



# Article Impact of Oil Sources on In Vitro Fermentation, Microbes, Greenhouse Gas, and Fatty Acid Profile in the Rumen

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**Abstract:** This study estimated the effects of oil sources on fermentation characteristics, greenhouse gas, microbial diversity, and biohydrogenation of fatty acids in the rumen. In vitro ruminal incubation was performed with 7 mg of oil source, 15 mL rumen buffer, and 150 mg of synthetic diet at 39 °C for 0, 3, 6, 12, and 24 h. Oil sources consisted of corn oil (CO; linoleic acid (C18:2n-6)), linseed oil (LSO; linolenic acid (C18:3n-3)), or Ca-salts (protected C18:2n-6). The ruminal gas was collected for CH<sub>4</sub> and CO<sub>2</sub> analysis. Incubated rumen buffer was sub-sampled for the analysis of microbial quantification, fermentation characteristics, and fatty acid profiles. The results showed that Ca-salt increased acetate (p = 0.013), while CO increased propionate (p = 0.007). *Fibrobacter succinogenes, Ruminococcus flavefaciens*, and *R. albus* increased (p < 0.05) with Ca-salt after 12 h of incubation, while *Streptococcus bovis* increased (p < 0.05) by LSO. The CO and Ca-salt resulted in the highest C18:2n-6 (p = 0.002), while LSO resulted in the highest C18:3n-3 (p = 0.001). The Ca-salt supplementation resisted biohydrogenation to some extent, decreased methanogenic archaea and protozoa, and exerted less toxic effects on fibrolytic bacteria.

Keywords: fatty acid; methane emission; oil sources; rumen fermentation; rumen microbes

# 1. Introduction

Research interest in the field of fatty acid (FA) supplementation for ruminants has increased in recent decades mainly with the desire to influence a number of physiological processes and to manipulate FA compositions of products derived from ruminants [1]. Those FAs having important biological functions as well as health benefits for both humans and livestock are of major concern, e.g., linoleic acid (C18:2n-6) and linolenic acid (C18:3n-3). These FAs were reported as essential FA for animals [1,2].

Supplementing oil sources to ruminant diet, poly-unsaturated FA (PUFA) in particular has been shown to reduce enteric methane (CH<sub>4</sub>) emission [3,4]. A depressing effect on ruminal methanogenesis was observed with linseed oil [5], sunflower seed oil [5,6], or canola oil [7]. The oil sources with free FAs in the rumen having antimicrobial effects [8,9] can change the proportional production of volatile FA (VFA) [9,10], and reduce CH<sub>4</sub> production [11]. Nevertheless, oil sources undergo extensive lipolysis and biohydrogenation of their PUFA by rumen microbes [12]. Calcium salt of FAs was found to be degraded to a



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**Copyright:** © 2022 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (https:// creativecommons.org/licenses/by/ 4.0/). lower extent in rumen [13], exert less toxicity on microbes [14], and thereby may potentially manipulate FA compositions at a physiological or product level. In our previous study, the effects of pure C18:2n-6, C18:3n-3, and their combination were studied [10]. However, the application of pure FAs is not practical at the farm level due to the expensiveness and the difficulties of handling and storage. The use of rich-oil sources can be an alternative solution as a rumen modifier instead of pure FA, which is more applicable for farmers. Even though varied rich oil can be used as a source of essential FA, there is limited information concerning its use.

In general, rich-oil sources of C18:2n-6 FA and C18:3n-3 FA have been used to demonstrate the effect on growth performance, milk production, and meat fatty acids [5–7]. Corn oil is known as a source of C18:2n-6 FA with a concentration of approximately 50.0–58.2% of total FA [15,16]. Besides that, linseed oil also contains C18:3n-3 FA of approximately 52.5–55.6% of total FA [17,18]. Calcium salt is formed by an ionic bond between the free carboxyl group of the FA and calcium ions [19]. The FA can bypass the rumen without disturbing the rumen microorganisms by the insoluble character of the calcium salt [20], which is commercially available and widely used as a fat supplement for lactating cow and beef cattle, which demand high-energy diets. Generally, the supplementation of pure essential FA was reported to reduce enteric CH<sub>4</sub> emission and biohydrogenation of essential FA, but it could present a negative effect on rumen microbes or fermentation [9,10]. Thus, protected FA was applied in the present study to investigate its effects on rumen fermentation. As a hypothesis, protecting C18:2n-6 FA with calcium salt was expected to reduce the negative effects of essential fatty acids on rumen microbes or fermentation.

