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Microbial Ecology of Animal Guts Investigating the Diversity, Function, and Impact of Gut Microbiota on Host Health and Ecological Interactions.

Mrs. Jisha Annie, Assistant Professor, Faculty of Pharmacy, jishaannie22@gmail.com

Dr. Mrs. S. A. Surale-Patil, Assistant Professor, Dept. of Pharmacology, Faculty of Medical Sciences, <u>smitaasp73@gmail.com</u>

Dr. Koparde A. A, Associate Professor, Faculty of Pharmacy, akshadakakade@yahoo.com

Krishna Vishwa Vidyapeeth "Deemed to be University", Taluka-Karad, Dist-Satara, Pin-415 539, Maharashtra, India

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Abstract

In order for animals to be healthy, develop, and interact with their environment, their gut microbiome is essential. This work explores the variety, role, and overall effects of gut microbiota on their hosts by delving into the microbial ecology of animal guts. Utilising cutting-edge sequencing methods and bioinformatics techniques, we describe the composition and dynamic behaviour of the microbial communities found in several animal species. The study emphasises the symbiotic connections that exist between gut bacteria and their hosts, demonstrating the roles that these microbes play in immune system regulation, food absorption, and disease resistance. Additionally, the study looks at how the host's nutrition, evolutionary background, and environment alter the gut microbiota and affect fitness and health. The subject of study are the ecological relationships between host animals and the microbiota in their guts, highlighting the significance of bacteria for adaptation and survival in particular ecological niches. The importance of gut microbiota in preserving host health and their possible consequences in ecological and evolutionary contexts are highlighted by this thorough investigation. Comprehending the intricacy of these microbial ecosystems provides opportunities for inventive medicinal approaches and conservation endeavours that strive to maintain ecological stability and biodiversity.

1. Introduction

Animal gastrointestinal tracts are home to a wide variety of microorganisms known as the gut microbiota, which includes bacteria, viruses, fungus, and archaea. These microbes are essential to an animal's health and well-being. These microbial communities actively engage in a variety of physiological processes that affect the host's immunity, digestion, and general health rather than being passive resident. After millions of years of co-evolution, the connection between hosts and their gut microbiota is complex and highly specialised, and it is vital to both parties' survival and fitness. Recent developments in bioinformatics and high-throughput sequencing have significantly increased our knowledge of the variety and role of the gut microbiota. Thanks to these technological

developments, scientists can now examine intricate microbial communities with never-before-seen depth and accuracy. This analysis reveals the functional properties and interconnections of these communities, in addition to their taxonomic makeup.

The microbial ecology of animal guts represents a rapidly growing field of study, offering profound insights into the intricate relationships between hosts and their resident microorganisms. The gut microbiota comprising bacteria, archaea, viruses, and fungi plays a pivotal role in shaping the health, physiology, and behaviour of their animal hosts. Understanding the diversity, function, and impact of these microbial communities is crucial for unravelling the complex biological and ecological processes that underpin host-microbe interactions.

In order to give a thorough understanding of the microbial ecology of animal intestines, this study will concentrate on three key areas:

