

# Forage Bioactive Compounds and Rumen Microbiota: Exploring Colonization, Density, and Diversity Dynamics in Dairy Cows

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## Abstract

*The rumen microbiota plays a pivotal role in the health, productivity, and sustainability of dairy cows, with forage serving as a fundamental dietary component influencing microbial dynamics. This review explores the intricate interactions between forage bioactive compounds – such as tannins, saponins, alkaloids, and flavonoids – and the rumen microbiota, focusing on their effects on colonization, microbial density, and diversity. Forage bioactives modulate microbial adhesion, biofilm formation, and growth patterns, driving shifts in microbial populations and functional guilds. These compounds exhibit selective antimicrobial properties, promoting beneficial microbes while suppressing pathogenic or methanogenic species. Additionally, the indirect effects of bioactive compounds, mediated through metabolic intermediates, pH regulation, and nutrient availability, further shape microbial community structure and function. This review synthesizes current knowledge on the molecular and ecological mechanisms underpinning these interactions, identifies research gaps, and highlights the implications of forage bioactives for sustainable dairy production. Understanding these dynamics provides valuable insights into optimizing forage-based diets to enhance rumen function, improve nutrient utilization, and mitigate environmental impacts.*

**Keywords:** Bioactive compounds, colonization, dairy cows, density, diversity, microbiota

## INTRODUCTION

The rumen microbiota of dairy cows represents a highly complex and dynamic ecosystem critical to nutrient digestion, fermentation, and overall animal productivity [1–3]. This intricate community of microbes, including bacteria, archaea, protozoa, and fungi, relies heavily on the composition and quality of dietary inputs [4, 5]. Forages, being the primary dietary component in dairy nutrition, not only serve as a vital source of energy and fiber but also introduce a variety of bioactive compounds with significant biochemical and ecological effects [6, 7]. These compounds, encompassing tannins, saponins, alkaloids, and flavonoids, have been shown to influence microbial activity and community structure, underscoring

the central role of forage in shaping rumen microbial dynamics [8–10].

Bioactive compounds in forages exert selective effects on microbial colonization, density, and diversity by modulating interactions at the microbial and host interface [11–13]. Their influence spans a range of processes, including microbial adhesion, biofilm formation, nutrient utilization, and interspecies competition [14–17]. These effects can alter the functional balance between cellulolytic, amylolytic, and methanogenic microbes, impacting rumen fermentation efficiency and overall cow performance. Despite growing recognition of these interactions, understanding the

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precise mechanisms by which forage bioactives regulate microbial dynamics remains incomplete. Additionally, factors, such as forage type, maturity, and preservation methods, further complicate these relationships, necessitating a comprehensive evaluation [18–21].

This review addresses a critical gap in knowledge by providing an in-depth analysis of how forage bioactive compounds influence colonization, density, and diversity dynamics within the rumen microbiota of dairy cows. Unlike previous studies, which have primarily focused on isolated effects, this work integrates insights across molecular, microbial, and ecological scales to offer a holistic perspective. By synthesizing evidence and identifying key interactions, this study aims to advance strategies for optimizing forage-based diets to enhance rumen function, dairy productivity, and sustainability, offering novel pathways for future research and practical applications.

## COMPOSITION OF BIOACTIVE COMPOUNDS

### Classes of Bioactive Compounds

Forage bioactives comprise diverse classes, including tannins, saponins, alkaloids, flavonoids, and terpenoids [8, 22]. Each class exhibits unique biochemical properties that interact differently with rumen microbes, influencing digestion, fermentation, and microbial metabolism. These compounds often act selectively, promoting beneficial microbes while suppressing harmful or methanogenic populations, thereby enhancing overall rumen function.

### Molecular Structures Influencing Microbial Interaction

The molecular structures of bioactive compounds, such as their polarity, charge, and functional groups, determine their mode of interaction with rumen microbes [23, 24]. These interactions can disrupt microbial membranes, inhibit enzymatic pathways, or modify interspecies signaling, directly influencing the colonization, density, and diversity of rumen microbial populations and their metabolic activities.

### Solubility and Bioavailability in the Rumen

The solubility and bioavailability of forage bioactive compounds in the aqueous rumen environment significantly affect their efficacy [25]. Compounds that dissolve readily can interact more effectively with microbial cells, while poorly soluble ones may remain less active. Enhanced bioavailability often correlates with improved microbial modulation and nutrient utilization efficiency.

### Concentration-Dependent Effects

The effects of bioactive compounds on rumen microbiota are highly concentration dependent. At optimal concentrations, these compounds can stimulate the growth of beneficial microbes or suppress harmful ones [26, 27]. However, at higher concentrations, they may exert toxic effects, disrupt microbial balance and reduce fermentation efficiency. Understanding these thresholds is crucial for dietary optimization.

## MODULATION OF RUMEN MICROBIAL COLONIZATION

### Effects on Adhesion of Microbes to Feed Particles

Forage bioactive compounds influence the ability of microbes to adhere to feed particles, a critical step in colonization [11, 14, 20]. By modifying microbial surface properties or binding sites on feed, these compounds can either enhance or inhibit microbial attachment, affecting the efficiency of fermentation and the subsequent breakdown of dietary components in the rumen.

