

Assessment of molasses-based additives for methane mitigation in beef cattle

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I certify that the intellectual content of this thesis is the product of my own work and that all the assistance received in preparing this thesis and sources have been acknowledged.

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Abstract

Enteric methane (CH₄) mitigation is a priority for pasture-based beef systems, where practical delivery of antimethanogenic additives to grazing cattle remains a key constraint. In Australia, enteric methane contributes a substantial share of the agricultural emissions profile, underscoring the need for scalable solutions suited to extensive systems. Molasses lick blocks (MLB) have been proposed as a feasible vehicle to regulate intake and enable more consistent dosing in grazing ruminants.

This thesis aimed to evaluate novel molasses-based additive formulations intended for incorporation into molasses lick blocks (MLBs) by (i) screening their *in vitro* antimethanogenic effects and (ii) quantifying *in vivo* responses in growing beef cattle over a 70-day feeding period. Specific ingredient identities are not disclosed due to commercial confidentiality and pending intellectual property protection; therefore, emphasis is placed on evaluating the performance of the formulated products rather than their proprietary components.

In vitro, four candidate formulations were assessed for their effects on rumen fermentation and methane production. One formulation (“Product 3”) displayed the strongest response. At 6% inclusion, CH₄ production was reduced by >90%, although this was accompanied by reduced *in vitro* dry matter digestibility. Additional testing showed that at 4% inclusion, Product 3 reduced CH₄ production by 62% without impairing digestibility, highlighting the importance of dose optimisation.

The *in vivo* experiment evaluated Product 3 delivered as a grain-based pellet to growing steers over 70 days, with target intakes of 0, 100, or 200 g per head per day. Methane emissions were measured using a GreenFeed system. Compared with steers receiving pellets without the product, CH₄ production tended to decrease by up to 10.7% ($P = 0.07$), CH₄ yield (g CH₄/kg DMI) was reduced ($P < 0.01$) by 11.7% and 7.5% at 100 and 200 g/day, respectively, and CH₄ intensity tended ($P = 0.06$) to decline by 16.8%. No detrimental effects were detected on dry matter intake, liveweight gain, or feed efficiency. Product intake was negatively correlated with CH₄ yield and intensity and positively correlated with water intake and average daily gain ($P < 0.05$), indicating acceptable palatability and compatibility with forage-based diets.

Collectively, these findings demonstrate a clear dose-sensitive antimethanogenic effect for Product 3, with >90% inhibition observed *in vitro* and approximately a 12% reduction in methane emissions *in vivo*, achieved without compromising animal performance. Several factors, however, limit direct extrapolation of these results to grazing systems. These include the additive stability and the structured feeding environment imposed by the experimental design.

In conclusion, Product 3 reduced methane yield by up to 11.7% *in vivo* over 56 days without adverse effects on intake or growth and achieved >90% inhibition *in vitro* at high inclusion, with substantial reductions at lower doses that preserved fermentation. These findings identify a promising dietary strategy for low-emission beef production while emphasising the need for optimisation of inclusion rates, assessment of response persistence, and confirmation of performance using final MLBs in grazing herds.

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Author Attribution Statement

I certify that this thesis is the result of my own work. I was primarily responsible for the research ideas, organisation, data analysis, and writing of all chapters and publications included in this thesis. The work was conducted independently under the supervision of Professor A. V. Chaves, Professor L. A. González, and Dr G. I. Simanungkalit.

Chapter 2 has been published in a peer-reviewed journal (Frontiers in Animal Science), and the published version is included in this thesis. All contributions by others to the research and preparation of this thesis have been appropriately acknowledged, either through authorship on the publication or in the acknowledgements section.

I certify that the intellectual content of this thesis is original and, to the best of my knowledge and belief, contains no material previously submitted, in whole or in part, for the award of any other degree or diploma at this or any other tertiary institution.

Maria Nikoloric

As supervisor for the candidature upon which this thesis is based, I can confirm that the authorship attribution statements above are correct.

Professor Chaves, A. V.

Generative AI Statement

During the preparation of the thesis, the author used ChatGPT to assist in the understanding of data analysis methods and Microsoft Copilot to support improvements in grammar, spelling, and overall readability, as well as to enhance the clarity and quality of the manuscript redaction. The author confirms that no text was directly generated or modified by generative AI during the preparation of the thesis. The author takes full responsibility for the submitted thesis, ensures the work is their own, and has used generative AI in accordance with university guidelines and policies (refer to the University of Sydney generative AI guide for researchers).

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Abbreviations

The following abbreviations are used throughout this thesis and have been defined at first use in each chapter.

A

ADF – Acid Detergent Fibre

ADG – Average Daily Gain

A:P – Acetate-to-propionate ratio

B

BCVFA – Branched-chain volatile fatty acids

Bovaer[®] – Commercial formulation of 3-NOP

C

CH₄ – Methane

CP – Crude Protein

CO₂ – Carbon Dioxide

CT – Condensed Tannins

D

DM – Dry Matter

DMD – Dry Matter Digestibility

DMI – Dry Matter Intake

DHA – Docosahexaenoic Acid

DFM – Direct-Fed Microbials

E

EID – Electronic Identification Number

EFSA – European Food Safety Authority

EO – Essential Oils

EPA – Eicosapentaenoic Acid

e.g. – For example

F

FAO – Food and Agriculture Organisation

F:C ratio – Forage-to-Concentrate Ratio

G

GC–MS – Gas Chromatography–Mass Spectrometry

GHG – Greenhouse Gases

GreenFeed® – Automated gas measurement system

H

H₂ – Hydrogen

HT – Hydrolysable Tannins

HP5-MS – GC column type

I

IVDMD – In vitro dry matter digestibility

In vivo – Experiments performed in live animals

In vitro – Experiments performed outside the organism

Intergado® – Electronic feeding system

L

LW – Live Weight

M

MCFA – Medium-Chain Fatty Acids

MCR – Methyl-coenzyme M Reductase

MLB – Molasses Lick Blocks

MPhil – Master of Philosophy

MRV – Measurement, Reporting and Verification

O

OM – Organic Matter

OMI – Organic Matter Intake

O₂ – Oxygen

P

P – Statistical significance (probability value)

PUFA – Polyunsaturated Fatty Acids

PTV – Programmed Temperature Vaporisation inlet

R

RFID – Radio-frequency identification

Rusitec – Rumen Simulation Technique

S

SRB – Sulphate-Reducing Bacteria

T

TCA cycle – Tricarboxylic Acid Cycle

TDU – Thermal Desorption Unit

Tenax TA – Adsorbent polymer for GC-MS

3-NOP – 3-Nitrooxypropanol

V

VFA – Volatile Fatty Acids

W

WSC – Water-Soluble Carbohydrates

Introduction

Enteric methane emitted by ruminant livestock is a major contributor to agricultural greenhouse gas emissions and remains a significant constraint for climate-smart livestock production (Beauchemin et al., 2022). In countries with large grazing-based cattle industries, such as Australia, enteric methane accounts for about 10% of national agricultural emissions (Greenwood et al., 2018) and represents an energy loss to the animal (Johnson and Johnson, 1995; Morgavi et al., 2010). Because methane is produced by ruminal anaerobic fermentation, particularly via methanogenic archaea, effective mitigation strategies must reduce emissions without disrupting rumen function, animal performance, or welfare (Hristov et al., 2022).

Nutritional manipulation has emerged as one of the most practical approaches for methane mitigation because dietary interventions can directly influence rumen fermentation pathways without requiring major changes in infrastructure, genetics, or management (Hristov et al., 2013). A wide range of dietary compounds, including synthetic inhibitors (e.g., 3-nitrooxypropanol), lipids, plant secondary metabolites, probiotics, enzymes and electron acceptors have been investigated for their capacity to modify fermentation and reduce methanogenesis (Beauchemin et al., 2022). Among these, plant-derived compounds such as essential oils, tannins, and saponins have attracted significant interest due to their natural origin, potential safety, and consumer acceptance (Patra and Saxena, 2011). These compounds can inhibit methanogens, modify protozoa, and redirect hydrogen towards alternative metabolic sinks (Benchaar and Greathead, 2011). Essential oils (EO) in particular, have shown potential to alter rumen fermentation and improve feed efficiency (Calsamiglia et al., 2007), and the combinations of EO may also increase microbial efficacy due to additive or synergistic effects between EO components (Benchaar and Greathead, 2011). Commercial essential-oil-based feed additives such as Crina[®] and Agolin[®] Ruminant have been developed to provide standardised blends of bioactive plant compounds for livestock. These products have demonstrated modest methane-mitigating effects —typically in the range of 5–10%—in controlled feeding trials without compromising dry matter intake or animal performance (Belanche et al., 2020). However, translating these effects into consistent *in vivo* methane reductions remains challenging, as responses depend on diet composition, additive dose, the length of supplementation and the delivery method (Benchaar and Greathead, 2011), reflecting the broader inconsistency often observed in responses to plant secondary metabolites (Calsamiglia et al., 2007; Jayanegara et al., 2012).

One of the greatest challenges for methane mitigation in extensive grazing systems is the reliable and consistent delivery of feed additives (Beauchemin et al., 2022). Unlike feedlot systems, grazing cattle cannot be individually dosed, making controlled intake difficult to achieve. As a result, additives must be incorporated into supplements such as pellets, molasses mixes, or lick blocks (Hegarty et al., 2021). However, relatively few studies have evaluated the methane-reduction potential of bioactive compounds delivered through such practical feeding systems. Molasses lick blocks, in particular, have been proposed as a promising vehicle for additive delivery because they can regulate intake and provide a more consistent supplementation pattern under grazing conditions (Hristov et al., 2025).

This thesis proposes that incorporating methane-inhibiting additives into molasses lick blocks could provide a practical mitigation strategy for pasture-based beef systems. The primary objective was to assess the formulation of novel MLB-based products designed to improve cattle productivity while reducing enteric CH₄ emissions. These products contained known anti-methanogenic compounds; however, specific ingredients cannot be disclosed due to commercial confidentiality and patent restrictions. Therefore, the emphasis is placed on the performance of the formulated product rather

than individual compounds. Importantly, this study evaluated the MLB-based formulations themselves—not hardened commercial lick blocks— which require further validation under grazing conditions.

Chapter 1 presents a literature review of safe dietary compounds with demonstrated or potential methane mitigation effects in ruminants. It examines the mechanisms by which these compounds influence rumen fermentation, hydrogen dynamics, methanogenic archaea, and overall fermentation pathways. Particular attention is given to the limitations and practical challenges associated with applying dietary strategies in grazing systems, including difficulty to measure intake, palatability concerns, and challenges in delivering consistent doses to free-ranging cattle. The literature review also identifies MLB technology as a promising vehicle for targeted additive delivery.

Chapter 2 evaluates four novel molasses-based additives formulations in vitro to assess their effects on fermentation and CH₄ production. Based on this result, the most effective formulation (Product 3) was subsequently incorporated into grain-based pellets and tested in vivo with growing steers. Results demonstrated reductions in methane yield and tendencies for lower CH₄ production and intensity, without adverse effects on dry matter intake or growth performance. These findings highlight Product 3 as a promising strategy; however, high inclusion levels posed potential risks for rumen fermentation, underscoring the need for dose optimisation. Chapter 3 provides a general discussion and conclusion that synthesises insights from both the literature and the experimental findings. It also highlights the limitations of the study and suggests possible directions for future research, along with the main conclusion of the thesis.

This thesis contributes to the growing body of work exploring nutritional strategies for low-emission beef production. Combining laboratory screening with controlled in vivo experimentation provides insight into the dose-dependent responses, practical constraints, and potential applications of molasses-based additive formulations. The finding also highlights areas requiring further investigation to explore higher dosages and extended feeding periods to better understand the trade-off between methane abatement, digestibility and overall productivity. It is also important to note that the products assessed were the formulations intended for manufacturing MLB, rather than the final commercial blocks. Therefore, the final blocks must be tested under grazing conditions to ensure optimal intake, followed by evaluating their effects on growth performance and methane emissions in practical production systems.

Chapter 1: Literature review on safe dietary compounds for mitigating enteric methane emissions in ruminants: mechanisms, efficacy, and future perspectives

1.1. Introduction

The search for safe, practical, and scalable strategies to mitigate enteric methane (CH₄) emissions has increasingly focused on dietary compounds that can be incorporated into conventional feeding systems. Among these, commercial feed additives have gained substantial attention because they can be integrated into existing production practices and provide measurable, reproducible reductions in CH₄ output (Beauchemin et al., 2022; Hristov et al., 2022). Prominent examples include the synthetic inhibitor 3-nitrooxypropanol (3-NOP; marketed as Bovaer®), which directly inhibits the methyl-coenzyme M reductase enzyme in methanogenic archaea (Hristov et al., 2022) and essential oil blends such as *Agolin*® and *Crina*®, which reduce the abundance of hydrogen-producing bacteria

and protozoa that are symbiotically associated with methanogenic archaea (Benchaar et al., 2008). Other promising approaches include electron acceptors and alternative hydrogen sinks, lipid supplementation, and probiotic or enzyme formulations that redirect hydrogen metabolism toward alternative sinks such as propionate formation (Ku-Vera et al., 2020; Roques et al., 2024).

Although many of these additives have demonstrated methane-reducing potential, their efficacy and safety vary considerably depending on ruminant species, diet composition, management system, and environmental conditions (Hegarty et al., 2021; Miller et al., 2023). Furthermore, microbial adaptation and the long-term persistence of CH₄ mitigation remain key uncertainties that may affect durability of response (Beauchemin et al., 2022; Hristov et al., 2022). These challenges underscore the need for a comprehensive synthesis of recent evidence to determine which dietary compounds offer the most reliable and sustainable mitigation potential under commercial conditions.

This review provides an integrated assessment of safe dietary compounds for mitigating enteric CH₄ emissions in ruminants. Safety is defined based on the absence of adverse effects on animal health, feed intake, nutrient digestibility, rumen fermentation characteristics, and overall animal performance, as well as minimal negative impacts on the environment. Emphasis is placed on additives with demonstrated or emerging commercial applicability, comparing their mechanisms of action, efficacy, dose–response relationships, potential side effects on the animal and the environment, and regulatory or economic considerations (see Table 1). The review also highlights critical knowledge gaps and future research priorities to support the development of integrated, science-based, and sustainable strategies for methane abatement in diverse livestock systems.

1.2. Methanogenesis in the rumen

1.2.1. Rumen microbial ecology

The rumen is a specialised anaerobic fermentation chamber that hosts a complex and dynamic microbial ecosystem vital for digesting fibrous plant materials in ruminants. This ecosystem mainly consists of bacteria, archaea, protozoa, fungi, and viruses, which work together to break down structural carbohydrates such as cellulose and hemicellulose, as well as starch, proteins, and lipids (Morgavi et al., 2010; Huws et al., 2018). The fermentation process produces volatile fatty acids (VFA)—primary acetate, propionate, and butyrate—which are the major energy source for the host animal. It also generates gaseous by-products, including carbon dioxide (CO₂), methane (CH₄), and hydrogen (H₂). Under typical rumen conditions, CO₂ accounts for the majority of the gas produced (approximately 60–70%), followed by CH₄ (around 25–35%), while H₂ is present only in small amounts due to its rapid utilisation by methanogenic archaea (Hook et al., 2010).

Rumen bacteria group, represents nearly 50-75% of the total microbial population (Minato et al., 1966). They play key roles in carbohydrate degradation, proteolysis, and lipid metabolism. Among them, fibrolytic species such as *Ruminococcus albus*, *R. flavefaciens*, and *Fibrobacter succinogenes* are responsible for the enzymatic hydrolysis of plant cell wall components, releasing sugars that are subsequently fermented to VFAs and gases. Other groups, such as amylolytic (*Streptococcus bovis*, *Selenomonas ruminantium*) and proteolytic bacteria (*Prevotella* spp.), specialise in starch and protein degradation, respectively.

Ciliate protozoa contribute significantly to feed particle breakdown and starch storage, but are also associated with the production of hydrogen and form close symbiotic relationships with methanogens (Finlay et al., 1994). Anaerobic fungi aid in fibre digestion by penetrating plant cell walls, facilitating bacterial access to structural polysaccharides (Gruninger et al., 2014). Bacteriophages and viruses regulate microbial populations through lytic and lysogenic interactions, influencing community stability and nutrient turnover (Anderson et al., 2017).

During fermentation, the production of acetate and butyrate by cellulolytic and saccharolytic

by increasing the redox potential; therefore, efficient disposal of H₂ is critical for maintaining rumen fermentation balance. This disposal occurs through interspecies hydrogen transfer, where methanogenic archaea use H₂ to reduce CO₂ to CH₄ (Morgavi et al., 2010). Other microbes, including fumarate- or nitrate-reducing bacteria and propionate-producing species, can also serve as alternative hydrogen sinks, though methanogenesis is typically the dominant pathway under normal rumen conditions.

Methanogenic archaea and their roles

Methanogenic archaea represent approximately 1–4% of the rumen microbial population but play a pivotal role in maintaining anaerobic stability by consuming excess H₂. The predominant methanogens belong to the genera *Methanobrevibacter*, *Methanomicrobium*, *Methanosphaera*, and *Methanobacterium*, all of which utilise the CO₂ reduction pathway with H₂ as the primary electron donor (Janssen and Kirs, 2008; Hook et al., 2010). This pathway contributes to approximately 90% of total CH₄ formation, although minor contributions arise from the methylotrophic pathway, where methanol or methylamines serve as substrates (Poulsen et al., 2013). *Methanosphaera stadtmanae*, for instance, utilises methanol derived from pectin fermentation, producing CH₄ via the reaction.

Methanogens are closely associated with protozoa and fungal hyphae, often attaching to their surfaces or residing in symbiosomes, enabling efficient H₂ transfer (Finlay et al., 1994). These associations explain why defaunation (the removal of protozoa) can reduce CH₄ production in some systems, as it limits local H₂ availability to methanogens (Newbold et al., 2015).

Molecular studies using 16S rRNA sequencing and metagenomics have shown that *Methanobrevibacter ruminantium* and *Methanobrevibacter gottschalkii* clades dominate in cattle and sheep (Henderson et al., 2015). However, community composition can vary with diet, animal species, and geographical region, influencing CH₄ emission potential. For example, high-concentrate diets generally reduce total methanogen abundance due to lower rumen pH and greater propionate formation (Tapio et al., 2018).

Microbial interactions and methane yield

Methanogenesis is intricately linked to the fermentation pattern of the rumen. When acetate and butyrate are the dominant VFAs produced, more reducing equivalents (H₂) are generated, favouring CH₄ formation. In contrast, propionate production acts as a competitive hydrogen sink, consuming H₂ and thereby reducing CH₄ emissions (Ungerfeld, 2020). Therefore, dietary interventions that shift fermentation towards propionate generally achieve partial CH₄ mitigation.

The total CH₄ yield (g CH₄/kg dry matter intake) is thus a reflection of microbial community composition, substrate availability, and hydrogen flow dynamics. Maintaining a stable, functional microbiota is essential for animal performance; hence, any CH₄ mitigation strategy must balance the reduction of methanogenic activity with preservation of overall fermentation efficiency and nutrient digestibility (Hristov et al., 2013).

1.2.2. Biochemical pathways of methane production

Methanogenesis in the rumen is primarily driven by hydrogenotrophic archaea, which consume molecular H₂ generated during microbial fermentation of carbohydrates to reduce carbon dioxide CO₂ into methane CH₄, thereby maintaining low H₂ partial pressures and enabling efficient rumen fermentation (Hook et al., 2010; Ungerfeld, 2020). Alternative hydrogen sinks also contribute to ruminal H₂ metabolism, including reductive acetogenesis, propionate synthesis, and the reduction of nitrate or sulphate, which can divert H₂ from methanogenesis and reduce CH₄ emissions under specific dietary or microbial conditions (van Zijderveld et al., 2011). Emerging evidence indicates that

interventions targeting these alternative hydrogen sinks—through dietary supplementation, feed additives, or microbial modulation—can effectively mitigate enteric methane emissions while maintaining or even enhancing fermentation efficiency and animal productivity (Beauchemin et al., 2022).