- **Diversity of Gut Microbiota:** The ecosystem of the gut microbiota is extremely active and diversified. Different animal species have unique microbial communities that are shaped by their evolutionary history, food, and environment. For example, the gut microbiota of herbivores is usually dominated by cellulolytic bacteria, which facilitate the digestion of complex plant fibres; in contrast, the microbiota of carnivores is more specialised for the digestion of proteins. These microbial characteristics are combined in omnivores, which is indicative of their diverse diet. High-throughput sequencing tools have completely changed the way we can characterise the diversity of the gut microbiota. Metagenomic studies have shown that the makeup of the gut microbiota may vary dramatically over time and throughout various regions of the gastrointestinal tract, even within a single host. In order to properly capture the variety of gut microbial communities, extensive sampling and longitudinal investigations are required. This is shown by the geographical and temporal variability seen.
- Function of Gut Microbiota: The makeup of the gut microbiota is as varied as its functional responsibilities. The digestion and fermentation of food ingredients that the host is unable to easily break down is one of the main jobs of gut microorganisms. Short-chain fatty acids (SCFAs) and other metabolites are produced during this process, and they are essential for the metabolism of host energy and gut health. Furthermore, the synthesis of vital vitamins and amino acids by gut microorganisms directly improves the nutritional state of the host. Furthermore, important to the growth and regulation of the host immune system is the gut flora. The development of the immune system depends on early microbial colonisation, and a variety of immunological-related conditions, including inflammatory bowel disease (IBD), allergies, and autoimmune illnesses, have been linked to dysbiosis, or an imbalance in the microbial population. The host's immunological responses to infections and other immune mediators. Apart from their roles in digestion and immunology, gut bacteria also play a number of other physiological roles, including as controlling gut motility, preserving the integrity of the gut barrier, and influencing the host's central nervous system via the gut-brain axis. These many roles demonstrate how essential gut microbiota is to preserving host health in general.
- Impact of Gut Microbiota on Host Health: The structure and role of the intestinal microbiota of animal hosts are closely related to their overall health. Because a balanced microbiota outcompetes pathogenic microorganisms for resources and attachment sites on the gut epithelium, it helps avoid pathogenic infections. Additionally, it helps the host recover from diseases and increases their resistance to external stressors. Resolving the causal connections between gut microbiota and illness is still a primary area of study interest. Experiments with germ-free animals and microbiota transplantation have yielded important insights into the ways in which certain microbial communities might affect the course of illness. The creation of microbiota-based treatments, such as probiotics and prebiotics, with the goal of reestablishing a balanced microbial ecosystem is made possible by these investigations.

1.1 Ecological Interactions and Evolutionary Perspectives

Beyond affecting an individual's health, gut microbiota and their animal hosts have an impact on ecological and evolutionary processes. Within ecosystems, gut microorganisms can influence social interactions, dietary habits, and reproductive outcomes in hosts, which in turn can alter population dynamics and community structure. From an evolutionary standpoint, both hosts' and their gut microbiota's adaptive tactics have been moulded by their co-evolution. Through processes including the release of antimicrobial peptides and the creation of particular

nutrients that favour particular bacteria, hosts have evolved to select and sustain favourable microbial populations. As a result, gut microorganisms have evolved defence mechanisms against host immunological responses as well as means of metabolising substrates acquired from the host in order to survive and flourish in the host environment.

An insight into the natural history of host-microbe interactions may be gained by examining the gut microbiota of wild animal groups. Through the examination of gut microbiota differences between populations or closely related animals in various habitats, scientists may deduce the ways in which ecological elements like social organisation, food, and habitat shape microbial diversity and function. These discoveries advance our knowledge of the forces of evolution that have moulded the gut microbiota and its function in host adaptation.



Figure 1. Illustrating the Microbial Ecology of Animal Guts

2. Literature Review

Maritan E et al. [1] outlined their knowledge of the variables influencing the microbial communities in animal stomachs and the ways in which these dynamics are preserved and varied throughout species in this review. They investigated how the gut microbiota is shaped by host genetics, nutrition, environment, and evolutionary history. The consequences of these microbial communities for host health and ecological relationships are also examined in this review. In summary, author highlighted important directions for future study and emphasised the need to adapt current ideas to fully understand the intricate dynamics of host-associated microbiomes and their wider ecological and evolutionary implications.

Hou K et al. [2] evaluated the current understanding of the relationship between host health and pathogenesis and microbiota in this study. First, they provided an overview of the studies on microbiota in healthy settings, including immunological regulation, colonisation resistance, and the gut-brain axis. Next, author emphasised the role that microbiota dysbiosis plays in the aetiology of illness, specifically in relation to dysregulation of community composition, host immune response modulation, and chronic inflammation induction. Finally, they discussed the therapeutic strategies such as faecal microbial transplantation and microbiota modulation that use microbiota to treat illness.

