enzymes-an unexploited area for industrial biotechnology". *Bio-catalysis and Agricultural Biotechnology* (2019).
13. Govindarajan RK., *et al.* "Purification, structural characterization and biotechnological potential of tannase enzyme produced by *Enterobacter cloacae* strain 41". *Process biochemistry* 77 (2019): 37-47.
14. Borlee Bradley R., *et al.* "Quorum-sensing signals in the microbial community of the cabbage white butterfly larval midgut". *The ISME journal* 2.11 (2008): 1101-1111.
15. Yadav Ruby and Pratyosh Shukla. "Probiotics for human health: Current progress and applications". *Recent advances in Applied Microbiology*. Springer, Singapore (2017): 133-147.
16. Krishnan Muthukalingan., *et al.* "Insect gut microbiome-An unexploited reserve for biotechnological application". *Asian Pacific journal of tropical biomedicine* 4 (2014): S16-S21.
17. Thiyonila Berchmans., *et al.* "Dung beetle gut microbes: Diversity, metabolic and immunity related roles in host system". *International Journal of Scientific Innovations* 1 (2018).
18. Kannan Mani. "Research on Insect Science could be a Wonderful Opportunity for Next Generation". *Acta Scientific Biotechnology* 1.2 (2020).
19. Khyami Mohammad and Essam Alyamani. "Aerobic and facultative anaerobic bacteria from gut of red palm weevil (*Rhynchophorus ferrugineus*)". *African Journal of Biotechnology* 7.10 (2008).
20. Shi Weibing., *et al.* "Comparison of insect gut cellulase and xylanase activity across different insect species with distinct food sources". *Bioenergy Research* 4.1 (2011): 1-10.
21. Delalibera Jr Italo., *et al.* "Contrasts in cellulolytic activities of gut microorganisms between the wood borer, *Saperda vestita* (Coleoptera: Cerambycidae), and the bark beetles, *Ips pini* and *Dendroctonus frontalis* (Coleoptera: Curculionidae)". *Environmental Entomology* 34.3 (2005): 541-547.
22. Anbalagan Sankarappan., *et al.* "The midgut bacterial flora of the hard tick *Hyalomma anatolicum* (Acari: Ixodidae) from South India as determined by molecular analyses". *Turkish Journal of Veterinary and Animal Sciences* 38.5 (2014): 520-525.
23. Wada-Katsumata Ayako., *et al.* "Gut bacteria mediate aggregation in the German cockroach". *Proceedings of the National Academy of Sciences* 112.51 (2015): 15678-15683.
24. Chen Bosheng., *et al.* "Biodiversity and activity of the gut microbiota across the life history of the insect herbivore *Spodoptera littoralis*". *Scientific reports* 6.1 (2016): 1-14.
25. Somerville., *et al.* "Aseptic Rearing and Infection with Gut Bacteria Improve the Fitness of Transgenic Diamondback Moth, *Plutella xylostella*". *Insects* 10.4 (2019): 89.

26. Xiang Hui., *et al.* "Bacterial community in midguts of the silkworm larvae estimated by PCR/DGGE and 16S rDNA gene library analysis". (2007): 222-233.
27. Chen Bosheng., *et al.* "Gut bacterial and fungal communities of the domesticated silkworm (*Bombyx mori*) and wild mulberry-feeding relatives". *The ISME Journal* 12.9 (2018): 2252-2262.
28. Zitvogel Laurence., *et al.* "Anticancer effects of the microbiome and its products". *Nature Reviews Microbiology* 15.8 (2017): 465.
29. Ramos Azucena and Michael T Hemann. "Drugs, bugs, and cancer: *Fusobacterium nucleatum* promotes chemoresistance in colorectal cancer". *Cell* 170.3 (2017): 411-413.
30. Wu Na., *et al.* "Dysbiosis signature of fecal microbiota in colorectal cancer patients". *Microbial Ecology* 66.2 (2013): 462-470.
31. Hogenhout Saskia A., *et al.* "Insect vector interactions with persistently transmitted viruses". *Annual Reviews on Phytopathology* 46 (2008): 327-359.
32. Zhang Nan., *et al.* "Comparative genomic analysis of *Bacillus amyloliquefaciens* and *Bacillus subtilis* reveals evolutionary traits for adaptation to plant-associated habitats". *Frontiers in Microbiology* 7 (2016): 2039.
33. Shahzad Raheem., *et al.* "Plant growth-promoting endophytic bacteria versus pathogenic infections: an example of *Bacillus amyloliquefaciens* RWL-1 and *Fusarium oxysporum* f. sp. *lycopersici* in tomato". *Peer Journal* 5 (2017): e3107.
34. Storz Gisela and Regine Hengge. Bacterial stress responses. American Society for Microbiology Press (2010).
35. Sankarappan Anbalagan. "Effective Stream Health Assessment: Soil Microbes Versus Aquatic Insects". *Ecologia* 7.(2017): 1-11.
36. Arumugam Thangarasu., *et al.* "Inhibition of methicillin resistant *Staphylococcus aureus* by bacteriocin producing *Pseudomonas aeruginosa*". *International Journal of Peptide Research and Therapeutics* 25.1 (2019): 339-348.
37. Kok Ruben G., *et al.* "Characterization of the extracellular lipase, LipA, of *Acinetobacter calcoaceticus* BD413 and sequence analysis of the cloned structural gene". *Molecular microbiology* 15.5 (1995): 803-818.
38. Ng I-Son., *et al.* "Purification and characterization of extracellular lipase from *Acinetobacter radioresistens* CMC-2". *Journal of the Chinese Institute of Chemical Engineers* 30.5 (1999): 355-362.
39. Han Soo-Jin., *et al.* "Expression and characterization of a novel enantioselective lipase from *Acinetobacter* species SY-01". *Biochimie* 85.5 (2003): 501-510.
40. Hong Ming Chaun and Ming Chung Chang. "Purification and characterization of an alkaline lipase from a newly isolated *Acinetobacter radioresistens* CMC-1". *Biotechnology letters* 20.11 (1998): 1027-1029.
41. Wang Hai Kuan., *et al.* "A novel low-temperature alkaline lipase from *Acinetobacter johnsonii* LP28 suitable for detergent formulation". *Food Technology and Biotechnology* 49.1 (2011): 96-102.
42. Jia Baolei., *et al.* "A zinc-dependent protease AMZ-tk from a thermophilic archaeon is a new member of the archaemetzincin protein family". *Frontiers in microbiology* 6 (2015): 1380.
43. Jaouadi Bassem., *et al.* "Biochemical and molecular characterization of a detergent-stable serine alkaline protease from *Bacillus pumilus* CBS with high catalytic efficiency". *Biochimie* 90.9 (2008): 1291-1305.
44. Jain Deepti., *et al.* "Purification and characterization of halo-alkaline thermoactive, solvent stable and SDS-induced protease from *Bacillus* sp.: a potential additive for laundry detergents". *Bioresource Technology* 115 (2012): 228-236.

Assets from publication with us

- Prompt Acknowledgement after receiving the article
- Thorough Double blinded peer review
- Rapid Publication
- Issue of Publication Certificate
- High visibility of your Published work

Website: www.actascientific.com/

Submit Article: www.actascientific.com/submission.php

Email us: editor@actascientific.com

Contact us: +91 9182824667