1.2.3. Factors influencing methane production

Diet composition

The composition and quality of the diet are among the most influential factors determining enteric CH₄ emissions in ruminants. Diets rich in structural carbohydrates, particularly cellulose and hemicellulose from forages, stimulate the production of acetate and butyrate during ruminal fermentation—volatile fatty acids that generate metabolic hydrogen as a by-product (Beauchemin et al., 2020). This H₂ serves as a key substrate for methanogenic archaea, facilitating methane synthesis via the hydrogenotrophic pathway. Consequently, high-forage diets typically lead to elevated methane emissions due to the dominance of fibre-degrading bacteria and acetogenic fermentation routes (Knapp et al., 2014).

In contrast, diets with a greater proportion of concentrates (starch-based feeds) tend to reduce methane output by enhancing propionate formation—an alternative hydrogen sink that competes directly with methanogens for available H₂ (Hristov et al., 2013; Kumar et al., 2014). High-concentrate feeding can lower ruminal pH, suppressing the activity of methanogenic archaea and protozoa while favouring amylolytic bacterial populations (Kumar et al., 2014). Also, excessive concentrate inclusion can increase the risk of ruminal acidosis and compromise animal health, emphasising the need for balanced diet formulations (Hristov et al., 2022).

Beyond basal diet manipulation, supplementation with targeted feed additives—such as plant secondary metabolites, lipids, or synthetic inhibitors—has demonstrated substantial mitigation potential. These compounds can inhibit methanogenesis directly or alter fermentation patterns to divert H₂ flow away from methane formation (Beauchemin et al., 2020). Thus, precision diet formulation integrating optimised nutrient composition and additive use represents a key systems-level strategy for sustainable methane abatement.

Animal species and feeding systems

Feeding systems further modulate methane output by influencing feed availability, rumen fermentation dynamics, and animal behaviour. Total mixed rations, which provide a consistent mixture of forages and concentrates, tend to promote stable rumen conditions and can reduce methane yield compared to systems where forages and concentrates are offered separately (Beauchemin et al., 2020). In contrast, grazing systems exhibit greater variability in methane emissions due to differences in forage maturity, nutrient composition, and seasonal feed availability. High-quality pastures with elevated digestibility can improve energy utilisation and reduce emissions per unit of product (Charmley et al., 2015). Pasture management strategies such as rotational grazing, maintaining optimal sward height, and supplementing with high-energy feeds can further reduce methane intensity through improved forage quality and increased animal productivity (Gerber et al., 2013a).

Beyond dietary and management influences, genetic variation among and within species represents an emerging avenue for methane mitigation. Heritability estimates for methane production on an absolute basis (g CH₄/day) are moderate—around 0.29 for sheep and 0.40 for cattle—suggesting that methane output has a genetic component that could be exploited through selective breeding (Pickering et al., 2015). However, when expressed relative to feed intake (g CH₄/kg dry matter

intake, DMI), heritability declines substantially (0.13 - 0.19), indicating that methane yield is more strongly influenced by environmental and management factors (Pickering et al., 2015).

Despite the potential for genetic selection, practical challenges limit its application. Methane emissions fluctuate over time due to variation in DMI, feed fermentability, and animal physiological status, making it difficult to measure representative long-term phenotypes (Beauchemin et al., 2020).

Efforts are also underway to identify reliable proxies for CH₄ emissions that could enable large-scale genetic selection, including rumen microbial markers, milk fatty acid profiles, and specific metabolites (Negussie et al., 2017). However, the genetic correlations between CH₄ output and key production traits remain ambiguous. Breider et al. (2019) reported positive correlations between CH₄ production and milk yield, implying that animals genetically predisposed to higher productivity may also emit more methane. This relationship complicates the potential benefits of direct selection for lower methane output, as it could inadvertently reduce production efficiency. Consequently, indirect selection via productivity and feed efficiency traits may be a more practical route for mitigating emissions, since improving growth and milk yield typically reduces CH₄ intensity per unit of output (Beauchemin et al., 2020).

Overall, both feeding systems and genetic variation play critical roles in shaping methane emissions. While dietary strategies offer immediate and flexible mitigation potential, integrating genetic selection for feed efficiency and low-emitting phenotypes provides a long-term opportunity to achieve sustained reductions in enteric methane emissions across ruminant production systems.

Rumen pH, passage rate, and forage-to-concentrate ratio

Rumen environmental conditions, particularly pH, passage rate, and the forage-to-concentrate ratio, exert a profound influence on ruminal fermentation patterns and consequently on enteric methane production. Rumen pH plays a central regulatory role in shaping microbial community composition and activity. Optimal rumen pH (6.0–7.0) supports fibrolytic bacteria responsible for cellulose and hemicellulose degradation, leading to acetate production and higher hydrogen availability for methanogenesis (Kumar et al., 2014). When diets are rich in concentrates, increased fermentation of starches leads to greater production of lactic and propionic acids, which in turn lowers ruminal pH. A moderately reduced pH suppresses the growth and activity of methanogenic archaea and protozoa, both major contributors to H₂ generation and CH₄ formation (Hook et al., 2010). However, excessively low pH (<5.8) can induce subacute ruminal acidosis, impairing fibre digestion and overall feed efficiency (Beauchemin et al., 2020). Therefore, methane mitigation through dietary acidification must be balanced with maintaining an environment conducive to rumen health and productivity.

The rate of digesta passage through the rumen is another important determinant of methane yield, as it dictates the residence time available for microbial fermentation. Faster passage rates, often associated with higher feed intake and greater feeding frequency, reduce the exposure of substrates to methanogens, thereby decreasing total CH₄ output per unit of feed (Janssen, 2010). Shorter retention times can also limit protozoal colonisation and the establishment of methanogenic consortia on feed particles. In contrast, slower passage rates—typical of high-forage diets - favour extended microbial degradation of fibrous material, leading to enhanced acetate production and higher methanogenesis potential (Knapp et al., 2014). Moreover, physical factors such as particle size and effective fibre content influence rumen motility and liquid turnover, indirectly modulating fermentation kinetics and methane emission intensity (Hristov et al., 2013).

The forage-to-concentrate ratio (F: C) fundamentally alters ruminal fermentation pathways and the distribution of volatile fatty acids (VFAs). Diets with high forage inclusion stimulate cellulolytic fermentation and promote the formation of acetate and butyrate, which generate reducing equivalents H_2 that serve as substrates for methanogenesis (Hook et al., 2010; Knapp et al., 2014). Conversely, higher proportions of concentrates favour amylolytic fermentation, leading to increased propionate synthesis—a major alternative hydrogen sink that competes with methanogenesis for available H_2 (Morgavi et al., 2010). As a result, decreasing the F: C ratio typically reduces methane production per unit of feed intake. Nonetheless, excessively high concentrate levels may compromise rumen function and fibre utilisation, emphasising the importance of achieving an optimal dietary balance.

The relationship between forage proportion and methane yield is also influenced by forage quality and digestibility. Highly digestible forages, such as young temperate grasses or legumes, can enhance passage rate and reduce methane yield compared with mature, lignified forages (Benchaar, 2016). In mixed diets, incorporating moderate levels of rapidly fermentable carbohydrates (e.g., grains or molasses) can improve microbial efficiency and shift fermentation toward propionate without inducing acidosis (Hristov et al., 2013). Additionally, forage processing techniques, such as chopping or pelleting, may indirectly affect methane emissions by altering rumen retention time and fermentation site dynamics (Beauchemin et al., 2020).

Overall, maintaining optimal rumen pH, balancing forage and concentrate proportions, and managing feed particle size and turnover are critical strategies for regulating methanogenesis. These factors are interdependent: diet composition influences rumen pH and passage rate, while microbial adaptation modulates how efficiently energy is partitioned between methane production and alternative hydrogen-utilising pathways. Hence, an integrated understanding of rumen physicochemical dynamics is essential for designing effective, sustainable methane mitigation strategies in ruminant nutrition.

1.3. Classes of safe dietary compounds for methane mitigation

1.3.1. Plant secondary metabolites

Plant secondary metabolites are increasingly recognised as natural and safe feed additives for reducing enteric methane (CH_4) emissions in ruminants. They exert antimethanogenic effects primarily by inhibiting methanogens, modulating rumen protozoa, and redirecting hydrogen towards alternative pathways.

1.3.1.1. Tannins

Tannins are polyphenolic compounds that can be classified into condensed tannins (CTs), which are high molecular weight, and hydrolysable tannins (HTs), which have lower molecular weight. Condensed tannins primarily bind to dietary proteins, whereas hydrolysable tannins exert direct effects on rumen microbial populations (Aboagye and Beauchemin, 2019). Both types of tannins can reduce methane production. According to (Jayanegara et al., 2015), hydrolysable tannins are generally more effective in lowering methane concentrations than condensed tannins. The antimethanogenic effect of condensed tannins is largely attributed to a reduction in fibre digestion, while hydrolysable tannins primarily inhibit the growth and activity of methanogenic archaea.

In vitro studies demonstrated antimethanogenic activity of both HTs and CTs, by inhibition of methanogens or by targeting protozoa (Bhatta et al., 2009; Jayanegara et al., 2015). The in vivo effects depend on the dose, molecular weight, source and type (Jayanegara et al., 2012). A study by Jayanegara et al. (2011) found that the effects of tannins on methane emissions were more pronounced in in vitro settings compared to in vivo studies.

Grazing tannin-rich forages or adding extracts to the diet can also affect positive ruminants by preventing bloat, increasing N utilisation, lowering CH₄ production, acting as an antioxidant, controlling parasites, and improving animal production (Min et al., 2003; Gladine et al., 2007; Huang et al., 2018).

Acacia spp., particularly *Acacia mearnsii* and *Acacia nilotica*, are rich in condensed tannins and have been extensively studied for their antimethanogenic properties. Supplementation with *Acacia* leaf meal or extracts has been shown to reduce methane production in both in vitro and in vivo studies (Patra and Saxena, 2011; Aboagye and Beauchemin, 2019).

Quebracho is a tree native to South America and is a prominent source of condensed tannins. Quebracho tannins have demonstrated consistent reductions in methane emissions, likely due to their ability to bind dietary proteins and inhibit fibre-digesting microbes, indirectly limiting hydrogen availability for methanogenesis (Beauchemin et al., 2020). In vitro screening experiments report dose-dependent decreases in CH₄ production when quebracho CT is added to rumen incubations, with reductions often ranging from modest (10–20%) to large (>40%), depending on CT concentration and substrate (Battelli et al., 2023). Recent in vivo trials corroborate these findings but also illustrate important trade-offs: controlled dairy goat and cow experiments using commercial quebracho extracts (2–6% of diet DM) observed lowered methane yield and shifts in archaeal and bacterial community structure, yet some studies reported small decreases in fibre digestibility or dry matter intake at higher inclusion levels (Arik et al., 2024; Battelli et al., 2024). Comparative work suggests quebracho CTs tend to bind dietary protein more readily and may have a different impact on fibre-degrading populations than some other CT sources (e.g., *Acacia*) (Brutti et al., 2023). Very recent trials have begun pairing quebracho with other additives (e.g., essential oils or dietary lipids) to seek additive or synergistic methane mitigation while minimising negative effects on intake and productivity; early results are promising but indicate that reporting CT composition (molecular weight, degree of polymerisation) and standardising dose as % DM are essential for cross-study comparisons and practical recommendations (Foggi et al., 2022; Min et al., 2022).

Chestnut contains hydrolysable tannins and has been used in ruminant diets to suppress methane production. One proposed mechanism is that chestnut HTs act more directly on methanogenic archaea or their hydrogen supply, rather than predominantly inhibiting fibre-digesting bacteria as is often attributed to condensed tannins (Jayanegara et al., 2015). For example, an in vitro study comparing CTs and HTs found that while both reduced CH₄ production, the chestnut (HT) treatment was less effective than quebracho (CT) but did not reduce archaeal or protozoal counts in that short incubation (Battelli et al., 2023). Further in vitro experiments demonstrated that chestnut HT decreased methanogenesis and ammonia production while promoting acetate formation (Jayanegara et al., 2012; Cappucci et al., 2021). In vivo, supplementation with chestnut extract (4% DM) reduced methanogen abundance in lambs (da Costa, 2017) and decreased CH₄ emissions while improving milk performance in dairy cows (Prodanović et al., 2025), supporting its potential as a practical mitigation strategy.

In addition to conventional tannin sources, agro-industrial by-products such as grape marc have gained increasing attention as a methane mitigation strategy, particularly in Australia where the wine industry produces substantial quantities of this residue. Grape marc, consisting of grape skins, seeds, and stems remaining after wine production, is rich in condensed tannins and lipids, both of which contribute to its antimethanogenic properties. In vivo studies in dairy systems have demonstrated that dietary inclusion of grape marc can reduce enteric CH₄ emissions by approximately 10–20%, primarily through a combination of direct inhibition of methanogenic archaea and shifts in rumen fermentation pathways, including reduced hydrogen availability (Moate et al., 2014). Similarly, in vitro studies have reported that the efficacy of grape marc is influenced by its chemical composition, particularly tannin structure and fatty acid profile, which can vary

depending on grape variety and processing conditions (Hixson et al., 2018). Despite these benefits, high inclusion levels may reduce diet digestibility and metabolisable energy content, highlighting the importance of optimising inclusion rates to balance methane mitigation with animal performance. Given its availability and dual role as a feed supplement and waste valorisation strategy, grape marc represents a promising and regionally relevant option for sustainable methane mitigation in Australian ruminant systems.

Several other tropical and subtropical leguminous plants, such as *Leucaena leucocephala*, *Flemingia macrophylla*, and *Calliandra calothyrsus*, are rich in condensed tannins. These plants have been incorporated into ruminant diets to reduce methane emissions while providing protein-rich feed (Patra, 2013).

Despite their potential, the use of tannins as methane mitigation agents presents some limitations. Palatability: high concentrations of tannins can reduce feed intake due to their astringent taste, impacting animal performance (Beauchemin et al., 2020); Nutrient Digestibility: excessive tannin levels can bind to proteins and other nutrients, reducing their availability and digestibility. This binding action can lead to reduced nutrient absorption and may impair the efficiency of feed utilisation (Besharati et al., 2022). The safe inclusion level of hydrolysable tannins in ruminant diets is generally considered to be below 20–30 g/kg DM, while concentrations exceeding 50 g/kg DM may lead to adverse effects, including reduced intake, impaired digestibility, and potential toxicity (Makkar, 2003; Jayanegara et al., 2012). At very high levels (e.g., >100 g/kg DM), tannins can exert toxic effects such as liver and kidney damage, particularly hydrolysable tannins (Makkar, 2003); Animal adaptation: over time, rumen microbes may adapt to the presence of tannins, potentially diminishing their effectiveness. reducing their impact on methane production (Makkar, 2003).

While these plants are effective for methane mitigation (typical reduction 10–30%) (Patra and Saxena, 2011; Jayanegara et al., 2012), their inclusion in diets must be carefully managed to avoid negative effects on feed intake and nutrient digestibility. The type of tannin (condensed vs. hydrolysable), its concentration, and the method of processing (e.g., drying, ensiling) all influence the extent of methane reduction and animal performance (Makkar, 2003; Jayanegara et al., 2015; Besharati et al., 2022).

1.3.1.2. Saponins

Saponins are glycosides, which means they consist of a hydrophobic component called sapogenin linked to hydrophilic sugar molecules (Goel and Makkar, 2012). Growing evidence indicates that saponins can help reduce enteric CH₄ emissions in ruminants, primarily by influencing rumen protozoa and the methanogenic archaea associated with them (Goel and Makkar, 2023; Pepeta et al., 2024; Yanza et al., 2024).

One of the primary mechanisms by which saponins reduce methane emissions is defaunation—the suppression or elimination of rumen protozoa. Rumen protozoa play an important role in methane production because they harbor symbiotic methanogens on their surfaces and within their cells, providing H₂ through the fermentation of carbohydrates (Newbold et al., 2015). By reducing protozoal populations, saponins indirectly limit the availability of H₂ for methanogenesis (Goel and Makkar, 2012). Additionally, Saponins interact with sterols present in protozoal cell membranes, causing increased membrane permeability, leakage of intracellular components, and ultimately cell lysis (Kholif, 2023). This membrane-disrupting effect is selective, as ruminal bacteria generally lack sterols in their membranes and are less affected (Wallace, 2004). The reduction in protozoal biomass leads to a decline in methane yield due to the disrupted protozoa–methanogen symbiosis (Goel and Makkar, 2012).

Several plant species are rich in saponins and have been investigated for their antimethanogenic potential. Two major commercial and research-relevant sources include *Yucca schidigera* and *Quillaja saponaria*.

Yucca schidigera (a desert plant native to North America) produces steroidal saponins that have been shown to reduce rumen protozoa populations and methane production in vitro and in vivo (Makkar et al., 1998; Wang et al., 2012). Supplementation with *Yucca* extract has achieved methane reductions of 10–30%, depending on dose and diet type (Goel and Makkar, 2012).

Quillaja saponaria, a South American tree, contains triterpenoid saponins with similar properties. *Quillaja* extracts have demonstrated protozoal inhibition and reduction in methane yield without major negative effects on total volatile fatty acid production, suggesting that fermentation efficiency is largely maintained (Pen et al., 2006; Holtshausen et al., 2009).

Other saponin-containing plants such as *Sapindus mukorossi*, Tea (*Camellia sinensis*), *Trigonella foenum-graecum* (fenugreek), and *Enterolobium cyclocarpum* have also been explored for similar purposes (Jayanegara et al., 2014). The antimethanogenic efficacy of saponins is highly variable, depending on the saponin type, concentration, diet composition, and rumen microbial adaptation (Jayanegara et al., 2014). Some studies report that the methane-suppressing effect diminishes over time, likely due to microbial adaptation—rumen microorganisms can degrade saponins into sapogenins, which lack defaunating activity (Wang et al., 2012). This adaptation may occur within weeks of supplementation, suggesting that the persistence of the antimethanogenic effect is limited unless saponin sources or concentrations are varied periodically.

In addition, excessive inclusion of saponins may reduce feed intake or fibre digestibility, particularly when protozoal suppression is extensive (Patra and Saxena, 2009). Therefore, identifying optimal inclusion levels (often 0.5–1.5 g saponin/kg DM) and delivery methods remains critical for practical application (Goel and Makkar, 2012).

Beyond their role in methane mitigation, saponins may offer additional health benefits in ruminant production systems. Their antimicrobial and antioxidant properties can enhance rumen microbial balance, gut health, and immune function (Cheeke, 2000). By reducing protozoal populations, saponins may also help control rumen ammonia concentrations, improving nitrogen utilisation efficiency (Patra and Saxena, 2009; Wang et al., 2012). Moreover, saponins have been reported to reduce incidences of bloat, acidosis, and parasitic infections, contributing to improved animal welfare and productivity (Wallace, 2004; Goel and Makkar, 2012). However, these health-promoting effects are dose-dependent and can be reversed at excessive inclusion levels, underscoring the need for careful formulation and monitoring.

Saponins present a promising natural strategy for methane mitigation (typical reduction 5–25%) (Pen et al., 2006; Goel and Makkar, 2012) through selective inhibition of rumen protozoa and disruption of methanogen–protozoa associations. Extracts from *Yucca schidigera* and *Quillaja saponaria* are the most widely studied and commercially viable options. However, the transient nature of their effects due to microbial adaptation and variable responses across diets highlights the need for continued research into formulation stability, dosage optimisation, and potential synergies with other feed additives.