### Influence on Biofilm Formation by Rumen Microbes

Bioactive compounds can regulate biofilm formation, which is essential for stable microbial communities. These compounds may promote biofilm formation by beneficial microbes, fostering robust enzymatic activity, or inhibit pathogenic biofilms, reducing competition [11, 14, 20]. This modulation impacts nutrient breakdown, microbial resilience, and the overall efficiency of rumen fermentation processes.

### **Inhibition of Specific Microbial Species**

Bioactive compounds exhibit selective antimicrobial effects, inhibiting harmful or methanogenic microbes while promoting beneficial species, such as cellulolytic and amylolytic bacteria [11, 14, 20]. This selectivity reshapes microbial community composition, favoring species that enhance fiber digestion and nutrient utilization, thereby contributing to improve productivity and health in dairy cows.

### **Alteration of Colonization Dynamics**

These compounds can influence microbial colonization on the rumen epithelium, a key site for nutrient absorption and microbial interactions [11, 14, 20]. By altering microbial adherence or competition at this interface, forage bioactives regulate microbial density and diversity, impacting the functional balance within the rumen ecosystem and contributing to overall digestive efficiency.

## **EFFECTS ON MICROBIAL DENSITY**

### **Bioactive Compounds as Growth Promoters**

Forage bioactive compounds can act as prebiotics, stimulating the growth of beneficial microbes like cellulolytic bacteria. These compounds provide substrates or create favorable conditions for these microbes to thrive, enhancing fermentation efficiency and nutrient utilization [28–31]. This promotion supports a healthy microbial balance critical for optimal rumen function and dairy cow performance.

### **Growth Inhibition of Pathogenic Microbes**

Certain bioactive compounds selectively inhibit the growth of pathogenic or competitive microbes. This inhibition can occur through mechanisms such as disruption of microbial membranes, enzyme inhibition, or interference with nutrient uptake [28–31]. By suppressing harmful populations, these compounds help maintain a balanced microbial ecosystem, preventing dysbiosis and enhancing overall rumen health.

### **Changes in Microbial Density**

Bioactive compounds can influence microbial density by altering nutrient availability. For example, tannins may bind dietary proteins, reducing availability for some microbes while benefiting others adapted to utilize bound proteins [28–31]. Conversely, these compounds can release nutrients from indigestible complexes, reshaping microbial densities based on nutrient access.

### **Interactions with Microbial Growth Factors**

Forage bioactives interact with microbial growth factors and inhibitors, modulating microbial density. They may enhance growth by providing essential cofactors or inhibit growth by neutralizing microbial activators [28–31]. These interactions create a dynamic microbial environment, balancing the populations of fermentative microbes to optimize digestion and reduce undesirable byproducts like methane.

## **IMPACT ON MICROBIAL DIVERSITY**

### **Selective Pressure on Microbial Species Richness**

Bioactive compounds exert selective pressure on microbial species richness in the rumen. By favoring certain species over others, they shape the overall species composition, promoting the growth of microbes that are better adapted to the compound's chemical properties [28–31]. This selective pressure can increase or decrease microbial diversity, influencing the stability and resilience of the rumen ecosystem.

### **Role in Shaping Microbial Dominance**

The presence of bioactive compounds can influence microbial evenness and dominance in the rumen. Compounds, like tannins, may reduce the dominance of specific species, promoting a more even distribution of microbial populations [28, 31]. This balanced microbial environment is essential for maintaining stable fermentation processes, enhancing nutrient utilization, and preventing overgrowth of harmful microbes.

### **Promotion of Specific Microbial Guilds**

Forage bioactive compounds can selectively promote specific microbial guilds, such as cellulolytic, amylolytic, or proteolytic microbes, which are essential for digesting various dietary components [28, 31]. By encouraging the growth of these functional groups, bioactives improve rumen fermentation efficiency, fiber degradation, and overall nutrient utilization, contributing to enhanced dairy cow productivity.

### **Effects on Rare Microbial Taxa**

Bioactive compounds can influence rare microbial taxa, which often play critical, yet understudied, ecological roles in the rumen [28, 31]. These compounds may stimulate the growth of less abundant but functionally significant species, thereby enhancing the overall microbial network. The modulation of these rare taxa can improve nutrient cycling, fermentation efficiency, and microbial interactions within the rumen ecosystem.

## **DIRECT ANTIMICROBIAL EFFECTS**

### **Mechanisms of Microbial Cell Wall Disruption**

Bioactive compounds can directly disrupt microbial cell walls, cause leakage of cellular contents, and impair microbial growth [18, 32–35]. Compounds, such as saponins and tannins interact with lipid bilayers or protein structures on microbial membranes, leading to structural damage. This disruption can inhibit microbial colonization and activity, particularly of pathogenic or less beneficial species.

### **Inhibition of Microbial Enzyme Systems**

Many forage bioactive compounds inhibit microbial enzyme systems essential for nutrient degradation in the rumen. For example, tannins can bind enzymes, like cellulases, reducing the breakdown of plant fibers [18, 32–35]. This inhibition affects microbial metabolism, limiting the efficiency of rumen fermentation and altering microbial population dynamics by selectively suppressing certain enzymatic pathways.

