THE GI MICROBIOME IN DOGS. ALL DISEASES BEGIN IN THE GUT

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The gastrointestinal microbiome constitutes a sophisticated and ever-changing assembly of microorganisms that inhabit the digestive tracts of animals. This community comprises bacteria, viruses, fungi, and protozoa, exhibiting variations in relative abundance across different animal species and individuals. Established during birth and continuously shaped throughout life, this microbiome plays a pivotal role in regulating digestion, facilitating nutrient absorption, and modulating immune function. This discourse aims to provide an overview of the gastrointestinal microbiome, elucidating its composition, function, and significance in animal health, while also exploring potential applications of microbiome research in veterinary medicine.

Composition of the Gastrointestinal Microbiome:

The predominant microorganisms in the gastrointestinal microbiome are bacteria, classified into various taxonomic levels such as phyla, class, order, family, and genus. Along the gastrointestinal tract (GIT), bacterial diversity and abundance gradually increase, with the colon and feces harboring the most diverse microbial communities. In adult dogs and cats, the prevailing bacterial phyla include Firmicutes, Bacteroidetes, Proteobacteria, Fusobacteria, and Actinobacteria. However, relative abundances may vary due to factors like host genetics, diet, age, and methodological discrepancies in sample collection and analysis. Apart from bacteria, other microorganisms like viruses and fungi also contribute to the functionality of the microbiome.

Function of the Gastrointestinal Microbiome:

The gastrointestinal microbiome exerts significant influence over digestion, nutrient absorption, and immune regulation.

Digestion:

Key roles in digestion include the breakdown of dietary fibers, proteins, bile salts, and choline. Certain bacterial phyla, like Bacteroidetes and Firmicutes, aid in the digestion of complex carbohydrates, yielding short-chain fatty acids (SCFAs) that serve as crucial energy sources and modulators of metabolic processes. Additionally, the microbiome assists in bile salt deconjugation and choline metabolism. Polyphenols found in plants are activated by gut microbes, leading to various beneficial effects, including detoxification and pathogen suppression.

Immune System:

The microbiome interacts with immune cells in the gastrointestinal tract, producing molecules that either stimulate or suppress immune responses. Mucosal immune defenses involve epithelial barriers, antimicrobial peptide secretion, and mucosal

lymphoid structures like Peyer's patches. Pattern recognition receptors (PRRs) on epithelial cells recognize microbial components, initiating immune responses. These interactions help maintain immune tolerance and regulate inflammation, crucial for gastrointestinal health.

Gut-Brain Axis:

Communication between the gastrointestinal tract and the brain occurs through the enteric nervous system (ENS). The microbiome influences the ENS via neurotransmitter production, barrier integrity maintenance, and immune modulation. Dysbiosis in the gut has been linked to emotional responses and neurological disorders, highlighting the bidirectional nature of the gut-brain axis.

Analyzing the Gastrointestinal Microbiome:

Fecal samples are commonly used for microbiome analysis due to accessibility. Various techniques like 16S rRNA gene sequencing and shotgun metagenomic sequencing enable microbial identification and functional analysis. Analytical tools assess diversity (alpha and beta diversity), functional capabilities, and metabolite profiles. This multidisciplinary approach offers insights into microbial composition, activity, and their impact on host health.

Gastrointestinal Metabolome:

Comprising diverse metabolites produced during digestion, the gastrointestinal metabolome influences immunity, inflammation, and intestinal integrity. Short-chain fatty acids (SCFAs), tryptophan derivatives, and bile acids modulate immune responses, barrier function, and inflammation. Metabolite analysis, utilizing techniques like GC-MS and LC-MS, provides insight into microbial activity and host-microbiome interactions, aiding in disease diagnosis and therapeutic target identification.

Importance in Animal Health:

Diet profoundly influences the gastrointestinal microbiome, impacting microbial composition and metabolic activity. Dysbiosis is associated with gastrointestinal diseases like acute and chronic diarrhea, emphasizing the microbiome's diagnostic and therapeutic relevance. Strategies like prebiotics, probiotics, synbiotics, and fecal microbiota transplantation (FMT) offer potential treatments for dysbiosis-related conditions. Understanding the gastrointestinal microbiome's role in health and disease enhances veterinary practices, promoting holistic approaches to animal care.

In conclusion, the gastrointestinal microbiome is a vital determinant of animal health, modulating various physiological processes and immune responses. Dysbiosis contributes to a spectrum of health issues, underscoring the importance of microbiome research in veterinary medicine. By leveraging advancements in microbiome analysis and therapeutic interventions, veterinarians can optimize animal health outcomes, recognizing the microbiome as a cornerstone of holistic veterinary care.