1.3.1.3. Essential oils

Essential oils (EOs) are complex mixtures of volatile secondary metabolites extracted from aromatic plants. Their primary active components—such as thymol, eugenol, and cinnamaldehyde—belong to chemical groups including phenols, aldehydes, and terpenes, which confer strong antimicrobial and antioxidant properties (Calsamiglia et al., 2007; Benchaar and Greathead, 2011). Thymol and carvacrol are commonly derived from thyme (*Thymus vulgaris*) and oregano (*Origanum vulgare*), eugenol from clove (*Syzygium aromaticum*), and cinnamaldehyde from cinnamon (*Cinnamomum zeylanicum*). These compounds have been investigated for their potential to modulate rumen fermentation and reduce enteric methane emissions through selective inhibition of rumen microorganism (Benchaar and Greathead, 2011).

Commercial EO-based feed additives have been developed to provide standardised and safe inclusion

rates for livestock. Crina[®] Ruminants (DSM-Firmenich) and Agolin[®] Ruminant (Agolin SA/Alltech) are two of the most widely studied EO blend. Studies have reported modest but consistent reductions in methane emissions, generally between 5–10%, when these blends are administered at recommended dosages without negative effects on feed intake or animal performance (Belanche et al., 2020). Crina[®] Ruminants (DSM-Firmenich) is an EO-based additive formulated with thymol, eugenol, and vanillin. The combination is aimed at improving rumen fermentation and feed utilisation while reducing methane and nitrogen losses. Experimental work has evaluated CRINA[®] for effects on rumen fermentation, animal performance, and enteric methane. For instance, in a controlled Latin-square study with Brahman steers fed Rhodes grass hay, Crina[®] was compared to Monensin and an untreated group at a low inclusion rate. Crina[®] was supplemented at approximately 1 g/head per day, while monensin was included at around 200 mg/head per day. While Crina[®] altered fermentation profiles, it produced only modest or inconsistent reductions in methane production and changes in the methanogen community metrics under the tested conditions (Tomkins et al., 2015). Several in situ studies observed that the blend of essential oils inhibited protein degradation, although the reported changes were small and varied depending on the type of feed being degraded, the specific ration fed to the animals, and the duration of the adaptation period (Molero et al., 2004; Newbold et al., 2004). McIntosh et al. (2003) further suggested that this combination of essential oils likely inhibits a specific group of gram-positive bacteria known as hyper-ammonia-producing bacteria. However, most studies assessing Crina[®] have found no significant impact on microbial protein synthesis in the rumen (Wallace et al., 2002; Castillejos et al., 2005).

Agolin[®] Ruminant is a standardised EO blend containing coriander oil (*Coriandrum sativum*), geraniol, geranyl acetate, and eugenol. These bioactive substances have been documented to exhibit a wide range of antimicrobial effects, including activity against rumen microorganisms. (Benchaar and Greathead, 2011) and was certified by Carbon Trust Assurance LTD for reduction of CH₄ emissions and improvements of feed efficiency (Belanche et al., 2020). In a meta-analysis of in vivo studies, Belanche et al. (2020) found that Agolin supplementation reduced methane emissions by an average of 8.8% and increased feed efficiency by 2.5%, with consistent effects across diet types and production systems; However, based on additional research, the effects of Agolin supplementation are only temporary (Klop et al., 2017) or don't have an effect at all (Miller et al., 2023; Silvestre et al., 2023; Benchaar and Hassanat, 2025). The causes of these discrepancies are not well understood but might involve variations in experimental conditions (such as the composition of the diet), the feeding rate conveyed on a dry matter intake basis, and the traits of the cows (Cobellis et al., 2016; Benchaar and Hassanat, 2025). Agolin's effects appear largely indirect, through reduced hydrogen availability and selective inhibition of certain bacterial populations (e.g., *Ruminococcus albus* and *Methanobrevibacter spp.*). The blend's composition ensures efficacy at low inclusion rates (0.5–1 g/day per cow), minimising risks to palatability or rumen digestibility (Belanche et al., 2020).

The antimicrobial activity of EOs is attributed to their ability to disrupt microbial cell membranes, alter proton gradients, and interfere with enzymatic activity, particularly in Gram-positive bacteria (Benchaar et al., 2008). In the rumen, these effects can lead to shifts in the microbial ecosystem, reducing the abundance of hydrogen-producing bacteria and protozoa that are symbiotically associated with methanogenic archaea (Patra and Yu, 2012). Consequently, fermentation patterns may change toward a greater proportion of propionate and lower acetate, thereby decreasing the availability of H₂ required for methane formation (Calsamiglia et al., 2007).

Agolin and Crina specifically have been shown to enhance fermentation efficiency by stabilising rumen pH and reducing ammonia accumulation (Benchaar et al., 2008). However, excessive EO doses may depress overall fermentation and reduce fibre degradation (Patra and Yu, 2012). Prolonged exposure to EOs can also lead to microbial adaptation, as rumen microbial communities may adjust to the presence of EO compounds, diminishing their efficacy over time (Calsamiglia et al., 2007). Encapsulation technologies and rotation or blending of EO compounds have been proposed to

mitigate adaptation effects and sustain long-term activity (Belanche et al., 2020). Essential oils used in feed are generally recognised as safe (GRAS) when administered at low concentrations (U.S. Food and Drug Administration, 2023). Nonetheless, palatability issues and potential toxicity at high doses require careful consideration (Burt, 2004). Commercial products such as Crina® and Agolin® Ruminant are formulated to ensure safety, stability, and effective delivery, and are approved for use in ruminant nutrition in several jurisdictions.

Overall, EOs represent a promising natural feed additive for methane mitigation (typical reduction 5-20%) (Calsamiglia et al., 2007; Benchaar et al., 2008) and enhancement of rumen efficiency, though their effects are variable across studies due to differences in EO composition, diet type, and duration of exposure. Further long-term and large-scale in vivo trials are needed to fully understand the persistence of effects and the economic feasibility of implementation in grazing systems.

1.3.2. Electron acceptors and alternative hydrogen sinks

In the rumen, methanogenesis serves as the primary pathway for H₂ disposal, maintaining redox balance during fermentation. Alternative hydrogen sinks—compounds capable of accepting electrons generated during carbohydrate metabolism—can redirect reducing equivalents away from CO₂ reduction to CH₄, thereby lowering enteric methane emissions (Hristov et al., 2013; Ungerfeld, 2020). Among these, sulphate and fumarate are two of the most extensively studied electron acceptors.

Sulphate as an alternative electron acceptor

Sulphate (SO₄²⁻) reduction competes directly with methanogenesis for available H₂ in the rumen. Sulphate-reducing bacteria (SRB), such as *Desulfovibrio* spp. and *Desulfotomaculum* spp., utilise H₂ to reduce sulphate to hydrogen sulphide (H₂S). This process effectively diverts H₂ from methanogens, resulting in lower methane production (Li et al., 2013). In vitro studies have shown that adding sulphate to rumen cultures can significantly reduce CH₄ production, with inclusion levels typically ranging from approximately 5 to 20 mM sulphate in batch culture systems (Wu et al., 2015; Wu et al., 2021). Additionally, in vivo research using sulphate supplementation has demonstrated a reduction in methane emissions of 10–20%, depending on the dosage and sulphur content of the basal diet. (van Zijderveld et al., 2010; Silivong et al., 2011; Arif et al., 2016). Furthermore, it has been shown that incorporating sulphate into the ruminant diet increases the activity of fibre-digesting bacteria and rumen fungi (Wang et al., 2021).

However, the practical application of sulphate is constrained by the potential for hydrogen sulphide toxicity. Excessive production of H₂S can lead to polioencephalomalacia and respiratory distress in ruminants (Gould, 1998). Therefore, sulphate inclusion must be carefully managed to maintain total dietary sulphur below recommended limits (typically <0.4% of DM). Furthermore, high sulphate levels exceeding these dietary limits may impair palatability and affect mineral balance (Spears et al., 2011). Despite these limitations, strategic use of sulphate - particularly in low-sulphur diets - remains a promising, albeit safety-limited, methane mitigation strategy.

Fumarate as an alternative hydrogen sink

Fumarate, an intermediate of the tricarboxylic acid (TCA) cycle, serves as another effective electron acceptor in the rumen (Stewart et al., 1997). Through microbial reduction, fumarate is converted to succinate and subsequently to propionate, a VFA that acts as an alternative H₂ sink and provides an energy benefit to the host animal (Castillo et al., 2004). This pathway not only consumes reducing equivalents but also shifts fermentation toward propionate production, reducing hydrogen availability for methanogenesis (Ungerfeld, 2015a). In vitro and in vivo studies have shown that fumarate supplementation can reduce methane emissions depending on concentration and microbial adaptation (Lopez et al., 1999; Bayaru et al., 2001; García-Martínez et al., 2005; Newbold et

al., 2005). Despite its mechanistic efficiency, the practical use of fumarate is limited by cost and dose requirements. Effective methane mitigation often requires several grams of fumarate per kilogram of feed, which may be economically unfeasible for large-scale livestock operations (Hristov et al., 2013). Moreover, rapid fermentation of fumarate may decrease rumen pH, posing risks of subacute acidosis when combined with high-concentrate diets (Ungerfeld, 2015a). Consequently, fumarate is primarily considered useful as a research tool or component of combined mitigation strategies, rather than a standalone additive for commercial use.

Both sulphate and fumarate function by diverting hydrogen away from methanogenesis, thereby lowering methane yield. However, their practical application differs. Sulphate offers stronger competition for H₂ but carries toxicity risks, whereas fumarate provides an additional nutritional benefit through propionate formation but requires higher and costlier doses. The success of these compounds is highly dependent on diet composition, sulphur background, and microbial adaptation (van Zijderveld et al., 2010; Ungerfeld, 2015a).

Sulphate and fumarate are promising alternative hydrogen sinks that can decrease enteric methane emissions (typical reduction 5-15%) (Ungerfeld et al., 2007) by providing competing pathways for hydrogen utilisation. Sulphate reduction offers high theoretical potential but is constrained by toxicity risks, while fumarate reduction provides both mitigation and energetic benefits but is limited by economic and practical feasibility. Future research should explore controlled-release formulations to enhance their safety and cost-effectiveness in ruminant production systems.

1.3.3. Lipids and fatty acids

Lipids and fatty acids have been extensively studied as dietary interventions to mitigate enteric methane emissions in ruminants (Patra, 2013; Arndt et al., 2022). Their efficacy arises primarily through two mechanisms: direct inhibition of methanogenic archaea and indirect effects on rumen fermentation, particularly fibre digestion.

Lipids can reduce methane production by directly affecting methanogens, the archaea responsible for CH₄ synthesis. Unsaturated fatty acids can disrupt the cell membranes of methanogens, leading to reduced growth and activity (Patra, 2013). Additionally, the presence of lipids in the diet can decrease fibre digestion by coating feed particles and altering microbial populations, thereby reducing the availability of hydrogen, a substrate for methanogenesis (Beauchemin et al., 2008).

Medium-chain fatty acids (MCFAs), such as lauric and myristic acids, are particularly effective at inhibiting methanogens due to their strong antimicrobial properties. Unsaturated oils, including linoleic and α -linolenic acids, also contribute to methane mitigation by hydrogenation reactions in the rumen, which compete with methanogenesis for hydrogen (Jenkins et al., 2008).

Coconut oil, palm kernel oil, and their derivatives are major sources of medium-chain fatty acids, particularly lauric and myristic acids. These fatty acids exhibit direct inhibitory effects on methanogenic archaea and rumen protozoa, reducing methane formation (Patra, 2013). Coconut oil is among the most extensively studied lipid sources for mitigating enteric methane, with reported reductions ranging from 13% to 73%, depending on dosage, diet composition, and ruminant species. Its lauric-to-myristic acid ratio (2.6:1) aligns with the most effective ratios identified for methane inhibition *in vitro* (Soliva et al., 2004), supporting its potential for substantial *in vivo* efficacy. Palm kernel oil, which has a slightly higher lauric-to-myristic acid ratio (3:1), may provide even greater methane abatement potential; however, *in vivo* evidence is currently lacking (Hook et al., 2010). Nevertheless, supplementation with either oil frequently leads to reduced dry matter intake, which

can confound methane reduction results by lowering total feed fermentation (Jordan et al., 2006; Liu et al., 2011).

Linseed (*Linum usitatissimum*) is rich in α -linolenic acid (C18:3 n-3), which has shown potential to reduce methane emissions through suppression of methanogens and biohydrogenation pathways (Martin et al., 2010; Grainger and Beauchemin, 2011). Linseed oil or whole linseed supplementation at moderate levels typically decreases CH₄ yield (Martin et al., 2016). Similar effects have been observed with other omega-3-rich oils such as chia and perilla oil, indicating that n-3 fatty acids play a role in reducing hydrogen availability for methanogenesis (Grainger and Beauchemin, 2011). However, the efficacy of omega-3 supplementation varies depending on the form (oil vs. seed), dosage, diet composition, and adaptation period (Guyader et al., 2015). Excessive inclusion of unsaturated fats may impair fibre fermentation and reduce animal performance, highlighting the importance of balancing methane mitigation potential with production outcomes (Grainger and Beauchemin, 2011). Additionally, omega-3 enrichment of ruminant products (milk, meat) offers potential health benefits for human consumers, creating dual advantages of environmental and nutritional value (Shingfield et al., 2013).

Sunflower and canola (rapeseed) oils provide unsaturated fatty acids—mainly linoleic (C18:2 n-6) and oleic (C18:1 n-9)—which can act as alternative hydrogen sinks during rumen fermentation, thereby lowering methane production. Studies have shown reductions in CH₄ emissions including oil in the diet (Beauchemin et al., 2020). The inclusion of these lipids in ruminant diets has consistently demonstrated mitigation effects on CH₄ emissions, although the magnitude of reduction depends on the form and level of supplementation (Silveira et al., 2019). Canola and sunflower by-products, such as cakes and meals, provide these lipids in a matrix that is generally less inhibitory to fibre digestion than free oils, allowing for a more balanced approach between methane mitigation and rumen function (Arndt et al., 2022). Supporting this, (Brask et al., 2013) observed that methane intensity (L/kg organic matter intake) was 12.6% lower in cows fed rapeseed cake compared with those receiving rapeseed meal, likely due to differences in lipid content and fermentability between the two feed forms. Similarly, (Beauchemin et al., 2009) reported a 16% reduction in CH₄ emissions in dairy cows when dietary supplementation with canola meal. These findings collectively indicate that oilseeds and their by-products can effectively suppress ruminal methanogenesis through multiple mechanisms—such as the biohydrogenation of unsaturated fatty acids, reduced fibre degradation, and limited substrate availability for methanogens—while maintaining overall feed efficiency and animal performance.

Fish oils are rich in long-chain n-3 PUFAs such as eicosapentaenoic acid (EPA) and docosahexaenoic acid (DHA), which have potent antimethanogenic properties. These fatty acids can reduce methanogenesis directly through microbial inhibition and indirectly by decreasing ruminal protozoa populations. Inclusion of small amounts (up to 2–3% DM) of fish oil has achieved methane reductions, although high inclusion levels may impair fibre degradation (Kouřimská et al., 2014; Vargas et al., 2017).

Whole or processed oilseeds (e.g., flaxseed, cottonseed, sunflower seed) provide a slower lipid release than free oils, reducing potential negative effects on rumen fermentation. Grinding or crushing oilseeds enhances lipid bioavailability in the rumen (Beauchemin et al., 2020). Arndt et al. (2022) reported that both oil and oilseed supplementation reduced methane yield, indicating comparable mitigation effects when properly managed.

The antimethanogenic effects of dietary lipids in ruminants are highly variable (10-30%) (Benchaar et al., 2008) and depend on multiple factors, including lipid type, inclusion level, and animal species.

Medium-chain fatty acids (e.g., myristic acid) and polyunsaturated fatty acids (PUFAs) from fish, sunflower, linseed, and canola oils are generally the most effective at reducing methane emissions (Patra, 2013).

While lipids can significantly reduce methane emissions, excessive inclusion may impair fibre digestion and reduce dry matter intake, negatively affecting animal productivity (Eugène et al., 2008). Therefore, dietary strategies must balance the level and type of lipid supplementation to optimise methane reduction while maintaining feed efficiency and animal performance (Palmquist and Jenkins, 2017). Inclusion of fats at 4–6% of dietary dry matter has been reported to improve milk production in ruminants. This beneficial effect is largely attributed to the additional energy provided by lipids and their potential influence on milk fat synthesis. However, feeding diets with higher fat concentrations can disrupt rumen microbial activity, inhibit fibre-digesting bacteria, and reduce nutrient digestibility, ultimately impairing rumen fermentation and overall animal performance (Patra, 2013). The negative effects are particularly pronounced when fats are supplemented in forms that are readily bioavailable in the rumen, highlighting the importance of careful selection of fat type and inclusion level in ration formulation (Arndt et al., 2022).

Long-term effects of dietary lipids are less well documented. Some studies indicate persistent antimethanogenic effects (Jordan et al., 2006), whereas recent research on grazing dairy cows suggests that the effects of oilseed supplementation may be transient (Muñoz et al., 2021). The inhibitory effect of lipids is generally greater in concentrate-based diets than in forage-based diets, potentially due to the lower rumen pH in concentrate-fed animals, which enhances the impact of fatty acids on methanogens (Beauchemin et al., 2022).

Additional research is required to determine economically viable and sustainable fat sources, as well as their optimal inclusion levels, that can mitigate methane emissions without compromising feed digestibility or animal performance. Furthermore, studies are needed to evaluate the long-term effects of fat supplementation on the persistence of methane reduction.

1.3.4. Synthetic compounds and feed additives

1.3.4.1. 3-Nitrooxypropanol (3-NOP)

3-Nitrooxypropanol (3-NOP) is a small synthetic molecule that specifically targets the terminal step of biological methanogenesis. Mechanistically, 3-NOP acts as an inhibitory substrate for methyl-coenzyme M reductase (MCR), the nickel-containing enzyme in methanogenic archaea that catalyses the final reduction of methyl-coenzyme M to CH₄, forming stable complex that prevent further turnover (Martinez-Fernandez et al., 2016). Because this mechanism acts directly on methanogens rather than on hydrogen-producing bacteria, it results in targeted suppression of CH₄ formation without major disruption of overall rumen fermentation (Hegarty et al., 2021).

Multiple controlled studies and meta-analyses have confirmed the robust efficacy of 3-NOP in reducing enteric methane emissions across dairy and beef cattle, sheep, and goats. The typical dosage reported in studies is 123 mg 3-NOP per kg of dry matter intake (DMI), which has been demonstrated to result in a 23% reduction in emissions for beef and a 39% reduction for dairy (Hegarty et al., 2021). A global meta-analysis reported average methane reductions of 29 % (Kebreab et al., 2023). Similar to the results found by (Dijkstra et al., 2018), with minimal effects on dry-matter intake and feed efficiency. These effects have been reproduced under both high-forage and high-concentrate diets. The composition of the diet can influence how effective antimethanogenic additives are. For instance, research shows that the ability of 3-NOP to reduce methane emissions diminishes as the concentration of NDF in the diet increases (Kebreab et al., 2023), probably because

there is greater hydrogen production linked to acetate-type fermentation in the rumen (Hristov, 2023).

No strong evidence of microbial adaptation that restores methane production has been reported in studies (Hristov, 2023). Meta-regression analyses further indicate that baseline methane intensity and diet fermentability influence absolute reductions, but proportional effects are consistent across production systems (Kebreab et al., 2023).