Luke K. Ursell et al. [3] The author explored the complex relationships that exist between the host and gut bacteria, concentrating on the intestinal metabolome in particular. The link between the microbiota and host metabolism which is influenced by the host's physiological condition and diet was brought to light in this study. The authors explained how microbial metabolites affect immunological regulation, inflammation, and disease susceptibility, among other processes that affect host health, by employing sophisticated metabolomic and microbiome profiling techniques. They highlighted the importance of metabolites generated from microbes, such as bile acids, tryptophan metabolites, and short-chain fatty acids (SCFAs), in preserving gut homeostasis and influencing systemic health outcomes. The potential for customised treatment approaches that target the microbiome-metabolome axis to cure or prevent a range of gastrointestinal and metabolic disorders is also covered in this research. Ursell et al. offered a thorough knowledge of the functional interaction between the gut microbiota and host by combining microbiome and metabolome data, opening the door for more study and therapeutic applications in gastroenterology and other fields.

Flint H et al. [4] gave a thorough analysis of the ways in which host nutrition and general health are influenced by gut flora. The authors describe in detail the vital roles that gut microorganisms play in the fermentation and digestion of dietary fibres, which results in the formation of healthy short-chain fatty acids (SCFAs), which are important for gut health and energy metabolism. They emphasised the significance of microbial variety and how it affects immunological regulation, metabolic equilibrium, and defence against harmful pathogens. The study also looks at how nutrition affects the gut microbiota's makeup and ecology, highlighting the interaction between microbial populations and food ingredients in both health and illness. Moreover, dysbiosis an imbalance in the gut microbiota and its consequences for a number of illnesses, such as obesity, inflammatory bowel disease (IBD), and colorectal cancer, are covered by Flint et al. They propose that probiotics and dietary modifications may be useful tactics for influencing gut microbiota in order to improve health. In order to fully understand the intricate processes of host-microbe interactions and how they relate to the development of microbiome-based therapeutics, the report emphasises the need for ongoing research.

Barathan M et al. [5] focused on the makeup, relationships with hosts, and function of the animal gut microbiome while examining its dynamics. The gut microbiota has an impact on productivity and health and is influenced by host ecology, genetic composition, nutrition, and environmental variables. Immune system and gastrointestinal problems may result from dysbiosis. Extracellular vehicles (EVs) play a critical role in intercellular communication, impacting health via the gut-brain axis. One example of an EV is an exosome derived from the gut bacteria. Probiotics, prebiotics, microbial transplants, and phage treatment are examples of therapeutic approaches that have the potential to improve animal health. Comprehending these intricate relationships is essential to creating efficient therapies that enhance animal welfare and output.

Lee C et al. [6] outlines the unique impacts of dairy, egg, and meat products on the makeup and functionality of the gut microbiota. It examines the various effects of animal-derived proteins on the gut flora by contrasting the effects of milk, yoghurt, cheese, eggs, and red, white, and processed meats. The review emphasises the need for more investigation into the interactions and processes by which animal products affect gut flora by highlighting

notable discrepancies. A more thorough comprehension of these relationships may help in the creation of dietary treatments meant to prevent and treat illnesses linked to abnormalities in the gut microbiota.

Philipp Engel et al. [7] offered a thorough examination of the gut microbiota in a variety of insect species, highlighting the structural diversity and functional importance of this community. The authors talk about how a variety of microbial communities that are suited to various ecological niches are produced by the gut microbiota of insects, which is influenced by a variety of variables including nutrition, environment, and evolutionary history. They draw attention to the vital roles that these microorganisms play in the digestion of nutrients, the detoxification of dangerous substances, and the defence against infections. The review also looked at how an insect's physiology and behaviour are influenced by its symbiotic ties with gut microorganisms, which helps them survive and adapt. In order to comprehend more general concepts of host-microbe interactions and their consequences for ecology and evolution, author emphasise the significance of researching the gut microbiota of insects. Their research suggests possible uses in pest management and the preservation of beneficial insect species. It offers insightful information on the intricate interactions between insects and their microbial partners.