### **Disruption of Microbial Metabolic Pathways**

Bioactive compounds can interfere with microbial metabolic pathways by altering nutrient uptake or modulating enzymatic processes [18, 33–35]. For instance, alkaloids can disrupt nitrogen metabolism or interfere with energy production pathways, reducing microbial growth. These effects may limit the proliferation of specific microbial groups, including methanogens or other less desirable species, and promote a more balanced rumen environment.

### **Synergistic Effects with Other Components**

Bioactive compounds often exert synergistic antimicrobial effects when combined with other dietary components, such as fiber or protein. For example, the presence of certain flavonoids may enhance the antimicrobial properties of tannins, improving their ability to target harmful microbes [18, 33–35]. This synergy can lead to more effective modulation of rumen microbiota, improving fermentation efficiency and overall animal health.

## **INDIRECT EFFECTS VIA METABOLITES**

### **Influence on Microbial Metabolite Production**

Forage bioactive compounds indirectly affect microbial metabolite production by modulating microbial activity [36–38]. For instance, compounds, like tannins, may reduce the production of ammonia by inhibiting protein degradation, while promoting the production of volatile fatty acids (VFAs) through enhanced fermentation of fiber. These shifts in metabolite profiles influence rumen health, efficiency, and nutrient absorption.

### **Alteration of the Rumen pH Buffering Capacity**

Bioactive compounds can alter the rumen's pH buffering capacity by affecting the production and absorption of VFAs [36–38]. Some bioactives may promote the accumulation of specific VFAs, thereby lowering rumen pH, which can inhibit the growth of pathogenic microbes while favoring beneficial,

acid-tolerant species. This buffering effect plays a crucial role in maintaining optimal conditions for microbial function.

### **Interaction with Secondary Metabolites**

The interaction between forage bioactive compounds and microbial secondary metabolites can influence the overall microbial ecosystem. Bioactives can modify the production of antimicrobial peptides, organic acids, or other secondary metabolites that either enhance or inhibit the growth of specific microbial species [36–38]. This interaction shapes microbial competition and cooperation, impacting colonization and overall rumen functionality.

### **Feedback Effects of Microbial Metabolites**

The metabolites produced by microbes, such as VFAs and methane, can influence further microbial colonization and population dynamics. High levels of certain VFAs may suppress the growth of less desirable microbes, creating feedback loops that support the growth of beneficial species [36–38]. Additionally, microbial metabolites can alter the local environment of the rumen, influencing future microbial colonization patterns.

## **INTERACTIONS WITH HOST PHYSIOLOGY**

### **Modulation of Rumen Epithelial Secretions**

Forage bioactive compounds can influence the secretion of mucus and other antimicrobial peptides in the rumen epithelium, affecting microbial attachment to the epithelial surface [39–42]. Compounds, like tannins, may enhance the production of mucins, creating a more viscous environment that reduces microbial adhesion or selectively promotes attachment of beneficial species. This interaction helps regulate microbial colonization and stability.

### **Influence on Saliva Production**

Bioactive compounds can alter the production of saliva in dairy cows, which contains antimicrobial components, such as lysozyme and immunoglobulins [39–42]. Increased saliva secretion or changes in its composition may create an environment in the rumen that is less favorable to harmful microbes while supporting the growth of beneficial microbes, thus influencing microbial dynamics and enhancing overall rumen health.

### **Effects on Rumen Motility and Microbial Distribution**

The impact of bioactive compounds on rumen motility can influence microbial distribution within the rumen [39–42]. By modulating rumen contraction patterns, these compounds can alter the flow of ingested material, affecting how microbes are dispersed and where they colonize. This, in turn, affects microbial community structure, fermentation efficiency, and the digestibility of different forage components.

### **Interaction with Immune Responses**

Bioactive compounds can interact with the host's immune system, particularly in the rumen epithelium. These compounds may stimulate immune responses, such as the release of pro-inflammatory cytokines or antimicrobial peptides, which help regulate microbial populations by enhancing the clearance of pathogenic or unproductive microbes [39–42]. This immune modulation contributes to a balanced and healthy rumen microbiota.

## **SYNERGISTIC AND ANTAGONISTIC INTERACTIONS**

### **Synergistic Effects**

Bioactive compounds from different forage sources can act synergistically, enhancing their collective effects on rumen microbiota [43–46]. For example, tannins and saponins may work together to suppress pathogenic microbes while promoting beneficial species like cellulolytic bacteria. These synergistic interactions help to optimize rumen fermentation, improve nutrient digestion, and support a balanced microbial community that benefits dairy cow health and productivity.

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### **Antagonistic Effects Reducing Microbial Diversity**

Some bioactive compounds may have antagonistic effects that reduce microbial diversity in the rumen [43–46]. For example, certain flavonoids or saponins can inhibit the growth of a wide range of microbial species, both beneficial and harmful, leading to a less diverse microbial ecosystem. This reduction in microbial diversity can negatively impact fermentation efficiency and overall rumen health, as balanced microbiota is crucial for optimal digestion and nutrient utilization.

### **Interaction with Other Dietary Components**

Bioactive compounds interact with other dietary components, such as fibers and starch, affecting the rumen microbial ecosystem. For instance, compounds, like tannins, can bind to starch and proteins, altering their availability to microbes [43–46]. This interaction may favor fiber-degrading microbes while reducing the availability of starch to amylolytic bacteria. These shifts in microbial substrates can influence overall microbial activity and nutrient absorption efficiency.