Safety evaluations indicate that 3-NOP is rapidly metabolised in the rumen and animal tissues, and at recommended inclusion rates does not cause adverse effects on animal health, milk or meat safety, or consumer exposure (Bampidis et al., 2021). Several jurisdictions have completed regulatory evaluations and authorised the commercial product Bovaer® (active: 3-NOP) for use in ruminant feed under specified conditions. Regulatory approvals include—but are not limited to—the European Union (for certain dairy applications), Brazil, Chile, Canada, Australia, and the United States. Commercial rollout and on-farm implementation (via Bovaer® and equivalent formulations) are now documented in multi-site trials and company-sponsored large-scale trials showing real-world reductions consistent with experimental literature. However, labelling, permitted species, approved dose ranges, and target production categories vary by jurisdiction, and users should consult the relevant regulatory authorisation for precise use conditions.

Because 3-NOP acts by enzymatic inhibition, consistent daily intake is critical for maintaining efficacy; feed-delivery variability can attenuate its effect (Kebreab et al., 2023). Production responses are generally neutral. Research on grain-based diets indicated that incorporating 3-NOP did not provide any benefits for live weight gain, and there could be a decline in growth when high levels of 3-NOP are included (Hegarty et al., 2021). Long-term trials are still required to confirm persistence of efficacy, cost–benefit ratios, and potential interactions with other mitigation compounds such as nitrate or tannins (Hristov, 2023). Overall, 3-NOP currently represents the most technically validated and regulatorily approved chemical inhibitor of enteric methane formation (20–60%) (Hristov et al., 2015; Melgar et al., 2020) available. However, a practical limitation of this compound is its relatively short shelf-stability and susceptibility to degradation during feed processing and storage. Bampidis et al. (2021) reported losses of 16.7% of 3-NOP in pelleted feed after three months of storage at room temperature in sealed polyethylene bags. Although some exposure to oxygen and variable humidity was still possible during storage (Bampidis et al., 2021). Similarly, MLA-funded research has shown that the inherent volatility of 3-NOP can lead to variable losses during pelleting and storage, reducing the consistency of additive delivery to animals (Lawrence et al., 2024). These findings highlight the importance of careful formulation, processing, storage, and handling protocols to preserve the compound’s stability and ensure its intended antimethanogenic efficacy under commercial feeding conditions.

1.3.4.2. Ionophores (e.g., Monensin) – very efficient to decrease CH₄ in vitro but not in vivo

Ionophores such as monensin are carboxylic polyether antibiotics that alter rumen fermentation by selectively increasing the permeability of microbial cell membranes to monovalent cations (Na⁺, K⁺), thereby favouring Gram-negative bacteria and shifting fermentation toward increased propionate production at the expense of acetate (Russell and Strobel, 1989). These shifts reduce hydrogen availability for methanogenesis and can depress methane formation in vitro and, under some conditions, in vivo. Ionophores commonly increase propionate: acetate ratios, reduce ruminal ammonia accumulation, and can improve feed efficiency in beef finishing systems (Hegarty et al., 2021).

Recent meta-analyses focusing on beef cattle show transient methane reductions that depend on diet composition (forage vs concentrate), ionophore dose, and duration of administration; sustained

long-term mitigation is inconsistent (Cooke et al., 2024). Ionophore efficacy is typically greater in the short term or in high-grain diets where shifts to propionate are pronounced (Appuhamy et al., 2013), but adaptation of the rumen microbiome can reduce the magnitude of effect with prolonged exposure (Rychlik and Russell, 2002). A meta-analysis reported the most significant results during the first 79 days of monensin supplementation, particularly on high forage diets offered for *ad libitum* intake (Cooke et al., 2024). Consequently, monensin is regarded as nutritionally beneficial but not a reliable long-term methane mitigant under practical feeding conditions.

Ionophores are classified as veterinary antibiotics but are not used in human medicine; nevertheless, their use raises broader concerns about selection for antimicrobial resistance (AMR) determinants in the gut microbiome and potential co-selection of resistance to other antibiotic classes (McEwen & Fedorka-Cray, 2002). Experimental and observational studies demonstrate that ruminal bacteria can develop reduced sensitivity to ionophores, and shifts in bacterial community composition may select for tolerant strains (Rychlik and Russell, 2002). Several countries have limited or prohibited routine antibiotic use for growth promotion, but ionophores remain permitted in ruminant nutrition under veterinary supervision (Callaway et al., 2003). The potential antimicrobial resistance consequences and the limited, variable long-term methane mitigation observed with ionophores argue that they are a suboptimal strategy for sustained greenhouse-gas mitigation.

In summary, ionophores exert useful fermentation-modulating effects and transient methane suppression (5-10%) (McGinn et al., 2004; Guan et al., 2006), but microbial adaptation considerations restrict their role as a sustainable mitigation technology.

1.3.5. Probiotics, enzymes, and other emerging bioactives

Use of direct-fed microbials (DFM):

Direct-fed microbials (DFM; “probiotics” for ruminants) aim to alter rumen fermentation by introducing or enriching microbial taxa that (a) redirect metabolic hydrogen toward non-methanogenic products (e.g., propionate), (b) compete with hydrogenotrophic methanogens, or (c) stabilise rumen conditions (pH, redox) that disfavour methanogenesis (Newbold et al., 2015).

Key candidate organisms used are *Megasphaera elsdenii* and *Propionibacterium spp*; *M. elsdenii* consumes lactate and produces volatile fatty acids (VFA) such as butyrate and propionate; it has been proposed as a DFM to prevent acidosis during high-grain transitions and increase propionate formation (thereby acting as an H₂ sink relative to methanogenesis). Strain selection, dose, and dietary context strongly influence outcomes; reviews and experimental work report benefits for rumen stability and feed efficiency but inconsistent methane reductions and occasional associations with milk-fat depression in dairy cows (Beauchemin et al., 2020); Propionibacteria and other propionate-forming bacteria are of interest because propionate formation consumes reducing equivalents that would otherwise form CH₄ (Jeyanathan et al., 2016). Some in vitro and short-term in vivo studies show increased propionate and modest methane decreases when propionigenic DFMs are added (Vyas et al., 2014; Vyas et al., 2016), but effects are variable (Meale et al., 2014; Chen et al., 2020). Current data indicates that while propionibacteria can produce propionate to utilise hydrogen during fermentation, this process does not reduce CH₄ emissions (Hegarty et al., 2021).

Controlled in vitro studies and small-scale in vivo trials show that certain DFMs can shift fermentation toward propionate and reduce measured methane production under specific diets (low pH, high fermentability) (Oyebade et al., 2024). However, consistent meaningful methane reductions under commercial, long-term conditions are not yet well established; responses vary with basal diet, strain viability in the rumen, interaction with resident microbiota, and delivery format (capsules, boluses, feed premix) (Oyebade et al., 2024).

Enzyme supplements to alter fermentation pathways.

Exogenous feed enzymes (e.g., cellulases, xylanases, pectinases, amylases, and some dehydrogenases) are used to improve fibre and starch digestion and, in doing so, can alter fermentation kinetics and end-product profiles. Where enzymes increase the proportion of propionate or accelerate feed conversion, they may indirectly reduce H₂ availability for methanogens (Salem et al., 2023).

Experimental work and recent reviews show that tailored enzyme blends can improve nutrient digestibility and feed conversion in ruminants (Carrillo-Díaz et al., 2022). Reports of direct, reproducible methane reductions attributable solely to enzymes are limited (Palangi and Lackner, 2022). Many enzyme trials focus on performance (DMI, ADG, milk yield) and fibre digestibility rather than methane as a primary endpoint. Where methane was measured, reductions are often small and diet-dependent (higher potential in low-quality forages where enzyme-induced increases in fermentability shift VFA profiles) (Salem et al., 2023).

Enzyme effectiveness depends on substrate accessibility, enzyme specificity, and rumen residence time; encapsulation or targeted delivery may increase efficacy.

The industrial-scale production of anaerobic bacteria remains technically demanding and substantially more expensive than the conventional cultivation of microorganisms under aerobic conditions (McAllister et al., 2011). This constraint currently limits the range of DFM that can be developed for practical, on-farm application (Gallo et al., 2015; Ban and Guan, 2021). Moreover, maintaining the viability of obligate anaerobes during storage and distribution poses a significant challenge, as these organisms are highly sensitive to oxygen exposure and environmental stress (Hegarty et al., 2021). Continued technological innovation in microbial formulation and stabilisation is therefore critical to broaden the potential use of anaerobic strains in methane mitigation strategies. For novel DFM candidates, *in vivo* evaluation is particularly important (Doyle et al., 2019), as microbial interactions and rumen responses observed in live animals often differ from those detected *in vitro*. Such studies are essential to assess potential shifts in the rumen microbiome and to ensure that no unintended effects occur on animal health, productivity, or product safety. Feed additive approval requires detailed evidence of safety, efficacy, and stability, and considerable variation exists in product quality and strain identity among commercial DFM preparations (Hegarty et al., 2021). In addition, the persistence and consistency of the antimethanogenic response must be validated through long-term animal trials, given that microbial adaptation or loss of efficacy may occur over time. Like other nutritional mitigation approaches, DFM require continuous daily administration (Alazeh et al., 2012) to maintain their beneficial effects, highlighting the need for cost-effective, stable, and user-friendly formulations suited to commercial production systems (Beauchemin et al., 2022).

Overall, while DFM represent a promising biological tool for improving rumen efficiency and mitigating enteric CH₄ emissions, their widespread use will depend on developing more robust strains, clarifying mechanisms of action, and demonstrating consistent economic returns under commercial production systems (Jeyanathan et al., 2014; Beauchemin et al., 2022).

1.4. Integrated approaches and synergies

The combination of feed additives may produce additive or synergistic effects. Combining compounds can target different steps or groups in ruminal H₂/CH₄ production (e.g., inhibit methanogenesis directly vs redirect H₂ to alternative sinks), lower the dose required of any single additive (reducing cost or side-effects), and reduce the likelihood of microbial adaptation to one single mode of action. However, combinations can also be antagonistic (no extra benefit) or produce

negative effects on intake, digestibility, animal production or animal health (del Prado et al., 2025; Hristov et al., 2025). Trials combining 3-NOP with other additives (different feed types, cottonseed, monensin) show mixed outcomes—some additive gains, some neutral interactions—indicating diet context matters (Kebreab et al., 2023). Studies of tannin + saponin blends report greater CH₄ reductions than either alone in some trials (e.g., reduced CH₄ per kg digested NDF), suggesting beneficial interactions between plant secondary metabolite classes (Cardoso-Gutierrez et al., 2021; Cuervo et al., 2024).

Ionophores alter fermentation (shift VFA profile toward propionate, a hydrogen sink). Combining ionophores with direct methanogenesis inhibitors can be additive, but results are inconsistent — some studies show no extra benefit when combined with 3-NOP or monensin+3-NOP (Romero-Pérez et al., 2017; Muñoz et al., 2024). Probiotics that enhance propionate producers or enzymes that alter fibre degradation may complement chemical inhibitors by improving H₂ fluxes, but evidence is early and variable (del Prado et al., 2025). Thus, optimal additive combinations should be matched to diet type to maximise mitigation efficiency without impairing animal performance.

In grazing-based systems, where direct feed supplementation is often impractical, the delivery method becomes a critical factor influencing the consistency and efficacy of additive intake (Beauchemin et al., 2022). Molasses-based lick blocks have emerged as a practical strategy for administering methane-mitigating compounds under extensive conditions (Imaz et al., 2020). These blocks provide a controlled-release vehicle that promotes steady consumption and uniform exposure across animals, while their molasses base enhances palatability and energy supply (Zhao et al., 2022). The lick block approach also allows the integration of multiple additive types, potentially facilitating additive or synergistic methane reduction effects when formulated appropriately for the prevailing diet composition and production system (Hristov et al., 2025).

Interaction of feed additives with diet composition

The efficacy of methane-mitigating additives is strongly influenced by diet composition, particularly the forage-to-concentrate (F: C) ratio, which determines ruminal fermentation pathways, hydrogen (H₂) availability, and microbial community structure (Hook et al., 2010; Patra and Yu, 2012). Diets rich in forage promote fibrolytic fermentation and acetate production, generating more H₂, whereas high-concentrate diets enhance propionate formation, a natural hydrogen sink, thereby lowering baseline methane yield (Beauchemin et al., 2020). Consequently, the same additive can produce markedly different outcomes depending on the diet matrix; The effectiveness of plant secondary metabolites depends on both their chemical form and the substrate fermented. In high-forage diets, condensed tannins can form complexes with fibre and proteins, potentially reducing digestibility but also inhibiting methanogens and protozoa (Jayanegara et al., 2015). In concentrate-based diets, their effects on methane are often smaller because the baseline methanogenesis is already reduced. Saponins show stronger protozoa-suppressing activity in forage-based diets where protozoal populations are higher (Patra and Yu, 2015). Similarly, essential oils (EO) tend to show greater inhibition of methanogens in high-forage diets due to longer rumen retention time, allowing sustained antimicrobial contact (Cobellis et al., 2016). Electron acceptors such as sulphate, nitrate, and fumarate compete with CO₂ for available H₂. Their response depends heavily on ruminal H₂ partial pressure, which is generally higher in forage-based systems (Ungerfeld, 2020). Consequently, fumarate or nitrate supplementation tends to be more effective under high-forage conditions, where hydrogen availability is abundant. In contrast, in high-concentrate diets with inherently low H₂ production, these compounds may yield limited additional CH₄ reduction or may cause redox imbalances if excessive doses are used. The inclusion of unsaturated fatty acids can reduce CH₄ through biohydrogenation and inhibition of protozoa and methanogens. However, the magnitude of inhibition depends on diet composition. Lipid supplementation is generally more effective in high-forage diets, where hydrogen availability and microbial diversity allow greater scope for redirection

of reducing equivalents (Beauchemin et al., 2020). In concentrate-rich diets, high lipid levels can depress fibre digestion and animal performance, offsetting the CH₄ benefits (Grainger and Beauchemin, 2011). The response to 3-nitrooxypropanol (3-NOP) is diet-dependent. Studies show higher efficacy in high-concentrate diets, where the additive is more stable and less degraded in the rumen (Romero-Perez et al., 2015). In forage-based systems, 3-NOP effectiveness can decline due to increased rumen passage time and microbial adaptation (Melgar et al., 2020). Conversely, ionophores (e.g., monensin) are more effective in concentrate-based diets, where they favour propionate-producing bacteria and suppress hydrogen formation, although responses are moderate when H₂ availability is already low (Russell and Houlihan, 2003). Probiotic and enzyme supplements interact with diet fibre content. Fibrolytic enzymes and certain propionate-producing bacterial strains exhibit greater methane-mitigating effects in high-forage diets by enhancing fibre breakdown while redirecting reducing equivalents (McAllister and Newbold, 2008). In concentrate diets, such additives may have minimal CH₄ effects but can improve feed efficiency and rumen health, indirectly influencing emissions intensity.

Additive or synergistic effects among compounds often depend on diet composition. For instance, 3-NOP combined with fumarate or lipids shows additive reductions in methane emissions in high-forage diets due to complementary hydrogen sinks, while tannin–saponin mixtures have greater effects under grazing systems where plant diversity supports microbial modulation (Guyader et al., 2015). Conversely, combinations involving ionophores and 3-NOP in concentrate diets may yield no additive effect, reflecting overlapping mechanisms (Romero-Perez et al., 2015). Thus, optimal additive combinations should be matched to diet type to maximise mitigation efficiency without impairing animal performance.

Systems-level mitigation: feed formulation, grazing management, and precision nutrition.

Methane mitigation in ruminant systems requires a holistic or systems-level approach, integrating nutritional, management, and technological strategies. Beyond individual feed additives, feed formulation, grazing management, and precision nutrition can collectively optimise rumen function, reduce methanogenesis, and improve productivity efficiency per unit of emission (Beauchemin et al., 2020). By improving feed conversion efficiency, animal productivity, and resource use, total methane output can be reduced even where absolute emissions per animal remain similar. Integration across scales—from rumen microbiome to whole-farm management—is essential to achieve sustainable, long-term mitigation compatible with animal welfare and profitability (Arndt et al., 2022).

1.5. Challenges and knowledge gaps

Microbial adaptation and long-term efficacy.

Many additives (EOs, saponins, 3-NOP, ionophores) show rapid methane reductions in short trials, but sustained efficacy over months or entire production cycles is poorly documented (Hegarty et al., 2021). The rumen microbiome is resilient and can adapt by selecting for resistant taxa or alternative H₂ pathways (Beauchemin et al., 2020). Microbial shifts and adaptation over time remain a major concern. Optimal dosing schedules, carrier matrices, and interactions with feed processing that influence persistence in the rumen are under-researched. Also, for some compounds (e.g., volatile essential oils), loss during feed processing or via rumen turnover reduces long-term exposure.

Variability among animal species and production systems.

The efficacy of methane-mitigating additives varies widely among animal species, diets, and production systems. Because methane production is closely linked to feed intake behaviour and diet composition, factors such as feed quantity, meal size, and feeding frequency can markedly influence

additive performance (Hristov et al., 2025). Significant differences in both methane yield and the rumen microbiota responsible for methanogenesis have been observed between animals consuming high-concentrate diets and those maintained on forage-based rations (Miller et al., 2023). Much of the existing evidence is derived from controlled studies in sheep or dairy cattle, which may not reflect responses under extensive grazing or in beef production systems (Hegarty et al., 2021). Moreover, breed-related differences in rumen physiology and feeding behaviour may further alter additive responses (Hegarty et al., 2021).

Rumen microbial composition also shifts with age and physiological stage, influencing methane output and responsiveness to additives (Miller et al., 2023). Despite its potential importance, few studies have investigated whether early life antimethanogenic interventions can produce lasting reductions in methane emissions. Evidence suggests that supplementation initiated soon after birth could influence rumen microbial development and lower methane production later in life (Yáñez-Ruiz et al., 2015; Beauchemin et al., 2020). Key knowledge gaps remain in understanding additive responses across beef versus dairy systems, grazing versus confined feeding, and across breeds and developmental stages.

Safety, residue, and consumer perception.

Some feed additives can negatively affect feed intake or nutrient digestibility—for example, high levels of tannins or lipid inclusion may reduce fibre degradation or alter milk composition. However, systematic evidence linking methane mitigation to animal performance across diverse production contexts remains limited. For synthetic inhibitors and concentrated plant extracts, data on residue occurrence in milk, meat, or offal, as well as their metabolic fate, are incomplete (Ku-Vera et al., 2020), and maximum residue limits have not been fully established for many compounds. Ionophores, widely used as feed additives in several countries, pose additional challenges due to concerns about antimicrobial resistance and differing national regulations, contributing to uncertainty and consumer distrust (Ahmed et al., 2024). Furthermore, transparent information regarding additive safety, labelling, and environmental benefits is often lacking, and consumer willingness to pay for verified methane-mitigation claims remains poorly understood.

Regulatory and policy barriers.

Even where efficacy is established (3-NOP has strong evidence), the regulatory pathways are complex and vary by country (Hegarty et al., 2021). For many additives (plant metabolites, alternative hydrogen sinks), evidence is weaker, especially for field implementation. There is a lack of harmonised measurement, reporting and verification (MRV) systems for methane mitigation in livestock. Additionally, many grazing systems (extensive) have little on-farm evidence, limiting scale-up.

Additives hold substantial potential to reduce enteric methane, but their role at scale is constrained by microbial adaptation, context dependency (diet, species, system), safety/residue questions and fragmented regulatory and market mechanisms. Prioritising long-term, mechanistic, and system-level research together with harmonised regulatory pathways and credible MRV will be essential to move from promising trials to safe, repeatable, and widely adopted mitigation strategies.