Recently, there has been limited information concerning a comparison of the effects of corn oil, linseed oil, and protected C18:2n-6 FA using calcium salt as a rumen modifier in an in vitro study. Therefore, the present study was conducted to investigate the effects of oil sources of FAs containing C18:2n-6, C18:3n-3, and protected C18:2n-6 on rumen fermentation, greenhouse gas emission, microbial diversity, and biohydrogenation of FA through in vitro technique.

# 2. Materials and Methods

# 2.1. In Vitro Incubation

Animal care and handling in the present study was approved by the Animal Ethic Committee of Gyeongsang National University, Korea (GNU-191011-E0050). Two rumen cannulated Hanwoo heifers fed rice straw and commercial concentrate at a ratio of 8:2 were used to collect rumen fluid. The crude protein (CP) and neutral detergent fiber (NDF) of rice straw were 5.40% and 63.85%, respectively. CP and NDF of commercial concentrate containing ground corn, barley meal, soybean meal, rice bran, corn gluten feed, and so on were 12.51% and 47.51%, respectively. Approximately 1 L of rumen fluid was collected from both animals before morning feeding, filtered with double layers of cheesecloth, and bulked to obtain homogenous inoculum. Then, rumen fluid (600 mL) was mixed with anaerobic culture medium (1200 mL) at a 1:2 ratio according to Adesogan et al. [21], which mainly consisted of micro- and macro-minerals with trypticase. An anaerobic condition was maintained during mixing by continuous flushing of carbon dioxide  $(CO_2)$  into the container of the rumen buffer in accordance with previous studies [22,23]. A fat-free synthetic diet containing 411 g cellulose (Sigma, C6413, St. Louis, MO, USA), 411 g starch (Sigma, S4180, St. Louis, MO, USA), and 178 g casein (Sigma, C3400, St. Louis, MO, USA) per kg DM was used as substrate [24]. Dietary treatments contained either corn oil (C18:2n-6; CO), linseed oil (C18:3n-3; LSO), or Ca-salts of rapeseed oil (protected C18:2n-6; Ca-salt). The FA compositions of oil sources are illustrated in Table 1. The Ca-salt of FA contained 28.50% C18:2n-6. Therefore, the Ca-salt treatment was considered as a rich source of protected n-6 FA in the present study. Ruminal incubation was performed in 50 mL glass serum bottles containing 150 mg of synthetic diet, 7 mg of each oil source (CO vs. LSO vs. Ca-salt), and 15 mL of rumen

buffer. The application ratio of oil sources demonstrated the energy diet for beef cattle. Incubations were conducted at 39 °C for 0, 3, 6, 12, and 24 h with 5 replications and 5 blanks for each incubation time period. The blanks only contained 15 mL of rumen buffer without the synthetic diet and oil sources.

Item _	Treatment <sup>1</sup>					
	СО	LSO	Ca-Salt			
C10:0	0.01	0.01	0.02			
C12:0	0.02	0.02	0.10			
C14:0	0.06	0.06	0.28			
C14:1	0.00	0.00	0.01			
C15:0	0.01	0.02	0.03			
C16:0	11.5	5.30	12.4			
C16:1	0.04	0.03	0.05			
C17:1	0.03	0.05	0.00			
C18:0	1.94	3.84	3.59			
C18:1cis-9	29.7	19.2	50.8			
C18:2n-6	55.0	15.6	28.5			
C18:3n-3	1.22	55.7	2.32			
C20:0	0.40	0.04	0.53			
C20:1	0.00	0.00	0.99			
C20:3	0.12	0.15	0.32			

Table 1. Fatty acid compositions of oil sources (% total fatty acid).

<sup>1</sup> CO, corn oil (C18:2n-6); LSO, linseed oil (C18:3n-3); Ca-salt, calcium salt of FA (protected C18:2n-6).