### **Effect on Microbial Cross-Feeding Relationships**

Bioactive compounds can influence microbial cross-feeding relationships within the rumen. Some compounds may enhance the growth of certain microbes that produce metabolites beneficial to other species, fostering a cooperative microbial network [43–46]. Conversely, other bioactives may inhibit these cross-feeding interactions, disrupt microbial collaboration, and reduce fermentation efficiency. Understanding these dynamics is essential for optimizing rumen microbiota and improving nutrient utilization in dairy cows.

## **TEMPORAL DYNAMICS**

### **Short-Term Vs. Long-Term Effects**

Bioactive compounds can have different effects on microbial populations depending on exposure duration. In the short term, these compounds may disrupt microbial growth or alter community composition, but over time, microbial populations may adapt [41, 47]. Long-term exposure may lead to the development of microbial resistance or stabilization of new, more efficient microbial communities that can thrive in the presence of these compounds, ultimately influencing rumen function.

### **Seasonal Variations**

The concentration of bioactive compounds in forages varies seasonally, impacting the rumen microbiota's exposure to these compounds [41, 47]. For instance, certain compounds, like tannins, may be more concentrated in certain plants during dry periods, influencing microbial activity. Seasonal variation in bioactive compound availability can lead to shifts in microbial composition, affecting fermentation patterns and nutrient utilization during different times of the year.

### **Adaptation of Microbial Communities**

Over time, microbial communities in the rumen can adapt to consistent exposure to specific bioactive compounds [41, 47]. For example, microbes that are less sensitive to tannins or other bioactives may become more prevalent, leading to a shift in community composition. These adaptive changes can optimize microbial function under persistent exposure, contributing to improved digestion and overall rumen efficiency.

### **Reversible and Irreversible Microbial Shifts**

Bioactive compounds can cause both reversible and irreversible shifts in microbial populations. In cases of short-term exposure, microbial changes may revert once bioactive compounds are no longer present [41, 47]. However, prolonged exposure can lead to irreversible microbial shifts, such as the establishment of more resilient microbial species or strains, that are better adapted to the bioactive compounds. These long-term shifts can have lasting effects on rumen microbial diversity and functionality.

## **ENVIRONMENTAL FACTORS INFLUENCING EFFECTS**

### **Impact Under Different Feeding Systems**

The effects of forage bioactive compounds on rumen microbiota can vary significantly between grazing and confinement feeding systems [29, 48, 49]. In grazing systems, cows have access to a more

diverse range of plant species with varying bioactive compounds, which can lead to a broader microbial diversity. In contrast, confinement systems may provide a more uniform diet, potentially limiting microbial adaptation and reducing the impact of bioactive compounds on microbial dynamics.

### **Effects of Rumen Temperature and pH Changes**

Rumen temperature and pH fluctuations can influence the effectiveness of bioactive compounds on microbial populations. A decrease in rumen pH, such as during high-concentrate feeding, can enhance the antimicrobial effects of certain bioactives, like tannins, which may thrive in more acidic environments [29, 48, 49]. Additionally, temperature changes, such as during heat stress, may alter microbial metabolism, affecting how bioactive compounds interact with rumen microbes and their subsequent effects on digestion and fermentation.

### **Interaction with the Forage**

Bioactive compounds are present in both water-soluble and structural fractions of forages, with different implications for microbial interactions. Water-soluble compounds, like saponins and flavonoids, are more readily available to microbes and can exert immediate effects on microbial growth and fermentation [29, 48, 49]. Structural compounds, such as tannins and lignins, may be released more slowly during rumen fermentation and can have a more sustained impact on microbial populations, altering microbial growth over time.

### **Influence of External Stressors**

External stressors, such as heat stress or nutritional deficiencies, can influence the impact of bioactive compounds on rumen microbiota. Heat stress can affect rumen motility, microbial distribution, and fermentation, potentially enhancing or diminishing the effects of bioactive compounds [29, 48, 49]. Similarly, nutritional deficiencies may alter the microbial response to bioactives, with certain microbes thriving or being inhibited based on the availability of specific nutrients in the rumen. These factors collectively influence how bioactive compounds interact with the rumen microbiota under diverse environmental conditions.

## **ROLE IN METHANOGENESIS**

### **Inhibition of Methanogenic Archaea**

Bioactive compounds can inhibit the colonization of methanogenic archaea in the rumen. Compounds, such as tannins and saponins, have been shown to disrupt the growth of methanogens by affecting their ability to attach to rumen particles or reducing their metabolic activity [16, 50, 51]. This inhibition can decrease methane production, improve the efficiency of rumen fermentation, and reduce the environmental impact of dairy farming.

### **Effects on Hydrogen-Utilizing Bacteria**

Certain bioactive compounds affect hydrogen-utilizing bacteria, which play a crucial role in the reduction of hydrogen, a precursor for methane production. By modulating the activity of these bacteria, bioactives can influence hydrogen availability for methanogens, reducing methane production [16, 50–53]. For example, flavonoids and tannins may enhance the growth of hydrogen-consuming microbes, thereby indirectly limiting the substrates available for methanogenesis.