1.6. Future directions

Genetic and microbiome-based approaches.

Combining host genetics and rumen microbiome selection offers a promising pathway for achieving permanent and cumulative reductions in methane emissions by breeding low-emitting animals and promoting the establishment of beneficial microbial communities. Recent genomic–microbiome

association studies have highlighted the potential of integrating host genetic markers with microbial traits to enhance methane mitigation outcomes, although large-scale, multi-breed validation trials are still required to confirm the heritability and stability of these traits (Roques et al., 2024). Moreover, evidence suggests that early-life interventions -such as inoculating young ruminants with desirable microbial consortia or exposing them to low-emission microbiomes soon after birth - could facilitate the stable colonisation of beneficial microbes and reinforce the host's genetic predisposition toward lower methane production (Meale et al., 2021; Pérez-Enciso et al., 2021). Such strategies may help establish a persistent low-emission phenotype from the earliest stages of rumen development, complementing genetic selection approaches and increasing the long-term effectiveness of methane mitigation programs (Beauchemin et al., 2022).

Precision feeding technologies.

Sensors, automated feeders, and precision-livestock-farming systems can tailor additive dosing and diet composition to individual animals or cohorts, improving efficacy and cost-efficiency while minimising negative production trade-offs (van Empel et al., 2016; Morey et al., 2023). By continuously monitoring feed intake, rumen function, and animal performance, precision feeding technologies enable dynamic diet adjustments that optimise nutrient use and reduce excess fermentable substrates (Hristov et al., 2025), thereby limiting hydrogen production and subsequent CH₄ formation. These systems can also ensure the precise and consistent delivery of methane-mitigating additives at effective doses, preventing under- or over-supplementation. Furthermore, real-time data integration from sensors tracking rumen pH, gas emissions, and feed efficiency allows for early detection of inefficiencies or metabolic imbalances that could elevate CH₄ output (Morey et al., 2023). Collectively, these technologies enhance feed conversion efficiency, lower methane yield per unit of intake, and support adaptive management for sustained mitigation. Field studies and reviews increasingly identify precision-livestock-farming systems as a practical near-term strategy to scale methane reduction across commercial production settings (Papakonstantinou et al., 2024).

Integration with carbon credit and sustainability frameworks.

Linking verified on-farm methane reductions (e.g., from additives like 3-NOP/Bovaer) to credible carbon markets or in setting programs can create revenue streams for producers, but requires harmonised MRV, conservative baselines and robust regulatory oversight. Policy briefs and market analyses stress the need for science-based accounting to avoid double-counting. In this context, molasses-based lick block technologies could play a valuable role by enabling the practical deployment and quantifiable delivery of methane-mitigating additives under grazing conditions. Because lick blocks can provide consistent, traceable additive intake and are easily monitored at the herd and paddock level (Imaz et al., 2020), they may facilitate more reliable estimation of emission reductions, improving eligibility for carbon credit schemes in extensive ruminant systems (Callaghan et al., 2020). Integrating such feed-delivery strategies within verified emission-reduction methodologies could therefore enhance both the scalability and credibility of livestock-based mitigation initiatives.

1.7. Conclusion

Enteric methane mitigation remains a critical challenge for achieving climate-neutral livestock production. This review highlights that a wide range of dietary additives—plant secondary metabolites (tannins, saponins, and essential oils), lipids and fatty acids, electron acceptors, and synthetic compounds such as 3-nitrooxypropanol (3-NOP) and ionophores—can effectively reduce ruminal methane emissions through diverse mechanisms. However, their efficacy is influenced by

diet composition, animal species, production system, and duration of use, with long-term adaptation of the rumen microbiome remaining a major uncertainty.

Among the evaluated options, 3-NOP and lipid supplementation currently show the most consistent methane-reducing potential without major negative effects on productivity. This is reflected in recent policy developments, such as Denmark, where regulatory frameworks from 2025 require larger dairy farms to adopt methane-reducing feed strategies—including 3-NOP additives and increased dietary fat—as part of the country’s drive to meet ambitious greenhouse-gas targets (Landbrugsstyrelsen, 2024). In comparison, saponins, essential oils, tannin and lipid supplementation offer moderate but context-dependent benefits. Emerging strategies such as microbiome-based interventions, probiotics, and enzyme supplementation show promise but require further validation under field conditions (Yu et al., 2021).

Future progress will depend on integrated, systems-level approaches that combine additive use with optimised feed formulation, precision nutrition, improved animal genetics, and grazing management (Vargas et al., 2022). Sustained investment in long-term, multi-system research, harmonised regulatory frameworks, and credible carbon-accounting mechanisms is essential to ensure that methane mitigation strategies are both effective and sustainable, contributing to resilient and climate-smart livestock systems (Papakonstantinou et al., 2024).

In addition to the dietary strategies discussed throughout this review, the practical delivery of methane-mitigating compounds remains a key consideration for effective implementation under commercial conditions (Beauchemin et al., 2022). Among the emerging approaches, the use of molasses-based lick blocks represents a promising strategy to facilitate the controlled intake of feed additives in grazing ruminants (Imaz et al., 2020; Hristov et al., 2025). Lick blocks serve as a convenient and cost-effective vehicle for the sustained release of these compounds helping to ensure a consistent supply and uniform consumption among animals. This delivery method is particularly advantageous in extensive production systems, where conventional feed supplementation is often impractical (Beauchemin et al., 2020). By integrating nutritionally balanced ingredients with targeted additives, lick blocks can enhance both methane mitigation efficacy and animal productivity while supporting ease of adoption at the farm level (Hristov et al., 2025). Building upon the mechanistic understanding and safety evaluation of dietary compounds outlined in this review, the following chapter forms the experimental foundation of this thesis, presenting an *in vitro* and *in vivo* assessment of molasses-based lick block formulations containing methane-reducing additives.

Table 1. Safe dietary compounds for mitigating enteric methane in ruminants: a comparative overview

Compound/Class	Mode of action	Typical CH ₄ reduction (%)	Experimental system	Advantages	Limitations / risks	References
Tannins (condensed, hydrolysable)	Bind proteins; inhibit methanogens and protozoa; reduce H ₂ availability	10–30%	In vivo (cattle, sheep) and in vitro	Naturally occurring; improve protein utilization	May reduce fibre digestibility and intake at high doses	(Patra and Saxena, 2009; Jayanegara et al., 2012)
Saponins	Defaunation (protozoa reduction); disrupt microbial membranes	5–25%	In vitro and in vivo	Plant-derived; potential health benefits	Microbial adaptation; variable effects	(Pen et al., 2006; Goel and Makkar, 2012)
Essential oils (thymol, eugenol, cinnamaldehyde)	Antimicrobial activity; shift VFA pattern toward propionate	5–20%	Mainly in vitro; some in vivo	Multifunctional; natural preservative effect	Adaptation and inconsistent responses	(Calsamiglia et al., 2007; Benchaar et al., 2008)
Fumarate / malate	Alternative H ₂ sink via propionate formation	5–15%	In vitro and in vivo	Safe organic acids	Limited effect in vivo; cost	(Ungerfeld et al., 2007)
Lipids / fatty acids (e.g., coconut oil, linseed oil)	Suppress methanogens; reduce fibre digestion	10–30%	In vivo (dairy and beef cattle)	Improve energy density	Depress fibre digestibility; possible milk fat depression	(Machmüller et al., 2003; Beauchemin et al., 2008)
3-Nitrooxypropanol (3-NOP)	Inhibits methyl-coenzyme M reductase in	20–60%	In vivo (cattle, sheep); supported	Proven efficacy; approved additive (Bovaer®)	Cost; stability under storage	(Hristov et al., 2015; Melgar et al., 2020)

Compound/Class	Mode of action	Typical CH ₄ reduction (%)	Experimental system	Advantages	Limitations / risks	References
	methanogens		by in vitro studies			
Ionophores (Monensin)	Shift rumen fermentation toward propionate	5–10%	In vivo (feedlot cattle, dairy cows)	Improves feed efficiency	Reduced long-term effect due to adaptation; antibiotic concerns	(McGinn et al., 2004; Guan et al., 2006)
Probiotics / Direct-fed microbials (<i>Propionibacterium</i> , <i>Megasphaera elsdenii</i>)	Compete for H ₂ via propionate pathway	5–15%	In vivo (rumen and feedlot trials)	Can enhance gut health	Variable persistence; diet-dependent	(Newbold et al., 2015)
Plant oils / extracts (emerging)	Disrupt methanogens; redirect H ₂	5–25%	In vitro and pilot in vivo	Natural and renewable	Inconsistent efficacy	(Hook et al., 2010)

Chapter 2: Assessment of molasses products containing additives to reduce methane emissions from beef cattle

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Assessment of molasses products containing additives to reduce methane emissions from beef cattle

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Methane (CH₄) emissions from ruminants significantly contribute to agricultural greenhouse gases. Strategies to reduce these emissions without compromising productivity are urgently needed. This study evaluated the effectiveness of a novel product incorporated into molasses containing additives and grain-based pellets to reduce CH₄ emissions and improve productivity in beef cattle. Four products were screened *in vitro* for their efficacy in reducing CH₄ production. The most effective (Product 3) was then incorporated into grain-based pellets and offered to steers via GreenFeed units at target intakes of 0, 100, and 200 g/head per day. Dry matter intake (DMI), CH₄ production, CH₄ yield (g CH₄/kg DMI), CH₄ intensity, water intake, and average daily gain (ADG) were measured. Product 3 reduced *in vitro* CH₄ production by over 90% at a 6% inclusion rate ($P < 0.05$). *In vivo* study, steers supplemented with product-containing pellets tended ($P=0.07$) to reduce CH₄ production by 10.7% compared to the control. Methane yield was reduced ($P < 0.01$) by 11.7% and 7.5% for the 100 and 200 g/day supplementation levels, respectively. Methane intensity tended ($P=0.06$) to be 16.8% lower in supplemented groups. Product intake was negatively correlated with CH₄ yield and intensity and positively associated with water intake and ADG ($P < 0.05$). The inclusion of Product 3 in supplements beef cattle fed high forage diet significantly reduced methane yield and tended to reduce CH₄ production and intensity, without negatively impacting performance. Product 3 shows potential as a viable CH₄ mitigation strategy for high forage diet systems. However, further research is needed to assess the long-term effects and potential microbial adaptation associated with higher or prolonged dosing.

KEYWORDS

enteric methane, greenhouse gas, cattle, molasses, pellet supplementation

1 Introduction

Agriculture is responsible for 19.1% of Australia's total greenhouse gas (GHG) emissions, with livestock methane (CH₄) emissions contributing to 57% of the agriculture sector, approximately 10% of the country's overall emissions (Department of Climate Change, 2022). The vast Australian landscape spans 769 million hectares that vary in soil and plant types as well as temperature zones. This diverse environment supports the grazing of approximately 97% of Australia's beef cattle. Due to these environmental differences, there are variations in feed quality and availability across regions, which affect animal performance throughout the year. For instance, in certain regions, forages lack energy, crude protein, vitamins, and/or minerals (Dixon et al., 2020), leading to inefficient nutrient utilization (Leng, 1990) and, eventually, low animal performance. To address these challenges, improving nutritional management through efficient forage use has been identified as a viable strategy for reducing GHG emissions (Eugène et al., 2021). Improving the quality of the pasture is one of the most common approaches, however, it is not always easy or economically feasible to achieve. This context underscores a strategy for reducing the amount of CH₄ produced per unit of feed consumed, thereby mitigating GHG emissions through dietary supplementation of grazing animals. Numerous studies have explored the supplementation of additives to ruminant diets, aiming to decrease enteric CH₄ production. For instance, 3-NOP supplementation has been shown to reduce ruminant CH₄ emissions ranging from 18 to 39% (Almeida et al., 2021). *Asparagopsis* included from 0.05%-3% of OM intake could reduce methane emissions between 9 and 98% (Wasson and Hristov, 2022). However, the implementation of additives as a mitigation strategy is currently challenged by multiple factors including consistency of results, lack of information of responses in different production systems, cost-benefit, price of additives, and policies and legislation, amongst others. For example, inconsistencies in CH₄ reduction across a range of studies, negative impacts on other fermentation parameters or DMI and challenges in integrating them into practical feeding regimes, particularly in grazing systems (Patra et al., 2017; O'Reilly et al., 2021).

Research has highlighted the potential of essential oils (EO) to modify rumen microbial fermentation and boost ruminant feed efficiency (Calsamiglia et al., 2007). Those are typically obtained from plants through steam distillation or solvent extraction techniques (Simon et al., 1990; Greathead, 2003) although the plant material can also be fed and included in diets (Ku-Vera et al., 2020). Essential oils have anti-nutritional and antimicrobial characteristics to protect plants, prevent herbivory and attract pollinators, and can also assist in modifying the rumen microbiome to reduce the production of CH₄ (Patra et al., 2017; Garcia et al., 2020; Honan et al., 2021). In addition, the combinations of various EO may also increase microbial efficacy due to additive or synergistic effects between EO components (Benchaar and Greathead, 2011). Despite these promising findings, integrating EO into grazing systems remains a challenge.

Lick blocks (LB) technology could deliver anti-methanogenic additives in grazing animals, which is traditionally used as a livestock strategic supplementation around the world to provide energy, protein, minerals, or therapeutic substances to improve animal performance (Makkar et al., 2007). The LB is practical to deliver, easy to manufacture and store, capable of containing feed additives with anti-methanogenic properties, and economically feasible (Imaz et al., 2019). The objective of the present study was to evaluate LB formulation to reduce CH₄ emissions and enhance the performance of beef cattle fed high forage diet. An important aspect of LB is that the ingredients can be tailored for different regions, animal requirements, seasons, and production systems. Four proprietary formulations were initially evaluated *in vitro*, and the best candidate was tested *in vivo*.

2 Material and methods

2.1 Feed formulation

The products that have been used in the experiments of the present study comprised proprietary mixes of natural products and essential oils sourced from the *Myrtaceae*, *Rutaceae*, *Poaceae*, *Amaryllidaceae*, and *Lamiaceae*. The formulation is proprietary protected (Australian Provisional Patent Application No. 2023902943). All compounds are used in the human food chain as ingredients and are generally regarded as safe and the rate of feeding does not differ from concentrations used in food manufacturing. Free ranging cattle and sheep browse plants that contain the bioactive compounds identified in the products from the *Myrtaceae*, *Rutaceae* and *Poaceae*. The product is manufactured in accordance with excluded nutritional or digestive (ENDS) products and the Australian Stockfeed Regulation (Authority, 2024).

2.2 *In vitro* study

The *in vitro* trial was conducted to screen potential products for reducing CH₄ emissions. A commercial LB manufacturing company (AgCoTech Holdings Pty. Ltd., Crestmead, QLD, Australia) provided four different proprietary products that were assessed at different concentrations. The products evaluated in this project contain natural plant ingredients, including vegetable oil, essential oils, condensed tannins, and tea saponins. The DM and ash results of the 4 products is presented in Table 1.

All animals were cared for in accordance with the guidelines of the University of Sydney Research Integrity Animal Ethics Committee (approval #2022/2180).

Three cannulated adult non-lactating crossbred beef cows fed with grass hay (13% CP; 60% aNDF) at maintenance level were employed for rumen fluid collection. These collections took place between two and three hours after feeding in the morning. Rumen fluid samples were taken from the dorsal, anterior ventral, medium ventral, posterior dorsal, and posterior ventral rumen areas and put

TABLE 1 Dry matter and ash content for products 1, 2, 3 and 4 used to manufacture molasses products containing additives assessed *in vitro* for their anti-methanogenic properties.

	Dry matter content, % as-fed	Ash, %DM
Product 1	49.7	41.5
Product 2	33.9	10.3
Product 3	30.1	44
Product 4	53.5	40.9

into a prewarmed thermos filled to the lid to eliminate air in the headspace and immediately transported to the laboratory. The fluid was not purged with CO₂ because there was no headspace after closing the lid. Rumen fluid was anaerobically transferred into a 2-L flask mixed with two parts of a phosphate buffer and kept at 39°C until *in vitro* incubation (Forwood et al., 2019).

The incubations design was a completely randomized design (CRD) with three incubation bottles (replicates) per treatment per run and two incubation runs. The base substrate for the incubation was a ratio of 100% grass hay (60% aNDF). Products 1, 2, and 4 were dried in a forced air oven at 55°C for 72h and grounded to 1-mm. Product 1 and 2 powders were added into F57 ANKOM filter bags (Ankom Technology, Macedon, NY USA) in 0, 5, 25, 50, 10 and 200 mg, corresponding to concentrations of 0, 1, 5, 10, 20, and 40% respectively, replacing the substrate to make up 0.5g DM in total. The product 4 powder was added in 0, 50, 100, and 200 mg, corresponding to concentrations of 0, 10, 20, and 40%, respectively, replacing the substrate to make up 0.5g DM. Product 3 was evaluated in the original liquid form and was added to the substrate bag containing 0.5g DM before the addition of rumen inoculum in 0, 0.5, 1.0, and 1.5 mL corresponding to 0, 2, 4, and 6% concentrations of total volume (rumen fluid and buffer). All treatments were incubated in triplicate bottles using the batch fermentation method described in (Meale et al., 2012). Each run also contained three blank bottles with no substrate or treatments. On the incubation day, amber serum vials were warmed in the incubator at 39°C for 60 minutes and gassed with CO₂ before adding 25 mL of a 2:1 mixture of buffer saturated with CO₂: rumen fluid (Forwood et al., 2019), after that process, the bottles were immediately closed with rubber stoppers and placed in an incubator set at 39°C with a rotatory shaker at 120 oscillations per min.

Batch culture bottles were removed from the incubator after 24h. To determine CH₄ concentration at that timepoint, 25 mL of fermentation gas was collected from the headspace of bottle using a syringe and transferred into a 12 mL evacuated extainer for later measurement by gas chromatography. A water displacement apparatus was used to measure the remaining gas in the bottle. Total gas production was calculated by adding the 25 mL of CH₄ sample to the reading from the water displacement instrument (Fedorah and Hruday, 1983). Then serum bottles were opened and immediately placed on ice to stop microbial fermentation, and pH was immediately tested (Activon Model 209, Gladesville, NSW, Australia). A 1.5 mL subsample of each bottle culture was placed into a 2 mL Eppendorf tube with 300 mL of metaphosphoric acid

(20% w/w) and refrigerated at -20°C for subsequent volatile fatty acid (VFA) measurement following procedures described by (Wang et al., 2000; Chaves et al., 2006). ANKOM bags were removed from bottles, washed twice using a washing machine at short cycle (10 minutes), and cleaned to remove culture media, then dried for 24h at 60 °C for the estimation of *in vitro* dry matter digestibility (IVDMD).

2.3 *In vivo* study

Animal ethics approval was obtained from the Animal Ethics Committee of The University of Sydney (Approval # 2293). This animal feeding trial was carried out with the product that had the greatest reduction in CH₄ production from all products tested in the *in vitro* trials.

2.3.1 Animal feeding

For 70-d period, a completely randomized design (CRD) experiment was conducted using 45 Angus steers with an initial live weight (LW) of 233 ± 41.3kg (mean ± SD). The cohort included 36 Angus and 9 Angus × Charolais steers. The 45 steers were stratified into three LW strata (15 steers each), and within each stratum, steers were randomly assigned to three groups of 15 steers each. Upon arrival, all steers were vaccinated against BVD virus, major clostridial diseases (Ultravac[®] 5 in 1, Zoetis Australia Pty Ltd., Rhodes, New South Wales, Australia), and BRD (Bovilis[®] MH + IBR, Intervet Australia Pty Ltd., East Bendigo, Victoria, Australia). The animals were fed an oaten hay-based diet for two weeks before the treatments were introduced (as an adaptation period) and throughout the entire experiment. Each of the 3 treatments had 15 animals assigned to it: 1) control group (supplemented with pellets without product), 2) pellets containing the product for a target intake of 100g of product per head per day and, 3) pellets containing the product for a target intake of 200g of product per head per day treatment of the product incorporated into pellets. All pelleted sources were delivered through the GreenFeed[®] system (C-Lock, 2023).