## 2.2. Sampling and Chemical Analysis

After each incubation time, the headspace gas pressure from incubation bottles was measured with a pressure transducer (Fisher Scientific, Traceable<sup>TM</sup>, Friendswood, TX, USA) attached with Luer-lock three-way stopcock, and then bottles were placed into the ice to stop microbial activity, as described by Honkanen et al. [25]. Total gas volume was calculated from gas pressure (psi) according to Mauricio et al. [26]. The gas sample was collected in vacutainer tubes without additives for  $CH_4$  and  $CO_2$ analysis using a multi-gas analyzer (Yes Plus LGA, Critical Environment Technologies, Canada Inc., Delta, BC, Canada). After the bottle was opened, an incubated sample was sub-sampled at 1 mL for microbial quantification using PCR. Additionally, 2 mL rumen buffer was collected for FA analysis. After that, the incubated sample was centrifuged at  $2568 \times g$  for 15 min to separate incubated rumen buffer and the remaining diet [22,23]. The incubated rumen buffer was sampled for analyses of pH, ammonia-N  $(NH_3-N)$ , total VFA, and FA profiles. The pH was determined by using a pH meter (SevenEasy, Mettler Toledo, Greifensee, Switzerland). The NH<sub>3</sub>-N was measured according to Chaney and Marbach [27] through distillation of the sample in a Buchi apparatus (B-342, BUCHI, Flawil, Switzerland) followed by titration with 0.1N H<sub>2</sub>SO<sub>4</sub> in a burette. For further analyses of rumen fermentation, the incubated rumen buffer was centrifuged at  $21,500 \times g$  for 15 min [22,23] and the supernatant was collected to determine the concentrations of VFA using HPLC (L-2200, Hitachi, Tokyo, Japan) fitted with a UV detector (L-2400, Hitachi, Tokyo, Japan) and a silica gel column (Metacarb 87H, Varian, Palo Alto, CA, USA) and concentrations were determined according to Muck and Dickerson [28].

The FA concentrations of oil source and rumen buffer were determined using the gas chromatography technique. One milligram of internal standard (C19:0) was added to the sample before freeze drying to calculate the total FA concentration. The two-step methylation procedure of Jenkins et al. [29] was used for the preparation of FA methyl ester (FAME), which was then analyzed using gas chromatography (450-GC, Varian, Palo Alto, CA, USA) equipped with an autosampler (CP-8400, Varian, Palo Alto, CA, USA), a flame ionization detector, and a Varian capillary column (CP-Sil 88, Palo Alto, CA, USA, 100 m  $\times$  0.25 mm  $\times$  0.2 µm). Hydrogen was the carrier gas. The injector and

detector were maintained at 230 °C. The oven temperature was initially set at 120 °C for 1 min, increased by 5 °C/min up to 190 °C, held at 190 °C for 30 min, increased again by 2 °C/min up to 220 °C, and held at 220 °C for 40 min. The peak of the samples was identified, and the concentrations were calculated based on the retention time and peak area of known standards.