### **Changes in Syntrophic Relationships**

Bioactive compounds can alter the syntrophic relationships between methanogens and other rumen microbes, particularly those involved in fermentation processes [54–56]. For instance, bioactives might stimulate the growth of bacteria that compete for hydrogen or other metabolites essential for methanogens, creating an environment where methanogenesis is reduced. These shifts in microbial interactions can lead to a more efficient and sustainable fermentation process, minimizing methane emissions.

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### **Reduction in Methane Emissions**

By influencing microbial populations, bioactive compounds can modify rumen fermentation pathways, leading to a reduction in methane emissions. For example, compounds, like tannins, can shift fermentation toward the production of more beneficial VFAs and away from methane [54–56]. These changes in microbial activity result in more efficient energy use by the cow, potentially improving both animal performance and environmental sustainability.

## **TECHNOLOGICAL APPLICATIONS**

### **Use of Bioactive Compounds in Feed Additives**

Bioactive compounds are increasingly being incorporated into feed additives to modulate the rumen microbiota and enhance digestion efficiency [33, 53, 57, 58]. For instance, tannins and saponins can be included in supplements to selectively inhibit pathogenic microbes while promoting beneficial species. These additives can improve nutrient utilization, reduce methane emissions, and support overall animal health, making them valuable tools in sustainable livestock production.

### **Development of Rumen Microbial Inoculants**

Rumen microbial inoculants, when combined with bioactive compounds, offer a novel strategy for enhancing rumen fermentation [33, 53–57]. These inoculants introduce beneficial microbes that can thrive in the presence of specific bioactives, improving the balance of the rumen microbiota. This synergistic approach could enhance digestion, reduce methane emissions, and improve feed efficiency, contributing to more sustainable and cost-effective dairy farming practices.

### **Bioactive Compounds in Precision Feeding**

Bioactive compounds can be incorporated into precision feeding strategies to optimize their effects on rumen microbiota [21, 42]. By tailoring the inclusion of specific bioactives based on individual animal needs, age, or health status, these strategies can improve microbial efficiency and nutrient absorption. Precision feeding ensures that dairy cows receive the right number of bioactive compounds at the right time, enhancing both animal performance and environmental sustainability.

### **Microbial Resilience Enhancement**

Bioactive compounds can be used to enhance microbial resilience in the rumen, particularly in response to environmental stressors such as heat or dietary changes. These compounds may strengthen the microbial community's ability to withstand challenges, maintaining microbial diversity and functionality [33, 57, 59, 60]. As a result, bioactives could play a key role in ensuring the stability and adaptability of the rumen microbiota, improving long-term productivity and health outcomes for dairy cows.

## **CONCLUSIONS**

Forage bioactive compounds significantly influence rumen microbiota in dairy cows, affecting microbial colonization, density, and diversity. These compounds, including tannins, flavonoids, and saponins, interact with rumen microbes in ways that can either promote beneficial microbes or inhibit harmful ones, improving nutrient utilization and feed efficiency. Modulating microbial diversity is crucial for efficient digestion and reducing methane emissions. Environmental factors, such as rumen pH and temperature, further affect the impact of these bioactives on microbial communities. Understanding these dynamics opens opportunities for using bioactive compounds in feed additives, precision feeding, and microbial inoculants, contributing to improved cow health, productivity, and sustainability. Future research should focus on long-term effects and practical applications for optimizing rumen microbiota and enhancing livestock management.

### **Future Directions**

Future research should focus on further elucidating the molecular mechanisms by which forage bioactive compounds influence rumen microbiota dynamics. This includes identifying specific microbial species targeted by bioactives and understanding how their interactions contribute to

improved feed efficiency and reduced methane emissions. Additionally, studies should explore the long-term effects of bioactive compound supplementation, considering potential adaptations in microbial populations and rumen function. There is also a need to investigate the synergistic and antagonistic effects of bioactive compounds when combined with other dietary components or microbial inoculants. Research into the environmental and management factors that influence bioactive compound effectiveness will be crucial for optimizing their use in various feeding systems. Finally, practical applications in precision feeding strategies and the development of sustainable feed additives should be prioritized to improve both productivity and environmental sustainability in dairy farming.