The three treatment groups were split into two pens (4m length × 20m width) groups, with one pen occupied by control and a second pen occupied by 100g and 200g treatments. The control pen was equipped with 4 and the treatment pen had 6 electronic feeders (Intergado[®] Ltd, Contagem, Minas Gerais, Brazil), to deliver the hay and measure individual feed intake. Each pen had a water trough with two cattle weighing scales measuring individual LW and estimating water intake from the change in weight of the animals while drinking water (Intergado[®] Ltd, Contagem, Minas Gerais, Brazil) and a GreenFeed[®] system with dual hoppers (C-Lock, 2023) to measure individual daily CH₄, H₂, and CO₂ production rate (g/d) and O₂ consumption rate (g/d), and to deliver the target dose of pellet per animal. Electronic feeders are mounted on load cells to measure individual feed intake and are equipped with a radio-frequency identification (RFID) reader to detect the presence of an animal by recording its electronic identification (EID) number. The electronic feeders automatically assigned the amount of feed

consumed to the unique EID number present at the feeder when the feed weight decreased (Chizzotti et al., 2015; Oliveira et al., 2018). Samples of the offered feeds were collected weekly and dried at 70°C for 24h for the determination of DM content. Animals were taken to the yard and weighed at the start of the experiment, at 14-d intervals throughout the trial, and on the last day of the experiment. The average daily gain (ADG) was calculated by dividing weight gain (Final– Initial weight) by 56 (days of the experiment).

2.3.2 Pellet manufacturing process

Pellets were manufactured in a low-temperature process to preserve the active ingredients of the product. Rice bran (66.7%) and canola meal (33.3%) were finely ground and proportionally blended with the liquid product to achieve the designated inclusion levels (0, 6.28%, and 11.74%) without adding the settling agent. Mechanical compression was used to form pellets via a 4.4mm × 12mm die with the maximum temperature maintained below 54°C. Following this, the pellets were air-dried and cooled under ambient conditions to reduce moisture content, improve pellet durability, and minimize volatilization of the active ingredients.

2.3.3 Product supplementation

The target intake was 0, 100, and 200g of product/d into a pellet amount of 1.5 kg/head.d. One ton of pellets was proportionally mixed with 0, 67, and 133kg of the liquid product, respectively. These pellets were fed at 1.5kg (0g product), 1.6kg (100g product), and 1.7kg (200g product) per head per day.

The GreenFeed system was configured to provide the target amount of pellets to entice the animals to visit the GreenFeed multiple times per day (C-Lock, 2023). The double hopper GreenFeed in pen with the two treatments provided the two doses of product incorporated into the pellets depending on the treatment allocated to the RFID identified by the system. The pellets were dropped into a tray in a semi-enclosed hood when the RFID tag of an animal was present within the hood of the GreenFeed. The visit frequency (visits/d), number of feed drops (drops/visit), and the interval between visits were adjusted to ensure that each steer across all groups received the targeted intake as follows:

Control: visit frequency = 6 visits/d (spaced at a minimum of 120min intervals); number of cups dropped/visit = 5 (at 40 s intervals); average cup drop = 42g.

100 g per head per day: frequency = 4 visits/day (120min interval); feed drops/visit = 6 (40 s interval); average cup drop = 70g.

200 g per head per day: visit frequency = 6 visits/d (spaced at a minimum of 120min intervals); number of cups dropped/visit = 7 (40 s interval); average cup drop = 41g.

2.3.4 Quantification of CH₄ emissions

Individual CH₄ emissions measured by the GreenFeed system are expressed as daily CH₄ production (g CH₄/d). Visit duration and the number of records per individual are critical for CH₄

measurement because CH₄ is typically belched at 40–120 s intervals (Hammond et al., 2016). The GreenFeed operation commences when the steer places its head inside the shroud (Hammond et al., 2015). Following this, the proximity sensor in the shroud will monitor the head position of the animal during each visit, which will also be used to dismiss all measures where animals stepped out from the GreenFeed. Air is continuously drawn through the shroud and past the neck of the animal at a precisely measured rate, and the CH₄ and carbon dioxide (CO₂) concentrations are quantified in the exhaust air stream (Velazco et al., 2016). As the GreenFeed system provides multiple short-term breath measures, 30 measurements over a two-week period (Koning et al., 2024) with a minimum of three min durations per visit are needed to achieve a minimal variance in CH₄ production rate per animal (Arthur et al., 2017). Data are logged and transmitted into the C-Lock Inc. data management system and can be downloaded through the C-Lock Inc. website interface (<https://greenfeed.c-lockinc.com>).

The chemical composition of the hay and treatment pellets is presented in Table 2, and the ingredients composition of pellets is shown in Table 3. Dry matter (DM) analysis was determined by oven-drying samples at 70°C for 48h, followed by hot weighing (Cunniff and Washington, 1995). Pellets were sent to be analyzed using AOAC methods of chemical analysis (Cunniff and Washington, 1995). Unless otherwise stated, all results are reported on a DM basis (Cunniff and Washington, 1995).

2.4 Compound odor analysis

Products compounds were identified through gas chromatography mass spectrometry (GC-MS) analysis of volatile compounds using a Gerstel Thermal Desorption Unit (TDU; Gerstel, Mülheim an der Ruhr, Germany). The samples were analyzed after drying for products 1, 2, and 4, and as received for product 4 as used for the *in vitro* studies. Milligram masses of each product were weighed into 200 µL micro vials, secured in glass thermal desorption liners and inserted into the TDU for analysis. When inserted in the TDU, the samples were purged with ultra-high purity helium (BOC Ltd, North Ryde, NSW, Australia) at 20°C for 1min to eliminate air from the sample and inlet. Samples were heated to 150°C by the TDU at a rate of 12°C/s with a helium flow of 75 mL/min. Thermal desorption (TD) products were carried by the helium through to a programmed temperature vaporization (PTV) inlet (CIS-4; Gerstel) installed in an Agilent 7890 GC (Agilent Technologies Pty Ltd, Mulgrave, Australia), which was used in solvent mode during the TD. The PTV inlet, containing a glass liner filled with Tenax TA, was held at 20°C during the TD using liquid CO₂ (BOC Ltd) as the cryogen. After 1min of TD, the CIS-4 was heated at 12°C/s to 300°C and held at that temperature for 5 minutes while the TD products were injected into the GC at either a 1:10 or 1:500 split ratio. TD products were separated on a HP5-ms capillary column (30m × 0.25mm × 0.25 µm film thickness; Agilent) for analysis by a mass selective detector (Model 5975C; Agilent). Ultra-high purity helium was used as carrier gas (flow rate through the HP5-ms column was 2.3 mL

TABLE 2 Chemical composition of Product 3 used to manufacture molasses products containing additives treatment pellets and oaten hay fed to growing steers during the *in vivo* trial.

	Unit	Product 3	Pellets			Oaten hay
			0 g	100 g	200 g	
Dry matter (DM)	%	38.4	91.3	92.4	90.3	92.0
Neutral detergent fiber (NDF)	% DM	NA	21.0	21.0	22.0	67.0
Acid detergent fiber (ADF)	% DM	NA	12.0	12.0	12.0	42.0
Crude protein (CP)*	% DM	14.4	23.1	22.8	22.7	5.9
Crude fat (ether extract)	% DM	25.3	15.7	15.5	15.3	–
Water soluble carbohydrates (WSC)	% DM	19.8	8.7	9.6	10.1	4.3
Dry matter digestibility (DMD)	% DM	NA	78.0	78.0	76.0	50.0
Inorganic ash	% DM	9.4	9.4	9.5	9.6	7.0
Organic matter (OM)	g/kg DM	906	906	905	904	930
Metabolizable energy (ME)**	MJ/kg DM	15.5	15.8	15.8	15.5	7.0
Total starch	% DM	NA	14.0	14.1	13.6	NA
Urea	g/100 g	2.79	<0.03	0.09	0.17	NA
Nitrate	mg/kg	29	13	42	46	NA
Aluminum	mg/kg	13	120	99	73	NA
Arsenic	mg/kg	<5	<5	<5	<5	NA
Boron	mg/kg	<4	10	11	10	NA
Calcium	%	0.66	0.30	0.37	0.44	NA
Cadmium	mg/kg	<0.2	<0.2	<0.2	<0.2	NA
Cobalt	mg/kg	0.52	0.20	0.21	0.24	NA
Chromium	mg/kg	0.23	0.67	0.62	0.54	NA
Copper	mg/kg	0.94	8.1	7.8	7.8	NA
Iron	mg/kg	55	200	160	150	NA
Potassium	%	0.70	1.6	1.6	1.7	NA
Magnesium	%	0.21	0.81	0.80	0.84	NA
Manganese	mg/kg	15	180	180	180	NA
Molybdenum	mg/kg	0.12	0.61	0.65	0.67	NA
Sodium	%	0.71	0.027	0.090	0.14	NA
Nickel	mg/kg	<0.7	0.91	0.88	1.0	NA
Phosphorus	%	0.031	1.9	1.8	1.9	NA
Lead	mg/kg	<2	<2	<2	<2	NA
Sulfur	%	0.12	0.34	0.33	0.34	NA
Selenium	mg/kg	<0.05	0.17	0.17	0.17	NA
Zinc	mg/kg	5.3	62	57	58	NA
Dietary cation-anion Difference (<i>calc</i>)	meq/kg	NA	200	200	220	NA

(Continued)

TABLE 2 Continued

	Unit	Product 3	Pellets			Oaten hay
			0 g	100 g	200 g	
Chloride	%	NA	0.028	0.13	0.22	NA
Monensin	mg/kg	<2	NA	NA	NA	NA

• *measured value from feed analysis using Dumas Combustion Method.

• **ME value for a liquid block was measured using bomb calorimeter; others were estimated using a following formula:

• ME = [(crude protein × 35) + (crude fat × 84.6) + (WSC × 35) kcal/kg] × 0.004184.

TABLE 3 Ingredients composition of control and treatment pellets used to manufacture molasses products containing additives fed to growing steers.

	Control	100 g	200 g
Daily allowance, kg per head per day	1.5	1.6	1.7
Rice, %	66.7	62.53	58.85
Canola, %	33.3	31.21	29.38
Product, % of total	0	6.25	11.76
Product, % in the pellet	0	6.66	13.33

min⁻¹. The initial oven temperature of the GC was 35°C, held for 5 minutes, then heated at a rate of 5°C/min to 160°C, then heated at a rate of 25°C/min to 300°C and held isothermally for 4.4min. The temperature of the GC-MS interface was 280°C, the MS ion source 230°C and the quadrupole 150°C. The detector, in electron impact mode (70 eV), scanned the range of 35–300 m/z. Operation of the

GC-MS was controlled by Agilent Chemstation (version E.02.01.117) and the TDU by Maestro (version 1.4.36.16; Gerstel).

The choice of 1:10 or 1:500 split ratio for a sample was determined by preliminary method testing the four products used for the *in vitro* study (Product 1, 2, 3 and 4), exhibiting significant quantities of VOCs. Hence, 2–5 mg of sample was used with a 1:500 split ratio. The remaining samples were injected at a 1:10 split ratio from 12–16 mg of sample.

Post-processing of the GC-MS data was performed as described in Lawson et al. (2020) except compound identification was made using the NIST14 database (Mikaia et al., 2014), and results were expressed as “present” or “not present”.

2.5 Statistical analysis

2.5.1 *In vitro* study

Data obtained from *in vitro* fermentations were analyzed as a completely randomized design using PROC MIXED of SAS software (9.4 version, SAS Institute Inc., Cary, NC), using treatment, and dose as fixed effects whereas the random effects were defined as the incubation run (experimental unit). The incubation run was employed as an error term to assess treatment effects on CH₄, rumen fermentation parameters, and *in vitro* dry matter digestibility (IVDMD). Treatment effects were determined by comparing treatment dose replicates means with control, using least squares mean linear hypothesis test (LSMEANS/DIFF) with

Dunnett adjustment, declaring significance at $P \leq 0.05$ and tendency at $0.05 < P \leq 0.10$.

2.5.2 *In vivo* study

Data from the GreenFeed system and Intergado[®] electronic feeders were statistically processed using R (R Core Team, 2023). Intake data from feeders was deleted if DMI that were above 3.5% of individual LW from that period. Methane emissions, LW and feed intake data were used to calculate average values for each animal throughout the 56d of the treatment period and data from the 30-d adaptation period were not used for analysis. There were 15, 13, and 14 steers for treatment 0, 100, and 200 g/d that had more than 30 measurements for the statistical analysis. Pearson correlation analyses were performed among CH₄ production and feed and product intake after the data was averaged for each across the treatment period (Charmley et al., 2015). Average daily product intake was calculated by multiplying the pellet intake measured by GreenFeed by the concentration of product in the pellet according to treatment. A CRD with ANOVA was used with the treatment group as the fixed effect and 45 animals as the experimental units. Differences between treatments across the groups were considered significant when $P \leq 0.05$ and tendency at $0.05 < P \leq 0.10$.

3 Results

The results from odor analysis of the four products (Table 4) are consistent with those commonly found in blends of plant secondary metabolites such as essential oils (EO). Product 1 contained 3-carene, eucalyptol, D-limonene, and p-cymene. Product 2 contained 2,3-butanediol, 3-carene, eucalyptol, D-limonene, b-longipinene, (1aR,7R,7aS,7bR)-1,1,4,7-tetramethyl-1a,2,3,5,6,7,7a,7b-octahydro-1H-cyclopropa[e]azulene, and 2,3-dimethyl-nonadecane. Product 3 analysis found the presence of 2,3-butanediol, D-limonene, and trans-1-methyl-4-(1-methylethyl)-2-cyclohexen-1-ol. Finally, the compounds found in product 4 were 3-carene, eucalyptol, and [1S, cis]-naphthalene,1,2,3,5,6,8a-hexahydro-4,7-dimethyl-1-[1-methylethyl].

3.1 *In vitro* study

The results of *in vitro* trial evaluating the four products are shown in Table 5. All variables were affected by treatment ($P < 0.05$). Product 1 improved IVDMD at concentrations of 5% and above ($P < 0.05$)

TABLE 4 Gas chromatography mass spectrometry (GC-MS) analysis of products 1–4 for odor compounds potentially reducing methane emissions in cattle.

Mean retention time	Product 1	Product 2	Product 3	Product 4	Compound ID	Compound class	Synonym
1.75	Not present	Not present	Not present	Present	Dimethyl sulfide	Organosulfur	–
13.79	Not present	Present	Present	Not present	3-Carene	Monoterpene	Bicyclo[4.1.0]hept-3-ene, 3,7,7-trimethyl-
14.71	Present	Present	Not present	Present	Eucalyptol	Oxygenated monoterpene	1,8-Cineole
15.02	Present	Present	Not present	Present	D-Limonene	Monoterpene	Cyclohexene, 1-methyl-4-(1-methylethenyl)-, (R)-
15.24	Present	Present	Present	Not present	p-Cymene	Alkylbenzene (related to monoterpenes)	Benzene, 1-methyl-4-(1-methylethyl)-
17.13	Present	Not present	Not present	Not present	Cyclohexene, 1-methyl-4-(1-methylethylidene)	Monoterpene	Terpinolene
27.78	Not present	Not present	Present	Not present	b-Longipinene	Sesquiterpene	(1S,2R,7S,8S)-2,6,6-Trimethyl-9-methylenetricyclo[5.4.0.0 ^{2,8}]undecane
29.93	Not present	Present	Not present	Not present	1H-Cycloprop[e]azulene, octahydro-1,1,4,7-tetramethyl-, [1aR-(1a,7a,7ab,7ba)]-	Sesquiterpene	(+)-Ledene
30.32	Not present	Present	Not present	Not present	Naphthalene, hexahydro-4,7-dimethyl-1-(1-methylethyl)-, (1S-cis)-	Sesquiterpene	d-Cadinene
35.16	Not present	Not present	Not present	Present	1-Hexadecanol	Alcohol	–
37.39	Not present	Present	Not present	Not present	Unknown	Very noisy spectra – column bleed	–

and did not affect other *in vitro* fermentation parameters ($P > 0.10$). Product 2 increased IVDMD by 10% and above whereas gas production increased, and pH decreased by 40% of the product only ($P < 0.05$). Product 3 was the most effective in all *in vitro* fermentation parameters, reducing CH_4 production by over 90% at 6% dose of the product compared to the control ($P < 0.01$), and decreasing pH at 2% to 6% concentrations. In addition, 6% concentration of product 3 also reduced VFA concentration and IVDMD ($P < 0.05$). The lowest dose of product 3 enhanced fermentation due to a 27% numerical increase of total VFA concentration and 55% ($P < 0.01$) increase of gas production compared to the control treatment. A dose of 4% v/v of product 3 reduced methane production ($P < 0.01$) without negatively affecting fermentation as suggested by no differences in gas production and VFA against the control ($P > 0.05$). The molar percentage of acetate was drastically reduced with high doses of product 3 whereas propionate and butyrate were increased ($P < 0.01$). Product 4 improved IVDMD at all the concentrations assessed ($P < 0.05$), but all the other parameters were not affected ($P > 0.05$). Butyrate molar proportion increased with all doses of product 3 whereas branch-chained volatile fatty acids (BCVFA) increased by 6% of product 3 compared to the control treatment ($P < 0.05$; Table 5). None of the other treatments affected butyrate or BCVFA ($P > 0.05$).

3.2 *In vivo* study

Table 6 presents the performance and CH_4 emissions of growing beef steers over the 56-d experiment. Total feed, hay, and pellet intakes were similar among treatments ($P \geq 0.28$, Table 6). In addition, ADG was similar for all dietary treatments ($P=0.35$). Product intake (g/d) was significantly different across the groups, with the greatest intake in the 200 g/d group ($P < 0.01$).

The number of cups delivered was the fewest for steers fed 100 g/d, whereas the GreenFeed average visit duration was shortest for steers in the control treatment ($P < 0.05$; Table 6). This resulted in no differences in total daily time with good data in the GreenFeed between 0 and 100 g/d ($P > 0.05$), but the 200 g/d had the longest daily GreenFeed data compared to the other treatments ($P < 0.05$; data not shown).

Results indicated a tendency for CH_4 production to decrease in the 100 and 200 g/d groups compared to the control treatment, with reductions of 10.7% and 6.3%, respectively ($P=0.07$; Table 6). Methane yield (g CH_4 /kg DMI) was lower ($P < 0.01$) in steers fed 100 and 200 g/d treatments, showing reductions of 11.7% and 7.5% compared to the control group (Table 6). Methane intensity (g CH_4 /kg ADG) also tended to decrease in the 100 and 200 g/d groups by 16.8% and 14.9%, respectively ($P=0.06$) compared to the control.

TABLE 5 *In vitro* gas production and fermentation parameters of four different candidate products for the manufacturing of lick blocks.