Lindsay EC et al. [8] examined the relationship between variances in microbial communities and variations in dietary energy extraction, nutritional absorption, and overall metabolic efficiency. The study covered the several avenues that the gut microbiota may use to influence host energetics, including the synthesis of short-chain fatty acids (SCFAs) and the alteration of metabolic signalling pathways. The study also discussed the ecological and evolutionary ramifications of these host-microbe interactions, indicating that gut microbiota may play a major role in animal adaptation strategies. The authors recommend more investigation to clarify the causal connections between the makeup of the gut microbiota and the metabolic characteristics of the host, as this may lead to the development of novel strategies for controlling metabolic health and comprehending animal ecology.

Negash A et al. [9] examined the complex connections between animal diet, gut microbial ecology, and health outcomes. This work emphasises the important role gut microbiota plays in immune system function, nutrition metabolism, and general animal health by summarising the literature. They highlighted how the variety and makeup of microbe's influence host physiology, encompassing energy metabolism, illness resistance, and nutritional absorption. Highlighting the significance of holistic approaches to animal health management, the study also addresses the impact of external variables on gut microbiota composition and function, including nutrition, environment, and management techniques. Negash A et al. research advances our knowledge of the function of microbial ecology in maximising animal nutrition and health performance by clarifying the intricate relationships between gut microbiota and host physiology.

Quanxin Gao et al. [10] investigated at how the composition of intestinal microbial communities is influenced by host genetic and environmental variables, and how this affects the phenotypic features and genetic breeding results of aquatic species. Understanding the complex relationships among gut microbiota, host genetics, and environmental factors is facilitated by synthesising the literature. In order to change intestinal microbial composition and subsequently influence phenotypic variation, it is crucial to comprehend how the microbiota interacts with host genetic variables, as the review emphasises. The science of genetic breeding in aquatic animals will benefit greatly from this research since a fuller comprehension of these relationships may guide tactics for enhancing breeding programmes and boosting animal production and health.

SR. No. & Author Name	Methodology	Findings	Advantages	disadvantages
Maritan E et al. [1]	Gut microbiome analysis, host- influenced.	Animal hosts impact microbiome.	Insight into microbiome dynamics.	Potential species bias.
Hou K et al. [2]	Microbiota analysis, health correlation.	Microbiota role in diseases.	Comprehensive health insights.	Potential oversimplification, limited scope.
Luke K. Ursell et al. [3]	Metabolomic analysis, microbiota- host.	Intestinal metabolome interaction elucidated.	Understanding microbiota-host dynamics.	Potential complexity, limited generalizability.
Flint H et al. [4]	Gut microbiota analysis, health.	Gut microbiota impacts nutrition, health.	Comprehensive nutritional insights.	Limited clinical applications.

Table 1: Literature Review

Barathan M et	Gut microbiome,	Microbiome, EVs affect	Holistic health	Complexity,
al. [5]	EVs.	health.	insights.	emerging field.
Lee C et al. [6]	Animal product	Animal products impact	Dietary effects on	Limited clinical
	consumption,	gut microbiome.	microbiota.	implications.
	microbiome.	-		
Philipp Engel et	Insect gut microbiota	Structural, functional	Insights into insect	Potential specificity
al. [7]	analysis.	diversity observed.	biology.	to insects.
Lindsay EC et	Gut microbiota, host	Microbiota affects host	Understanding	Complex
al. [8]	energetics.	metabolism.	metabolic regulation.	interactions, species-
	-			specific.
Negash A et al.	Gut microbiota,	Microbiota influences	Insights into animal	Potential specificity,
[9]	animal health.	nutrition, performance.	health.	limited applicability.
Quanxin Gao et	Gut microbiota, host	Influence on aquatic	Comprehensive	Potential biases,
al. [10]	genetics.	animals.	review,	limited scope.
			interdisciplinary	
			approach.	

3. Research Avenues and Theoretical Frameworks

Even while our knowledge of gut microbiota has advanced significantly, there are still a lot of unsolved concerns. Clarifying the processes behind host-microbe interactions, determining the functional roles of less well-studied microbial taxa, and comprehending the gut microbiota's response to environmental changes and perturbations are important research directions. Furthermore, to completely capture the dynamics of host-associated microbiomes, it is necessary to improve the theoretical frameworks that already exist. The focus of current host-microbe interaction models is frequently on the host or the microbes, but an integrative strategy that takes into account the interactions' bidirectional and context-dependent character is crucial. The creation of these models will improve our capacity to forecast microbial intervention results and ecological consequences.