## REFERENCES

1. Plaizier JC, Mesgaran MD, Derakhshani H, Golder H, Khafipour E, Kleen JL, et al. Review: Enhancing gastrointestinal health in dairy cows. *Anim.* 2018;12(S2):S399–S418. doi: 10.1017/S1751731118001921.
2. Cholewińska P, Czyz K, Nowakowski P, Wrostek A. The microbiome of the digestive system of ruminants – A review. *Anim Health Res Rev.* 2020;21(1):3–14. doi: 10.1017/S1466252319000069.
3. Diao Q, Zhang R, Fu T. Review of strategies to promote rumen development in calves. *Animals.* 2019;9(8):490. doi: 10.3390/ani9080490.
4. Maniaci G, Ponte M, Giosuè C, Gannuscio R, Pipi M, Gaglio R, et al. Cladodes of *Opuntia ficus-indica* (L.) as a source of bioactive compounds in dairy products. 2024; Elsevier. doi: 10.3168/jds.2023-23847.
5. McGaw L. Use of plant-derived extracts and bioactive compound mixtures against multidrug resistant bacteria affecting animal health and production. In: *Fighting Multidrug Resistance with Herbal Extracts, Essential Oils and Their Components.* 2025. pp. 291–311.
6. Mills S, Ross RP, Hill C, Fitzgerald GF, Stanton C. Milk intelligence: Mining milk for bioactive substances associated with human health. *Int Dairy J.* 2011;21(6):377–401. doi: 10.1016/j.idairyj.2010.12.011.
7. Nekrasov RV, Lozovanu MI, Laptev GY, Ilina LA, Yildirim EA, Tyurina DG, et al. Bioactive feed additive for the prevention of clostridial disease in high-yielding dairy cattle. *Agriculture.* 2023;13(4):786. doi: 10.3390/agriculture13040786.
8. Lu Q, Luo Q, Li J, Wang X, Ban C, Qin J, et al. Evaluation of the chemical composition, bioactive substance, gas production, and rumen fermentation parameters of four types of distiller's grains. *Molecules.* 2022;27(18):6134. doi: 10.3390/molecules27186134.
9. Kussmann M, Abe Cunha DH, Berciano S. Bioactive compounds for human and planetary health. *Front Nutr.* 2023;10:1193848. doi: 10.3389/fnut.2023.1193848.
10. Park YW. *Bioactive components in milk and dairy products.* Hoboken (NJ): John Wiley & Sons; 2009. p. 1–426. doi: 10.1002/9780813821504.
11. Arshad MA, ul Hassan F, Rehman MS, Huws SA, Cheng Y, Din AU. Gut microbiome colonization and development in neonatal ruminants: Strategies, prospects, and opportunities. *Anim Nutr.* 2021;7(3):883–895. doi: 10.1016/j.aninu.2021.03.004.
12. Amat S, Dahlen CR, Swanson KC, Ward AK, Reynolds LP, Caton JS. Bovine animal model for studying the maternal microbiome, in utero microbial colonization and their role in offspring development and fetal programming. *Front Microbiol.* 2022;13:854453. doi: 10.3389/fmicb.2022.854453.
13. Wang Y-L, Wang W-K, Wu Q-C, Zhang F, Li W-J, Li S-L, et al. In situ rumen degradation characteristics and bacterial colonization of corn silages differing in ferulic and p-coumaric acid contents. *Microorganisms.* 2022;10(11):2269. doi: 10.3390/microorganisms10112269.
14. Anadón A, Ares I, Martínez-Larrañaga MR, Martínez MA. Prebiotics and probiotics in feed and animal health. In: *Nutraceuticals in Veterinary Medicine.* 2019. pp. 261–285. doi: 10.1007/978-3-030-04624-8\_19.
15. Rabee AE, El Rahman TA, Lamara M. Changes in the bacterial community colonizing extracted and non-extracted tannin-rich plants in the rumen of dromedary camels. *PLoS One.* 2023;18(2):e0282889. doi: 10.1371/journal.pone.0282889.