	Control	Product 1					Product 2					Product 3			Product 4			SEM	P-value
	0%	1%	5%	10%	20%	40%	1%	5%	10%	20%	40%	2%	4%	6%	10%	20%	40%		Treat
Gas, mL/g DM	85	82	85.8	87.3	84.4	78.8	91.8	93.5	98	105.7†	117.0*	131.8*	102.4	76.5	91.7	92.4	96.1	8.42	<0.01
CH ₄ , %	14.8	15.1	15.4	15.5	13.2	12.7	15.3	14.4	14.7	15.2	14.6	15.6	5.6*	1.2*	15.2	14.8	14.3	1.4	<0.01
CH ₄ , mL/g DM	12.7	12.4	13.3	13.7	11.1	10	14.2	13.5	14.4	16.1	17.2	20.7*	6.8*	0.92*	13.9	13.7	13.8	2.25	<0.01
pH	6.89	6.89	6.88	6.84	6.86	6.9	6.86	6.81	6.74	6.69†	6.54*	6.48*	5.84*	5.00*	6.79	6.81	6.70†	0.05	<0.01
Total VFA, mM	66.4	68.9	66.9	65.3	67.3	58	72.8	69	67.9	75.7	88.2	84.5	61.2	16.5*	69.8	79.9	75.3	6.14	<0.01
% of total VFA																			
Acetate (A)	65.2	64.8	64.5	65	65.4	66.5	65.5	63.5	64.5	64.6	62.7	59	65.3	3.2*	67.6	65.8	64.9	2.01	<0.01
Propionate (P)	21.3	21.4	20.6	20.4	20.6	19.3	21.3	21.4	22	21.5	22.8	12.0*	11.5*	42.5*	20.5	20	20.9	2.15	<0.01
Butyrate	9.1	9.4	9.3	9.8	9.6	9.6	8.7	9.6	9.7	10.1	10.4	23.8*	20.6*	45.6*	8.2	10.5	10.7	1.87	<0.01
BCVFA	2.53	2.51	3.1	2.75	2.48	2.65	2.56	3.04	2.2	2.19	2.24	2.63	1.49	5.06*	2.18	2.12	2.03	0.707	<0.01
Ratio A:P	3.06	3.03	3.14	3.19	3.17	3.44	3.08	2.97	2.93	3	2.75	4.92	5.68	0.08	3.3	3.28	3.1	1.1	N/A
IVDMD, %	51.7	51.4	54.1*	54.4*	57.7*	63.9*	51.4	53.7	56.6*	59.7*	66.9*	49.5†	48.9*	44.7*	57.0*	60.2*	69.3*	0.72	<0.01

CH₄, methane; IVDMD, *in vitro* dry matter digestibility; VFA, volatile fatty acids; BCVFA, branched-chain VFA (iso-butyrate + iso-valerate); SEM, standard error of the means.

*Means differ ($P \leq 0.05$) from control treatment, † means tend to differ from control treatment ($P \leq 0.10$).

TABLE 6 Animal performance and greenhouse gases (GHG) emissions of growing beef steers after being offered pelleted with product 3 for 56 days of the experiment.

	Treatment g Product per head per day			SEM	P-value
	Control	100 g	200 g		
Total DMI, kg/day	6.93	6.90	7.01	0.272	0.96
Hay DMI, kg/day	5.74	5.60	5.85	0.251	0.78
Pellet DMI, kg/day	1.20	1.29	1.15	0.588	0.28
Targeted pellet intake, kg/day	1.50	1.60	1.70	–	–
Product intake, g/day	0.00 ^a	92.3 ^b	172.5 ^c	4.99	<0.01
ADG, kg/day	0.71	0.78	0.78	0.04	0.35
GreenFeed cup consumption, n#/day	31.8 ^a	20.0 ^b	32.2 ^a	1.03	<0.001
GreenFeed visits, n#/day	4.42 ^a	3.26 ^b	4.40 ^a	0.203	<0.001
GreenFeed visit length, sec/visit	257.9 ^b	331.8 ^a	361.6 ^a	15.69	<0.001
Time of GreenFeed visit, h in 24 h	9.0	11.3	12.0	0.230	<0.001
GreenFeed Total time visits, h	30.8	23.4	31.4	1.58	<0.001
GreenFeed total visits	358	234	308	12.6	<0.001
CH ₄ production, g/day	159	142 [†]	149 [†]	5.10	0.07
CH ₄ yield, g/kg DMI	24.0 ^a	21.2 ^b	22.2 ^{ab}	0.55	<0.01
CH ₄ intensity, log (g/kg ADG)	2.35	2.27 [†]	2.28 [†]	0.03	0.06
CH ₄ intensity, g/kg ADG	223.9	186.2 [†]	190.5 [†]	1.07	0.06
CO ₂ production, g/day	4492	4424	4479	146	0.94
H ₂ production, g/day	0.217	0.236	0.225	0.01	0.42
O ₂ consumption, g/day	3258	3449	3484	119	0.35
Water intake, L/day	28.9 ^a	37.0 ^{bc}	40.3 ^{cd}	2.31	<0.01

DMI, dry matter intake; ADG, average daily gain; SEM, standard error of the means.

^{a,b}means with different superscript differ at $P < 0.05$.

[†]means tend to differ from control treatment ($P \leq 0.10$).

No significant differences were observed between the 100g and 200 g/d treatments for any variable ($P > 0.10$). Additionally, there were no differences among groups ($P \geq 0.35$) in CO₂ and H₂ production (g/d) or oxygen consumption (g/d) as measured by the GreenFeed unit.

Daily water intake increased with the incremental dose of product supplementation, showing 28% and 39% increases on the steers treated with the 100 and 200 g/d treatments compared to the control steers, respectively ($P < 0.01$).

The correlation matrix among selected variables is illustrated in Figure 1. Average product intake throughout the trial was positively correlated with ADG ($P < 0.05$) and water intake ($P < 0.001$). Importantly, the average product intake was negatively correlated with CH₄ intensity ($P < 0.01$) and CH₄ yield ($P < 0.05$) and positive with O₂ consumption ($P < 0.05$).

4 Discussion

The objective of the *in vitro* studies was to assess different formulations of products containing a range of essential oils. Some of these products are volatile as demonstrated in the odor analysis, and others can bind with sugar and lipid moieties. It is important to note that products 1, 2, and 4 were dried at 55°C to improve dose accuracy. In contrast, product 3 was used ‘as received’ because it could not be dried. Drying may have affected the activity or concentration of some essential oils, which prevents us from making a direct comparison between the products. However, the objective of the *in vitro* trial was to select the product with the greatest effect on CH₄ production. It is important to note that the analysis of volatile compounds of each product was done after drying products 1, 2, and 4 as used *in vitro*. In addition, both pellet

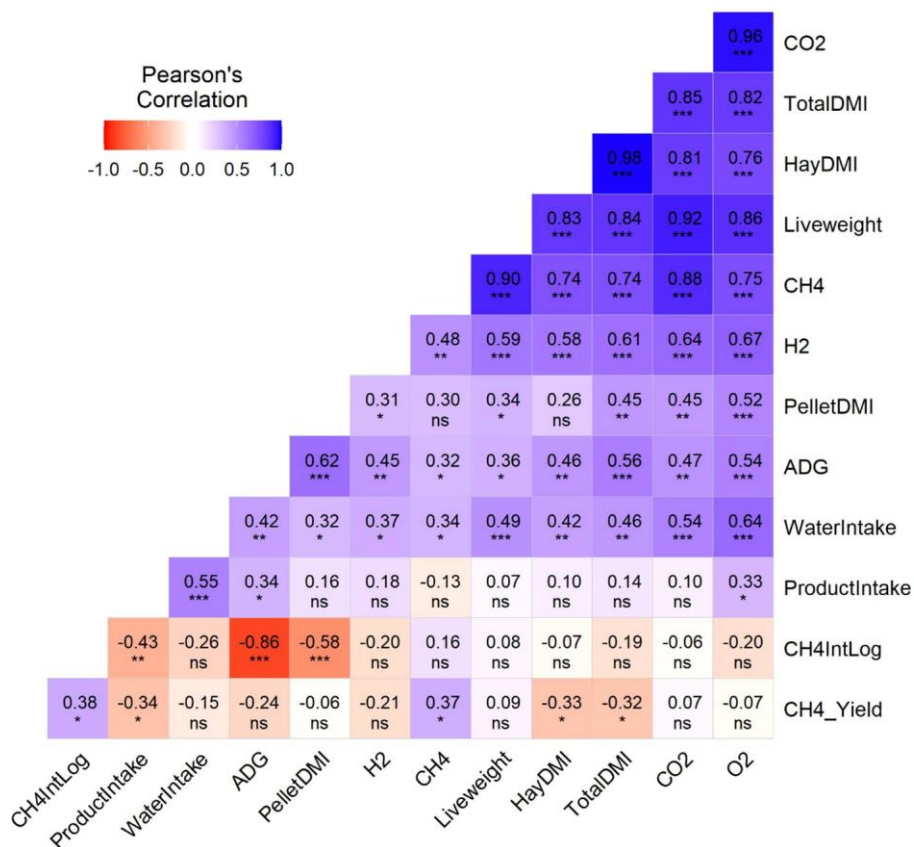


FIGURE 1 Correlation matrix of the performance and greenhouse gases (GHG) emissions of beef steers offered product supplement used to manufacture pellets with molasses containing additive during 56d of the experiment. ns P ≥ 0.05; *P ≤ 0.05; **P ≤ 0.01; and ***P ≤ 0.001. TotalDMI, Total Dry Matter Intake; HayDMI, Total Dry Matter Intake from Hay; PelletDMI, Dry matter Intake from Pellets; ADG, Average Daily Gain.

and lick block manufacturing (where the products are added) also involve increased temperatures. Furthermore, the ash content values obtained in this study are higher than typically expected for organic feed products. The exceptionally high ash content, particularly in Product 3 (44% DM), raises concerns about the composition of the material. While Mix 3 is described as a liquid-solid admixture, the ash content suggests a significant presence of inorganic material, potentially making it similar in composition to a clay slurry or mineral-rich mixture. Given this, further investigation into the mineral composition of the products may be warranted to confirm the nature of the ash content and its potential impact on fermentation dynamics.

Product 3 emerged as the standout performer in reducing CH₄ emissions *in vitro*, demonstrating a remarkable efficacy across all evaluated *in vitro* parameters. Notably, at a 6% dosage of product 3, it reduced CH₄ production by over 90% in comparison to the control group, highlighting its potent effect albeit the significant reduction in IVDMD. This product was then selected for the *in vivo* trial, where results indicated that steers treated with the product tended to have up to 10.7% reduction in CH₄ production. Moreover, CH₄ yield from steers treated with the product were up to 11.7% lower than those without the product in the pellet. Furthermore, CH₄ intensity tended to be 16.8% lower for the

animals fed Product 3 compared with the control, reflecting the efficiency of the product in reducing emissions.

Results from the volatile analysis showed compounds present in each product that are consistent with those commonly found in blends of plant secondary metabolites such as EO. Dimethyl sulfide was one of the compounds found to be present in Product 4, belonging to the organosulfur chemical group, which are common constituents of many plant species including those used in the present study and garlic oil (Honan et al., 2021). Organosulfur compounds have been suggested to reduce enteric CH₄ due to their toxic effects on methanogenic archaea (Soliva et al., 2011). In a study by Soliva et al. (2011), garlic oil was added to a mix of hay and soybean meal concentrate *in vitro* at a dose of 300 mg/L of incubation liquid and reported a 91% reduction in CH₄ production. Pawar et al. (2014) also screened garlic oil *in vitro* in a concentrate diet and reported reductions in CH₄ of up to 71.5% when included in a dose of 30 μL/mL of liquid incubation, though there was also a significant negative impact on overall fermentation noted. Patra and Yu (2015) used garlic oil in a forage-based diet *in vitro* and also reported a reduction in CH₄ but only by 29%, indicating that dimethyl sulfide may be more effective for use in concentrate diets compared to forage-based diets as used in the present study. However, Product 4 did not have a significant

reduction in CH₄ and IVDMD had a significant increase in the present study. These differences may be due to low concentration of the active ingredients in the present compared to previous studies, but this cannot be confirmed because dimethyl sulfide concentration was not measured, and it is also unclear whether this or other compounds present in garlic oil may be responsible for a reduction of CH₄ production.

Compounds belonging to the monoterpene and oxygenated monoterpene chemical groups were identified in all four products screened in the present study. As classes of secondary plant metabolites commonly found in EO, some oxygenated monoterpenes have been found to inhibit the metabolism and growth of microbes in the rumen (Garcia et al., 2020). Eucalyptol, an oxygenated monoterpene was identified in product 1, 2, and 4. Results from *in vitro* analysis of those products did not indicate any significant change in CH₄, pH and VFA in the present study which aligns with the results found in the literature. Eucalyptol oil was screened *in vitro* in a study by Colombini et al. (2021). It was added to a total mixed ration substrate in a dose of 200 mg/L of inoculum and incubated with measurements taken at 24 and 48h time points. The study found that there was no observable impact on CH₄ production at either time point, or on CH₄ concentration after 24h. However, there was a 7% reduction in CH₄ concentration after 48h (Colombini et al., 2021). The latter study did not observe any impact on pH or total volatile fatty acids (VFA) production, despite the changes in the relative abundance of rumen microbes. In particular, the Firmicutes-to-Bacteroidetes ratio was higher when eucalyptol was added compared to the control, a trend that has been previously associated with reduced feed conversion (Jami et al., 2014; Colombini et al., 2021). Colombini et al. (2021) did not measure IVDMD, but the reduced fermentation may have been a result of impaired microbial proportions and explained the reduction of CH₄ at a later time point.

Comparatively, the hydrocarbon monoterpenes, such as 3-carene or D-limonene identified in products 1, 2, 3, and 4, have been reported to increase rumen microbial activity in some cases (Kahvand and Malecky, 2018), which aligns with results from our experiments assessing product 3. For instance, the *in vitro* evaluation of product 3 showed a significant reduction of CH₄ at 4 and 6% v/v concentration, but no significant changes at 2% concentration. This could be attributed to an insufficient dose. Furthermore, feeding product 3 *in vivo* confirmed that CH₄ yield decreased and CH₄ production and intensity tended to decrease with product 3. In contrast, no change was observed on CH₄ production when product 2 was assessed *in vitro*, but there was an increase in gas production and IVDMD. It has been suggested that monoterpenes may be broken down and utilized as a source of carbon by rumen microbes (Benchaar et al., 2008; Garcia et al., 2020). This may explain the significant increase in IVDMD with increasing dose of products 1, 2, and 4 in the present study, all of which contained monoterpenes. Nonetheless, Product 3 also contained monoterpenes but showed a significant decrease in IVDMD especially when a greater dose was added. This decrease could be attributed to the reduction in pH which will be further explained in this discussion.

Sesquiterpene compounds were also found in Products 2 and 3. (Nunes et al., 2023) evaluated the effects of three EO from exotic plants at different levels, with *Hedychium gardnerianum* containing essential oils from the sesquiterpene family. Results from that study indicated that both gas and CH₄ production decreased significantly compared to the control treatment after incubation at medium and high but not at the lowest concentration. Similarly, a significant reduction in CH₄ production was observed at medium and high concentration of product 3, but gas production was not affected at medium and high concentrations and increased at the lowest concentration of product compared to the control treatment. Product 2 increased gas production by 28.9 and 37% at 20 and 40% of the substrate, although CH₄ was not affected. The variation in outcomes observed from differing treatments and concentrations indicates an interaction between the type of EO and its quantity, which affects both total gas and CH₄ production. The presence of active compounds within each EO, and their combination, can lead to distinct synergistic, antagonistic, or additive impacts on the results (Burt, 2004). Also, possible differences exist in rates at which microbial populations degrade these secondary metabolites (Broudiscou et al., 2007). Given these factors, additional studies are required to ascertain the degree to which the essential oils used in the present project can be degraded or metabolized within the digestive system of ruminants.

Previous research that has demonstrated successful CH₄ mitigation from EO supplementation suggests that a trade-off exists between CH₄ inhibition and overall fermentation (Durmic et al., 2014; Jahani-Azizabadi et al., 2019; Parra et al., 2023).

Although modifications in VFA profiles are often deemed beneficial, EO can sometimes lead to undesirable shifts in individual VFA proportions. For instance, Castillejos et al. (2006) observed that eugenol at 500 mg/L diminished propionate proportion without altering total VFA concentration. Moreover, Cardozo et al. (2005) demonstrated that the impact of EO on VFA profiles is contingent upon pH. The 6% concentration of Product 3 decreased the medium's pH to 5, seemingly at odds with the lower VFA concentration. These pH values are of particular concern as pH < 5.5 is considered within the threshold for subacute ruminal acidosis, and pH < 5.0 is considered acute ruminal acidosis (Penner et al., 2007). However, this can be attributed to the product's inherent acidity which was measured at 4.82 (data not shown), indicating that the low ruminal pH may be due to the increasing concentration of Product 3 instead of a result of fermentation products. It is important to point out that the dose rate used in the present *in vitro* study may not yield a significant pH change *in vivo* because the animal produces bicarbonate buffer and the dose is low compared to total rumen turnover and volumes. In contrast, 4% of product 3 reduced CH₄ by 62% without negatively impacting fermentation, and medium pH was 5.84. This underscores the significance of product 3 dose optimization and suggests that as an optimum concentration.

In the *in vivo* study, water intake increased with product 3 supplementation, potentially due to its higher mineral content, prompting animals to drink more to sustain high pellet intake. This phenomenon has been well-documented: cattle consuming

salt-enhanced supplements or saline water show increased water intake and enhanced renal diuresis (e.g., heifers consuming salty water exhibited water intake rising from 21.8 to 41.5 kg/day with increasing salt concentrations) (Visscher et al., 2013). Additionally, studies investigating the broader impacts of salt supplementation to low-quality forage diets found that increasing salt levels can alter forage intake, rumen fill, liquid kinetics, and fermentation patterns, further reflecting the physiological consequences of elevated salt intake (White et al., 2024). Despite this rise in water intake, no significant changes were noted in ADG or feed intake. This observation aligns with findings from various *in vivo* studies with EO fed to beef cattle where no effect was demonstrated on feed intake (Beauchemin and McGinn, 2006; Yang et al., 2010), suggesting no changes in rumen methanogenesis or adaptation of the microbes to EO. Cardozo et al. (2006) found that while a high dose of EO (a blend of 600 mg/day of cinnamaldehyde and 300 mg/d of eugenol) did not affect DMI, a lower dose (180 mg/d of cinnamaldehyde and 90 mg/d of eugenol) decreased DMI in beef heifers fed a high-concentrate diet. Similarly, Yang et al. (2010) reported that eugenol supplementation in growing heifers did not influence DMI. Conversely, Benchaar et al. (2006) observed that supplementation of 2–4 g/day of an EO mixture containing eugenol increased DMI in a digestion study, but this effect was not replicated in a growth study with beef cattle on a high-forage diet. Nonetheless, product 3 supplementation showed a trend for a reduction in CH₄ production and intensity, especially noticeable in 100 g/head.d and a significant reduction in CH₄ yield compared to the control group. Additionally, a negative correlation exists between the log of CH₄ intensity and ADG. This correlation may stem from the fact that Product 3 could be enhancing feed conversion, resulting in better weight gain without a proportional increase in methane production. Therefore, while the cattle gain more weight, the methane per unit of weight gain is reduced. These results indicate the potential for product 3 to contribute to lower CH₄ emissions from beef steers fed high-forage diets. The specificity of Product 3 in reducing CH₄ without affecting CO₂ and H₂ production further supports its targeted action, as corroborated in the literature on plant-based compounds, including EO, saponins, and tannins (Francis et al., 2002; Benchaar and Greathead, 2011; Wang et al., 2012; Honan et al., 2021).

5 Conclusion

This study evaluated four novel products *in vitro*, with Product 3 showing the strongest methane-reducing effect. At 6% (v/v), CH₄ production was reduced by over 90%, although digestibility was also compromised. At a lower inclusion level (4%), methane emissions were reduced by 62% without negatively affecting digestibility, emphasizing the importance of dose optimization.