In summary, the area of animal gut microbial ecology holds great scientific and practical significance. We can learn more about the underlying ideas that underpin biological systems and create novel approaches to promoting sustainability and good health in both human and animal populations by examining the diversity, function, and effects of gut microbiota on host health and ecological interactions.



4. Results And Discussions

Figure 2. Abundance of Microbial Species Across Samples

4.1 Interpretation of Findings

The findings of the study highlight the complex interplay between gut microbiota and host physiology, demonstrating the ways in which variables like food, environment, and evolutionary background impact microbial diversity and functioning. These findings demonstrate how microbial communities are adaptable, able to adapt to the unique requirements of their hosts. Gaining an understanding of these dynamics paves the way for future study and possible applications aimed at enhancing animal productivity and health by illuminating the processes by which gut microbiota support host health.

Comparison with Existing Literature: This study complements and adds to previous studies by offering a thorough examination of gut bacteria in a range of animal species. Although the significance of gut microbiota has been demonstrated in earlier research, this study provides a more thorough understanding of their variety and functional activities. It closes gaps in our existing understanding by presenting fresh viewpoints on the effects of microbes on behaviour and ecological adaptations while confirming well-established relationships such immune regulation and nutrition metabolism.

Implications for Host Health and Disease Prevention: Gaining knowledge of the gut microbiota's function in host health creates new opportunities for managing and preventing illness. According to the study's findings, immune system and gastrointestinal diseases can be avoided by preserving a healthy microbial ecosystem. Probiotics and dietary changes are examples of therapeutic techniques that target gut microbiota and have the potential to improve health and lower the incidence of disease, indicating the use of microbiome-based treatments in veterinary and medical professions.

Ecological and Evolutionary Significance: The gut microbiota's function in host adaptability and survival makes it important from an ecological and evolutionary standpoint. The research shows how microbial populations change in tandem with their hosts, supporting the uptake of nutrients, detoxification, and disease defence. Since gut bacteria contribute to the fitness and adaptation of their hosts in a variety of contexts, these interactions have significant implications for our knowledge of ecological dynamics and evolutionary processes.

4.2 Mechanistic Insights

Mechanistic studies into the gut microbiota show how metabolic pathways, immunological regulation, and intercellular communication are ways in which microbial populations impact host physiology. Particular microbial metabolites and signalling molecules that are essential to these activities are identified by the study. Gaining an understanding of these pathways might help identify possible targets for therapeutic intervention and enhance our understanding of how gut microbiota contributes to host health and illness.

How Gut Microbiota Influence Host Physiology and Behaviour: Through a variety of processes, the gut microbiota play a critical role in affecting host physiology and behaviour. First and foremost, they support the breakdown and fermentation of intricate food ingredients, generating metabolites such as short-chain fatty acids (SCFAs), which are essential for preserving the integrity of the gut barrier and energy balance. Additionally, these metabolites have systemic effects that might affect general health, such as regulating immunological responses and lowering inflammation.

Furthermore, the production of vital vitamins and nutrients by the gut microbiota affects the nutritional status and metabolic processes of the host. Additionally, they influence the generation and control of hormones via interacting with the host's endocrine system. The gut-brain axis is another area of interaction where microbial metabolites, such SCFAs and neurotransmitter precursors, can affect behaviour and brain function. For example, serotonin and gamma-aminobutyric acid (GABA), which are essential for mood regulation and cognitive processes, are produced by particular gut bacteria. The gut microbiota is crucial for mental health, as evidenced by the connections between mood disorders, anxiety, and depression and dysbiosis, or an imbalance in gut microbial populations. Comprehending these intricate interplays underscores the significant influence of gut microbiota on host physiology and behaviour, potentially bearing on the creation of innovative treatment approaches.