16. Angelova T, Krastanov J, Yordanova D, et al. Inhibitors of methanogenesis. *Bulg J Vet Med*. 2023. Available at <https://search.ebscohost.com/login.aspx?direct=true&profile=ehost&scope=site&authtype=crawler&jrnl=05147441&AN=164640279>.
17. Duarte VdS, Franklin FV, Krysmann A, Porcellato D. Longitudinal study of the udder microbiome of Norwegian Red dairy cows using metataxonomic and shotgun metagenomic approaches: Insights into pathogen-driven dynamics. *bioRxiv*. 2025. doi: 10.1101/2025.04.04.647183.
18. Ike KA, Adelusi OO, Alabi JO, et al. Onion peel and essential oils as feed additives in the diet of dairy cattle: In vitro fermentation, greenhouse gases, ruminal nutrient degradability, and volatile fatty acids. *Turk J Vet Anim Sci*. 2024. Available at <https://journals.tubitak.gov.tr/veterinary/vol48/iss6/3/>.
19. Hartonian P. The effects of rumen modifiers on the rumen and hindgut microbiota composition of lactating dairy cows. ProQuest Dissertations. 2024:31239043. Available at: <https://search.proquest.com/openview/8097d4856380b0035a45a4f4a848afdd/1?pq-origsite=gscholar&cbl=18750&diss=y>.
20. Aditya A, Rahaman SO, Biswas D. Impact of *Lactobacillus*-originated metabolites on enterohemorrhagic *E. coli* in rumen fluid. *FEMS Microbiol Ecol*. 2022;98(12):fiac128. doi: 10.1093/femsec/fiac128.
21. Ma J, Huangfu W, Yang X, Xu J, Zhang Y, Wang Z, et al. 'King of the forage'—alfalfa supplementation improves growth, reproductive performance, health condition and meat quality of pigs. *Front Vet Sci*. 2022;9:1025942. doi: 10.3389/fvets.2022.1025942.
22. Khiaosa-Ard R, Pacífico C, Mahmood M, Mickdam E, Meixner J, Trautinger L-S, et al. Changes in the solid-associated bacterial and fungal communities following ruminal in vitro fermentation of winery by-products: Aspects of the bioactive compounds and feed safety. *Anaerobe*. 2024;82:102893. doi: 10.1016/j.anaerobe.2024.102893.
23. Park YW, Haenlein GF. Milk and dairy products in human nutrition: Production, composition and health. Hoboken (NJ): John Wiley & Sons; 2013. doi: 10.1002/9781118534168.
24. Silva FG, Silva SR, Pereira AMF, Cerqueira JL, Conceição C. A comprehensive review of bovine colostrum components and selected aspects regarding their impact on neonatal calf physiology. *Animals*. 2024;14(7):1130. doi: 10.3390/ani14071130.
25. Patra AK, Saxena J. Dietary phytochemicals as rumen modifiers: A review of the effects on microbial populations. *Antonie Van Leeuwenhoek*. 2009;96:363–375. doi: 10.1007/s10482-009-9364-1.
26. Agulló V, Favari C, Pilla N, Bresciani L, Tomás-Barberán FA, Crozier A, et al. Using targeted metabolomics to unravel phenolic metabolites of plant origin in animal milk. *Int J Mol Sci*. 2024;25(8):4536. doi: 10.3390/ijms25084536.
27. del C Mondragon Portocarrero A, López-García M, Pérez-Sánchez A, et al. Substitutive effects of milk vs. vegetable milk on the human gut microbiota and implications for human health. *Nutrients*. 2024;16(18):3108. doi: 10.3390/nu16183108.
28. Guo J, Zhang Z, Guan LL, Zhou M, Yoon I, Khafipour E, et al. Postbiotics from *Saccharomyces cerevisiae* fermentation stabilize rumen solids microbiota and promote microbial network interactions and diversity of hub taxa during grain-based subacute ruminal acidosis (SARA) challenges in lactating dairy cows. *Front Microbiol*. 2024;15:1409659. doi: 10.3389/fmicb.2024.1409659.
29. Firkins JL. Reconsidering rumen microbial consortia to enhance feed efficiency and reduce environmental impact of ruminant livestock production systems. *SciELO Brasil*. 2010;13:49. doi: 10.1590/s1516-35982010001300049.
30. Cammack KM, Austin KJ, Lamberson WR, Conant GC, Cunningham HC. Ruminant nutrition symposium: Tiny but mighty: The role of the rumen microbes in livestock production. *J Anim Sci*. 2018;96(2):752–770. doi: 10.1093/jas/skx053.
31. Tedeschi LO, Muir JP, Naumann HD, Norris AB, Ramírez-Restrepo CA, Mertens-Talcott SU. Nutritional aspects of ecologically relevant phytochemicals in ruminant production. *Front Vet Sci*. 2021;8:628445. doi: 10.3389/fvets.2021.628445.
32. Oyama LB. Prospecting rumen bacteria for novel antimicrobials. 2015. Available at: <https://core.ac.uk/download/pdf/185314933.pdf>.