In vivo, incorporation of Product 3 into beef steer diets modestly reduced methane yield (–11.7%) without affecting feed intake or growth performance. While these results are promising, they should be interpreted cautiously given the short duration and scale of the trial.

Further research at higher dosages, over extended periods, and with incorporation into lick blocks is needed to clarify the balance between methane abatement, digestibility, and overall animal performance.

Data availability statement

The raw data supporting the conclusions of this article are not publicly available because they are subject to confidentiality and contractual obligations. In accordance with the governing agreement, such data cannot be disclosed or distributed without the prior written consent of both Meat & Livestock Australia (MLA) and Agcotech. Requests to access the datasets should be directed to the corresponding author/s.

Ethics statement

The experimental protocols and use of animals were approved by the Animal Ethics Committee of the University of Sydney for both the *in vitro* (2022/2180) and *in vivo* (ARA No. 2023/2293) trials. The study was conducted in accordance with the local legislation and institutional requirements.

Author contributions

MN: Data curation, Formal analysis, Investigation, Methodology, Writing – original draft. GS: Conceptualization, Data curation, Formal analysis, Investigation, Methodology, Project administration, Supervision, Validation, Writing – review & editing. MP: Formal Analysis, Investigation, Methodology, Writing – review & editing, Data curation. LG: Conceptualization, Data curation, Formal analysis, Funding acquisition, Investigation, Methodology, Project administration, Supervision, Writing – review & editing. AC: Formal analysis, Investigation, Methodology, Supervision, Writing – review & editing.

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

The author(s) declared that they were an editorial board member of *Frontiers*, at the time of submission. This had no impact on the peer review process and the final decision.

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Chapter 3: General Discussion

This chapter synthesises the key findings of this thesis, integrating results from both in vitro and in vivo experiments to evaluate the efficacy of MLB formulations containing anti-methanogenic ingredients. The discussion highlights practical constraints and identifies areas for future research to optimise the use of MLBs as a mitigation strategy in pasture-based beef systems.

3.1 Main findings

Reducing methane emissions on pasture-based systems remains a significant challenge. The literature review of this thesis (Chapter 1) evaluated several methanogenesis-inhibiting feed additives, while those show promise, practical and scalable delivery methods for grazing cattle are still lacking. Attempts to prolong the effectiveness of rumen methane inhibition -for example, the use of 3-NOP (Muetzel et al., 2019) formulations have been explored. Research has also examined ways to improve the stability of volatile additives within feed (Magnusson et al., 2020). Despite these advances, available evidence supporting reliable delivery of these additives remains limited and insufficient for widespread adoption in grazing production system (Hegarty et al., 2021). Among the emerging solutions, molasses-based carrier systems have been proposed as a practical means of improving additive stability and intake control under grazing conditions (Imaz et al., 2020; Hristov et al., 2025).

Chapter 2 of the thesis presents an experimental evaluation of novel molasses-based products, formulated as additive carriers containing essential oils and plant secondary metabolites, rather than molasses lick blocks per se. Results from the in vitro screening identified one formulation (Product 3) as the most promising additive. At the highest inclusion level (6%), Product 3 reduced methane production by more than 90% compared with the control group, although this was accompanied by a significant reduction in in vitro dry matter digestibility (IVDMD). At a lower inclusion level (4%), methane emissions were reduced by 62% without negatively affecting digestibility, emphasising the importance of dose optimisation. This initial screening effectively differentiated products based on anti-methanogenic potential and guided the selection of Product 3 for subsequent in vivo testing.

In contrast to the strong in vitro inhibition, the in vivo responses were more moderate but directionally consistent. Supplementation with the Product 3-pellet did not significantly reduce methane production, but it achieved reductions in methane yield (up to 11.7%) and tended to decrease methane intensity (16.8%). These reductions occurred without negative effects on dry matter intake, water intake, or average daily gain, indicating that the supplement was well-tolerated and compatible with forage-based diets. The positive association between supplement intake and both water consumption and weight gain suggests potential production benefits or, at minimum, no trade-off in growth performance.

3.2 Practical considerations and limitations

While the present study demonstrates that Product 3 has potential to reduce enteric methane emissions in beef cattle, several factors limit the interpretation and extrapolation of these findings. These limitations relate primarily to (i) additive stability and microbial adaptation, (ii) duration of supplementation and implications for long-term mitigation, and (iii) constraints associated with methane measurement using GreenFeed system.

3.2.1 Additive stability and microbial adaptation

The strong inhibition of methane production observed in vitro contrasted with the more modest reduction recorded in vivo, reflecting the fundamental differences between these two experimental systems rather than an inconsistency in additive efficacy. In vitro techniques are primarily intended

as screening tools that allow rapid identification of compounds with antimethanogenic potential under tightly controlled conditions. These systems are closed, short-term incubations in which additives, particularly essential oils, tannins, and saponins, are typically applied at relatively high concentrations and experience minimal degradation or host-mediated modulation. As a result, *in vitro* assays often report pronounced methane inhibition that may overestimate responses achievable under practical feeding conditions (Patra and Yu, 2015; Beauchemin et al., 2020).

In contrast, *in vivo* experiments are designed to validate the efficacy of selected candidate (e.g.: Product 3) under biologically complex and dynamic conditions, where rumen fermentation is influenced by feed intake patterns, digesta passage rate, microbial adaptation, compound metabolism, and host-diet interactions. The *in vivo* dose response evaluation conducted in this thesis therefore represents a more realistic assessment of the mitigation potential of a single promising product identified during *in vitro* screening. Lower effective concentrations of EO-based additives *in vivo*, combined with microbial adaptation and potential detoxification mechanisms, are well documented to attenuate antimethanogenic responses relative to *in vitro* systems (Calsamiglia et al., 2007; Durmic et al., 2021). Consequently, the more moderate but production-relevant methane reductions observed *in vivo* are consistent with previous literature and reinforce the importance of interpreting *in vitro* findings as indicative rather than predictive of *in vivo* outcomes. Chemical characterisation of Products 2 and 3 indicated the presence of sesquiterpene compounds, a class of EO constituents previously associated with antimethanogenic activity (Nunes et al., 2023). In that study, sesquiterpene-rich EO extracted from exotic plants reduced total gas and methane production at medium and high inclusion levels, but not at low concentrations. A comparable dose-dependent response was evident in the present *in vitro* experiment. Product 3 significantly reduced CH₄ production at medium and high inclusion rates, whereas gas production was unaffected at these levels and increased at the lowest dose. In contrast, Product 2 increased total gas production substantially without altering CH₄ proportion, suggesting stimulation of fermentative activity without concurrent inhibition of methanogenesis. These divergent responses across products and inclusion rates indicate a strong interaction between EO composition and dosage, affecting both fermentation dynamics and methane output. The presence of multiple bioactive compounds within each EO blend may result in synergistic, antagonistic, or additive effects, leading to divergent outcomes depending on concentration and microbial context (Burt, 2004). Further variability likely arises from differences in the extent to which rumen microorganisms degrade, transform, or detoxify secondary metabolites (Broudiscou et al., 2007). Such processes may substantially reduce the persistence and bioavailability of EO constituents *in vivo*, thereby limiting their capacity to sustain methane inhibition. The absence of direct measurements of EO degradation or rumen metabolite profiles in the present study represents a limitation, as it constrains interpretation of whether reduced *in vivo* efficacy reflects chemical instability, microbial metabolism, or both.

Beyond chemical factors, the adaptive capacity of rumen microbial communities presents a critical challenge to the long-term effectiveness of antimethanogenic additives (Patra and Yu, 2012). Rumen microorganisms are highly resilient and capable of adapting to antimicrobial compounds through shifts in community structure, induction of detoxification mechanisms, and functional redundancy among hydrogen-utilising pathways (Calsamiglia et al., 2007; Morgavi et al., 2010). Such adaptations may progressively diminish the inhibitory effects of plant-based additives, leading to declining efficacy during prolonged supplementation (Hegarty et al., 2021). The 56-day duration of the present *in vivo* trial may therefore have captured only the early phase of response, with longer-term exposure potentially resulting in further attenuation of the methane-reducing effect. Indeed, previous studies have reported time-dependent declines in efficacy for essential oil-based and

tannin-based additives, highlighting the challenge of sustaining antimethanogenic activity *in vivo* (Durmic et al., 2014; Patra and Yu, 2015).

An additional limitation related to the narrow margin between effective methane inhibition and undesirable impacts on rumen function (Patra and Yu, 2012; Hristov et al., 2025). Studies reporting successful methane suppression with EO supplementation consistently emphasise a trade-off between antimethanogenic efficacy and overall fermentation (Durmic et al., 2014; Jahani-Azizabadi et al., 2019; Parra et al., 2023). Although alterations in VFA profiles are often interpreted as a favourable indicator of improved rumen efficiency, EO can also induce less desirable shifts depending on dose and rumen conditions. For instance, (Castillejos et al., 2006) demonstrated that eugenol at 500 mg/L reduced propionate proportion without affecting total VFA concentration, while (Cardozo et al., 2005) showed that EO effects on fermentation pathways are strongly pH-dependent, with divergent responses observed under acidic versus neutral conditions. In the present study, the highest inclusion (6%) of Product 3 decreased medium pH to approximately 5.0, a value within the threshold for acute ruminal acidosis (Penner et al., 2007). This appears inconsistent with the observed reduction in VFA concentration; however, this response is likely attributable to the inherent acidity of Product 3, which had a measured pH of 4.82 (data not shown), rather than to excessive fermentative acid production. It is important to recognise that this pH may overestimate the magnitude of response expected *in vivo*, given the substantial buffering capacity provided by salivary bicarbonate and dilution within the much larger rumen volume and continuous digesta turnover (Beauchemin, 2018). Nevertheless, this response illustrates a key practical constraint: dose escalation to maximise methane suppression may inadvertently compromise rumen stability, either directly through additive acidity or indirectly via altered microbial activity. Notably, a 4% inclusion of Product 3 reduced CH₄ proportion by 62% without negatively affecting fermentation, with medium pH maintained at 5.84. This finding highlights the importance of dose optimisation, demonstrating that effective methane mitigation can be achieved without compromising rumen function when inclusion levels are carefully calibrated (Patra and Yu, 2012; Belanche et al., 2020). Despite their potential, optimal dosing strategies, suitable carrier matrices, and feed-processing interactions that influence additive stability and persistence in the rumen remain under-researched (Hristov et al., 2025). Addressing these factors is essential to improve the consistency, safety, and scalability of EO-based methane mitigation strategies in ruminant production systems.

3.2.2 Duration of supplementation and implications for long-term mitigation

The relatively short duration of the *in vivo* trial also limits confidence in the persistence of the observed methane reductions. Methane mitigation strategies based on plant secondary metabolites are particularly susceptible to time-dependent declines in efficacy, as rumen microbes adapt to repeated exposure (Patra and Yu, 2012).

This limitation has important implications for carbon accounting and mitigation claims. For a feed additive to contribute meaningfully to national greenhouse gas inventories or carbon credit schemes, its effects must be consistent, predictable, and durable over time (Hristov et al., 2015; Beauchemin et al., 2022). Short-term reductions measured over several weeks may not translate into reliable annual mitigation if microbial adaptation leads to diminishing responses.

Accordingly, the present findings should be viewed as evidence of short-term mitigation potential rather than definitive proof of long-term effectiveness. Extended feeding trials across production cycles are required to assess the stability of the methane response and to quantify any adaptation-related attenuation of efficacy (Hristov et al., 2022).

3.2.3 Constraints associated with GreenFeed methane measurements

Methodological constraints related to methane quantification using the GreenFeed system must also be considered. GreenFeed relies on voluntary animal visits, meaning that methane estimates depend on visit frequency, timing, and duration (Hammond et al., 2016b). Although minimum thresholds for visit numbers improve precision, differences in animal behaviour inevitably introduce sampling bias (Hammond et al., 2016a).

In the present study, treatment differences in visit duration and frequency were observed, particularly for steers receiving the 200 g/day dose. While this may reflect increased motivation to consume pellets, it complicates the interpretation of methane production data because animals with more frequent or longer visits contribute disproportionately to the dataset (Velazco et al., 2016). If treatment affects motivation to visit GF (e.g., due to palatability or novelty), the emissions dataset becomes behaviour-conditioned. Protocols recommend standardising attractant type and delivery, auditing visit distributions, and verifying that treatment differences in visitation do not drive apparent CH₄ responses (Hammond et al., 2015; Hammond et al., 2016b).

Furthermore, GreenFeed captures emissions during short sampling windows rather than continuously. Which increases variability compared with whole-day enclosure techniques as respiration chambers. This limitation is well documented, even though GreenFeed remains a validated method for estimating enteric methane emission in cattle (Hammond et al., 2016b; Arthur et al., 2017). Therefore, modest treatment differences, particularly those approaching statistical tendencies, should be interpreted with appropriate caution.

Regardless of the method used, methane measurement is affected by both systematic and random errors, which together define the uncertainty of the results (Gates et al., 2009; Tedeschi et al., 2022). Because uncertainty has not been evaluated for all published emissions values, comparisons across studies, assessment of data quality, and certification of emission reductions remain challenging. Establishing standardised methodologies for uncertainty assessment will be a critical step forward (Hristov et al., 2018). Further research is required to validate individual measurement techniques, assess their performance across diverse production scenarios, and establish calibration and standardisation protocols for existing methodologies (Tedeschi et al., 2022).

3.3 Future research directions

Building on the present findings, future work should prioritise field-relevant validation, persistence of the antimethanogenic effect, dose optimisation under voluntary intake, and clarification of the mechanism of action. The following studies are recommended:

3.3.1 Validate Product 3 in hardened MLB under commercial grazing:

Conduct paddock trials using finalised, hardened blocks to quantify true voluntary intake behaviour under grazing, including between-animal variability, the proportion of non-consumers over consumers, and intake stability across differing forage conditions. High variability in MLB intake is well documented and represents a major challenge for ensuring consistent additive delivery at the herd level (Bowman and Sowell, 1997; Imaz et al., 2019; Callaghan et al., 2020).

To strengthen inference, block-level disappearance should be paired with individual intake monitoring technologies (e.g. electronic identification–linked feeders or marker-based approaches) (Imaz et al., 2019) to link intake distribution with methane outcomes and animal performance. Integrating individual intake data with in-paddock methane measurement systems (e.g., GreenFeed® or portable accumulation chambers) (Hammond et al., 2015) would allow assessment of how

heterogeneous supplement consumption translates into variation in methane mitigation efficacy and growth responses at the animal and herd level (Charmley et al., 2015).

Collectively, such studies would provide critical evidence on the feasibility, robustness, and scalability of Product 3 as a practical methane mitigation strategy within commercial grazing systems, where self-regulated intake and environmental variability are unavoidable.

3.3.2 Assess persistence of the methane response over longer durations

Longer-term supplementation trials are required to determine whether the observed methane-reducing effect is maintained, attenuated, or lost over time as a result of rumen microbial adaptation. Although the present study extended over 56 days, adaptation of rumen microbial communities to antimethanogenic compounds—particularly plant secondary metabolites and essential oil-based additives—has been widely reported and may substantially reduce efficacy beyond short experimental periods (Patra and Yu, 2012; Cobellis et al., 2016).

Season-long studies conducted under production-relevant conditions would enable quantification of the temporal stability of methane suppression, as well as potential rebound effects following prolonged exposure. Such designs would also allow assessment of interactions between additive persistence, diet quality, and seasonal changes in pasture composition, which are known to influence both rumen fermentation and methane emissions (Hristov et al., 2013; Beauchemin et al., 2020; Hristov et al., 2022).

Critically, evaluating persistence over extended durations is essential to ensure that short-term mitigation responses translate into durable, cumulative emissions reductions that are compatible with national greenhouse gas emissions inventories and emerging carbon accounting frameworks. Without evidence of sustained efficacy, extrapolation of short-term methane reductions risks overestimating real-world mitigation potential (Gerber et al., 2013b; Hristov et al., 2015).

3.3.3 Clarify mechanisms of action

Complementary studies are required to elucidate how Product 3 influences rumen microbial ecology, hydrogen outputs, and fermentation pathways that ultimately regulate methanogenesis. Integrating microbial community profiling with metabolomic analyses would provide insight into shifts in rumen metabolic networks induced by supplementation (Morgavi et al., 2010; Ungerfeld, 2015b).

Such approaches would help determine whether methane suppression arises from direct inhibition of methanogens, redirection of metabolic hydrogen toward competing pathways or a broader restructuring of the rumen microbiome (Patra and Yu, 2012). This mechanistic resolution is particularly important for explaining the dose-dependent responses observed under controlled feeding and for determining whether any time-dependent attenuation of efficacy reflects microbial adaptation or functional redundancy within the rumen ecosystem (Beauchemin et al., 2020).

A clearer understanding of the mechanism will strengthen confidence in the durability and predictability of Product 3 under variable grazing conditions and will inform rational optimisation of dose, formulation, and delivery strategy in future field applications.

3.3.4 Evaluate economic feasibility and adoption potential

Finally, evaluation of Product 3 should extend beyond biological efficacy to include whole-farm economic and adoption analyses. Integrating additive cost, delivery infrastructure, labour requirements, and any effects on animal productivity, with estimates of methane abatement value (e.g. carbon credits or emissions intensity reductions) (Callaghan et al., 2020) would provide a

realistic assessment of commercial viability. Such whole-system approaches are essential to determine whether observed methane reductions translate into economically meaningful outcomes at the enterprise level (Beauchemin et al., 2020; Hegarty et al., 2021).

Equally important is understanding producer decision-making constraints, including variability in supplement intake, compatibility with extensive grazing management, perceived risk, and regulatory or market incentives. Previous studies highlight that adoption of methane mitigation technologies in extensive systems is strongly influenced by practicality, cost-effectiveness, and ease of integration into current management rather than mitigation potential alone (Almeida and Hegarty, 2021; Almeida et al., 2023).

Addressing these economic and behavioural dimensions, alongside technical performance, will be critical for scaling Product 3 beyond experimental settings and enabling durable, large-scale implementation within pasture-based livestock industries.

3.4 Conclusion

This thesis demonstrated that the molasses-based additive formulation Product 3 can reduce enteric methane emissions in beef cattle without negatively affecting performance. *In vitro* screening showed strong, dose-dependent inhibition of methane production, with reductions exceeding 90% at high inclusion levels, while moderate inclusion (4%) achieved a 62% reduction without impairing fermentation, supporting dose optimisation and guiding the selection of Product 3 for *in vivo* testing.

In the subsequent 56-day *in vivo* trial, Product 3 reduced methane yield by up to 11.7% under controlled feeding conditions, with no adverse effects on feed or water intake, or average daily gain. The positive associations between supplement intake, growth, and water consumption suggest good palatability and compatibility with forage-based diets. Overall, these findings indicate that Product 3 can effectively mitigate methane emissions while maintaining productivity, demonstrating its potential applicability in pasture-based beef production systems. Several methodological and practical limitations must be considered when interpreting these findings. The pellet-based delivery system does not reflect the intake variability and behavioural dynamics associated with free-access molasses lick blocks, the intended commercial delivery method. In addition, the known potential for rumen microbial adaptation to plant secondary metabolites highlights the need for longer-duration studies to assess persistence of the antimethanogenic effects over time.

Despite these limitations, this thesis provides valuable evidence supporting the potential of molasses-based additives containing essential oils as methane mitigation tools for grazing beef cattle. The demonstration of a clear dose-response relationship under controlled conditions provides a strong foundation for further formulation refinement.

Future research should prioritise longer-term *in vivo* evaluation, validation under grazing conditions using hardened lick blocks, and mechanistic investigations of rumen microbial and fermentation responses. These steps will be essential to confirm durability of mitigation effects and determine suitability for integration into emerging carbon accounting frameworks.

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