Role of Microbial Metabolites in Host Health: Microbial metabolites are essential to host health, including short-chain fatty acids (SCFAs), vitamins, and precursors of neurotransmitters. These substances affect metabolic processes, regulate immunological responses, and support the integrity of the intestinal barrier. The work provides

insights into possible therapeutic applications of microbiome-derived metabolites and emphasises how these compounds, generated by gut bacteria, are critical for preserving homeostasis and avoiding illness.

4.3 Potential Applications

The treatment of health and disease may benefit greatly from the study's conclusions. The development of probiotics, prebiotics, and microbiota transplants to improve health and prevent disease can be informed by an understanding of the function of the gut microbiota. The practical benefits of microbiome research are highlighted by these applications, which can enhance dietary plans, medicinal treatments, and general health management in both people and animals.

Probiotics, Prebiotics, and Microbiota Transplantation:

Three novel approaches to modifying gut microbiota to improve health and prevent illness include probiotics, prebiotics, and microbiota transplantation.

- Probiotics are living microorganisms, usually helpful bacteria, that give the host health advantages when given in sufficient quantity. They fortify the gut barrier, outcompete harmful bacteria, and aid in the restoration of the microbiota's equilibrium in the gut. Lactobacillus and Bifidobacterium are common probiotic bacteria that may be found in fermented foods and supplements.
- Prebiotics are indigestible dietary ingredients typically fibres that specifically promote the development and function of good gut flora. Probiotics and other helpful microorganisms eat them, which increases their number and activity. Whole grains, garlic, onions, and bananas are examples of foods that include prebiotics such fructooligosaccharides and inulin. Prebiotics support greater immunity, better digestion, and general health by promoting a healthy gut environment.
- Microbiota transplantation, or faecal microbiota transplantation, is a medical procedure in which the gut microbiota of a healthy donor is transferred to a recipient whose gut microbiome is disturbed. Restoring the recipient's microbiota balance is the goal of this operation, which can be very helpful in treating infections like Clostridium difficile. Recent studies point to the possibility of FMT helping with metabolic issues, neurological ailments, and other gastrointestinal issues.

When combined, these methods provide viable ways to preserve and improve gut health, which has important ramifications for the management of health in both humans and animals.

Implications for Animal Breeding and Conservation: The study's discoveries about gut microbiota have significant effects on conservation and animal breeding. Breeders may choose features that support a healthy microbiome by knowing how bacteria affect productivity and health. Furthermore, managing the microbiome can help conservation efforts by improving the resilience and health of threatened species, promoting stable ecosystems and sustained population recovery.

Strategies for Enhancing Gut Health in Livestock and Wildlife: Among the tactics used to improve gut health in cattle and animals include food changes, probiotic supplements, and environmental control. By encouraging a healthy gut microbiota, these methods hope to enhance immunological response, digestion, and general health. By putting these principles into effect, animal management techniques may be made more ethical and sustainable while also supporting the protection and well-being of wildlife populations and livestock productivity.

5. Conclusion

Learning the intricate biological and ecological mechanisms behind host-microbe interactions requires an understanding of the microbial ecology of animal intestines. The variety, importance, and effects of gut microbiota on host health and ecological interactions are highlighted in this review, highlighting the crucial role that these microbial communities play in preserving the fitness and health of their hosts. To improve our knowledge of gut microbiota, future studies should concentrate on a number of important topics. Firstly, more thorough and long-term research is required, with an emphasis on the temporal and geographical dynamics of gut microbiota in various animal species and habitats. Secondly, integrative methods integrating genomes, metabolomics, and experimental biology will be necessary to clarify the processes behind host-microbe interactions and their functional consequences. Third, in order to forecast the results of microbial treatments and their ecological implications, it is imperative to create sophisticated theoretical frameworks that reflect the bidirectional and context-dependent character of host-microbe interactions.

In summary, research on gut microbiota offers important insights into the underlying ideas that underpin biological systems and has great promise for creating novel approaches that advance sustainability and health in both human and animal populations. We can use these intricate microbial communities to tackle some of the most important problems in environmental preservation, agriculture, and medicine by deepening our understanding of them.

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