33. Adetunji AO, Price J, Owusu H, Adewale EF, Adesina PA, Saliu TP, et al. Mechanisms by which phytogetic extracts enhance livestock reproductive health: Current insights and future directions. *Front Vet Sci.* 2025;12:1568577. doi: 10.3389/fvets.2025.1568577.
34. Kwoji ID, Aiyegoro OA, Okpeku M, Adeleke MA. Multi-strain probiotics: synergy among isolates enhances biological activities. *Biology (Basel).* 2021;10(4):322. doi: 10.3390/biology10040322.
35. Gomes V, Hoffmann C, Castro-Tardón DI, Ramos dos Santos FC, Suárez-Retamozo S, Hurley DJ. Vertical transfer of gut microbiota from dam to neonate calf in the early life. *Sci Rep.* 2024;14:72296. doi: 10.1038/s41598-024-72296-0.
36. Taschuk R, Griebel PJ. Commensal microbiome effects on mucosal immune system development in the ruminant gastrointestinal tract. *Anim Health Res Rev.* 2012;13(1):129–141. doi: 10.1017/S1466252312000096.
37. Szumacher-Strabel M, Cielak A. Dietary possibilities to mitigate rumen methane and ammonia production. *Greenh Gases – Capturing, Util Reduct.* 2012. doi: 10.5772/32105.
38. Christensen RG. Improvement of nutrient utilization efficiency, ruminal fermentation and lactational performance of dairy cows by feeding birdsfoot trefoil. 2015. Available at: <https://search.proquest.com/openview/38db5af54e625c1cf46392a4d60b5656/1?pq-origsite=gscholar&cbl=18750>.
39. Zhang J, Shi H, Wang Y, Li S, Cao Z, Ji S, et al. Effect of dietary forage to concentrate ratios on dynamic profile changes and interactions of ruminal microbiota and metabolites in Holstein heifers. *Front Microbiol.* 2017;8:2206. doi: 10.3389/fmicb.2017.02206.
40. Wang F, Sha Y, He Y, Liu X, Chen X, Yang W, et al. Age differences in ileum microbiota density: VFAs and their transport-related gene interactions in Tibetan sheep. *Fermentation.* 2024;10(10):509. doi: 10.3390/fermentation10100509.
41. Law SR, Mathes F, Paten AM, Alexandre PA, Regni R, Reid C, et al. Life at the borderlands: Microbiomes of interfaces critical to One Health. *FEMS Microbiol Rev.* 2024;48(2):fuae008. doi: 10.1093/femsre/fuae008.
42. Kaur H, Kaur G, Gupta T, Mittal D, Ali SA. Integrating omics technologies for a comprehensive understanding of the microbiome and its impact on cattle production. *Biology (Basel).* 2023;12(9):1200. doi: 10.3390/biology12091200.
43. Szumacher-Strabel M, Cieślak A. Essential oils and rumen microbial populations. In: *Diet Phytochem Microbes.* 2012. pp. 285–309. doi: 10.1007/978-94-007-3926-0\_10.
44. Callaway TR, Lillehoj H, Chuanchuen R, Gay CG. Erratum: Callaway TR, Lillehoj H, Chuanchuen R, Gay CG. Alternatives to antibiotics: A symposium on the challenges and solutions for animal health and production. *Antibiotics (Basel).* 2021;10:81024. doi: 10.3390/antibiotics10081024.
45. Ike KA, Okedoyin DO, Alabi JO, Adelusi OO, Wuaku M, Olagunju LK, et al. The combined effect of four nutraceutical-based feed additives on the rumen microbiome, methane gas emission, volatile fatty acids, and dry matter disappearance using an in vitro batch culture technique. *Fermentation.* 2024;10(10):499. doi: 10.3390/fermentation10100499.
46. Firkins JL, Yu Z. Ruminant nutrition symposium: How to use data on the rumen microbiome to improve our understanding of ruminant nutrition. *J Anim Sci.* 2015;93(4):1450–1470. doi: 10.2527/jas.2014-8754.
47. Adnane M, Chapwanya A. Microbial gatekeepers of fertility in the female reproductive microbiome of cattle. *Int J Mol Sci.* 2024;25(20):10923. doi: 10.3390/ijms252010923.
48. Cholewińska P, Górnjak W, Wojnarowski K. Impact of selected environmental factors on microbiome of the digestive tract of ruminants. *BMC Vet Res.* 2021;17:xxx. doi: 10.1186/s12917-021-02742-y.
49. Ban Y, Neves LA, Guan LL, McAllister T. Modifying the rumen environment to reduce greenhouse gas emissions. In: *Reducing Greenhouse Gas Emissions from Livestock Production.* 2022. pp. 287–330. doi: 10.1201/9781003048213-11.
50. Angelova T, Krastanov J, Yordanova D, Mihaylova M. Inhibitors of methanogenesis. *Review.* 2023. doi: 10.5555/20230293722.

51. Ebeid HM, Mengwei L, Kholif AE, Hassan F-U, Lijuan P, Xin L, et al. Moringa Oleifera oil modulates rumen microflora to mediate in vitro fermentation kinetics and methanogenesis in total mixed rations. *Curr Microbiol.* 2020;77:1271–1282. doi: 10.1007/s00284-020-01935-2.
52. Cieslak A, Szumacher-Strabel M, Stochmal A, Oleszek W. Plant components with specific activities against rumen methanogens. *Animal.* 2013;7(Suppl 2):253–265. doi: 10.1017/S1751731113000852.
53. Zeng X, Chen Y, Li W, Liu S. Application of fenugreek in ruminant feed: Implications for methane emissions and productivity. *PeerJ.* 2024;12:e16842. doi: 10.7717/peerj.16842.
54. Hosen Z, Islam MR, Naidu R, Biswas B. ‘Geophagy’ and clay minerals: Influencing ruminal microbial fermentation for methane mitigation. *Microorganisms.* 2025;13:866. doi: 10.3390/microorganisms13040866.
55. Wei H, Yang D, Chen Y, Yang G. Rumen fermentation and microbial profiles of sheep fed a high-concentrate diet supplemented with ellagic acid. *J Anim Sci Vet Med.* 2023;1:0355. doi: 10.11829/j.issn.1001-0629.2022-0355.
56. Yáñez-Ruiz DR, Belanche A. Plant secondary compounds: Beneficial roles in sustainable ruminant nutrition and productivity. In: *Improving Rumen Function.* 2020. p. 727–774. doi: 10.19103/as.2020.0067.25.
57. Jack A, Fawole O, Chen J. Microalgae application in feed for ruminants. In: *Handbook of Food and Feed from Microalgae: Production, Applications, Regulation, Sustainability.* 2023. p. 397–409. doi: 10.1016/B978-0-323-99196-4.00042-5.
58. Attwood GT, Reilly K, Forster RJ. Applications of the soil, plant and rumen microbiomes in pastoral agriculture. *Front Nutr.* 2019;6:107. doi: 10.3389/fnut.2019.00107.
59. Gao Z, Liu B, La S, Li D, Zhu X, Sun H, et al. Alfalfa hay substitution for wheat straw improves beef quality via rumen microflora alteration. *Heliyon.* 2023;9:e20803. doi: 10.1016/j.heliyon.2023.e20803.
60. Uyanga VA, Ejeromedoghene O, Lambo MT, Alowakennu M, Alli YA, Ere-Richard AA, et al. Chitosan and chitosan-based composites as beneficial compounds for animal health: Impact on gastrointestinal functions and biocarrier application. *J Funct Foods.* 2023;109:105520. doi: 10.1016/j.jff.2023.105520.