## 2.3. DNA Extraction, Primer Information, and Real-Time PCR Analysis

The DNA extraction procedure involved a physical disruption of cells by a mini bead-beater (BioSpec Products, Bartlesville, OK, USA) followed by isolation and purification using a commercial DNA extraction kit (QIAamp DNA mini kit, Qiagen, Germantown, MD, USA). A 700 mL aliquot of homogenously incubated rumen sample was transferred into a 1.5 mL microcentrifuge tube. Then, 180 mL of buffer animal tissue lysis (ATL) and 20 mL of proteinase K (supplied in the QIAamp DNA Mini Kit) was added into the tube, followed by vortexing. The tubes were then incubated at 56 °C for 12 h for cell lysis in a heating block (HB-48, Daihan Scientific Co., Ltd., Seoul, Korea). Next, the DNA purification protocol was followed as described in the manufacturer's instruction manual (Qiagen, Germantown, MD, USA). A NanoDrop Spectrophotometer (Thermo Scientific, ND-1000, Santa Clara, CA, USA) was used to determine the genomic DNA concentrations in the extracted samples, and the concentrations of all template DNA samples were adjusted to 50 ng/ $\mu$ L. PCR conditions were 95 °C for 5 min, followed by 35 cycles of 95 °C for 15 s and 56~60 °C for 30 s. The universal primers for bacteria and species-specific primers for Fibrobacter succinogenes, Ruminococcus flavefaciens, Ruminococcus albus, Streptococcus bovis, rumen methanogenic archaea, and ciliates (Entodinium) are given in Table 2. Real-Time qPCR was performed using Bio-Rad C1000 Touch<sup>TM</sup> Thermal cycler Real-Time PCR detection system (CFX96<sup>TM</sup> Real-Time system, Bio-Rad Laboratories, Inc., Hercules, CA, USA), with fluorescence detection of SYBR Green Real-Time Master Mix (TOYOBO Co. Ltd., Osaka, Japan). The values of cycle threshold (Ct) after Real-Time PCR were used to determine fold change of different microbial populations relative to the control (universal bacteria). Abundance of these microbes was expressed by the equation: relative quantification =  $2^{-\Delta CT(Target) - \Delta CT(Control)}$ , where Ct represents threshold cycle. The ribosomal protein gene for rumen bacteria was used as a reference gene for internal control as done by Kim et al. [30]. All quantitative PCR mixtures (final volume of 20  $\mu$ L) contained forward and reverse primers (200 nM), SYBR Green Master Mix, DNA template (50 ng), and sterilized distilled water. A negative control without template DNA was used.

Target Organism	Sequence (5'-3')	Annealing Temp (°C)/Size (bp)	Reference
Bacteria	F-CGGCAACGAGCGCAACCC R-CCATTGTAGCACGTGTGTAGCC	55.3/130	Denman and McSweeney [31]
Fibrobacter succinogenes	F-GTTCGGAATTACTGGGCGTAAA R-CGCCTGCCCTGAACTATC	61.0/121	Denman and McSweeney [31]
Ruminococcus falvefaciens	F-GAACGGAGATAATTTGAGTTTACTTAGG R-CGGTCTCTGTATGTTATGAGGTATTACC	57.7/132	Denman and McSweeney [31]
Ruminococcus albus	F-CCCTAAAAGCAGTCTTAGTTCG R-CCTCCTTGCGGTTAGAACA	59.3/175	Koike and Kobayashi [32]
Streptococcus bovis	F-CTAATACCGCATAACAGCAT R-AGAAACTTCCTATCTCTAGG	57.0/869	Tajima et al. [33]
Methanogenic archaea	F-TTCGGTGGATCDCARAGRGC R-GBARGTCGWAWCCGTAGAATCC	56.0/140	Denman et al. [34]
Ciliates (Entodinium)	F-GAGCTAATACATGCTAAGGC R-CCCTCACTACAATCGAGATTTAAGG	55.0/180	Skillman et al. [35]

Table 2. Oligonucleotide primers used for Real-Time PCR assay.

### 2.4. Statistical Analysis

The study was a completely randomized design. All data were analyzed using PROC ANOVA of Statistical Analysis System (SAS) ver. 9.3 [36]. Its model was  $Yij = \mu + Ti + eij$ , where Yij = response variable,  $\mu$  = overall mean, T = effect of treatment i, and eij = error effect. In addition, rumen pH, NH<sub>3</sub>-N, acetate, propionate, CH<sub>4</sub>, CO<sub>2</sub>, rumen microbes, and 18-carbon FA data were also analyzed using PROC GLM of SAS to test the significant levels of dietary treatment, incubation time, and the interaction between dietary treatment and incubation time. Its model was  $Y_{ijk} = \mu + \alpha_i + \beta_j + (\alpha\beta)_{ij} + e_{ijk}$ , where  $Y_{ijk}$  = response variable,  $\mu$  = overall mean,  $\alpha_i$  = the effect of dietary treatment,  $\beta_j$  = the effect of incubation time,  $(\alpha\beta)_{ij}$  = the interaction effect between dietary treatment and incubation time,  $\alpha\beta_{ij}$  = the interaction effect between dietary treatment and incubation time,  $\alpha\beta_{ij}$  = the interaction effect between dietary treatment and incubation time,  $\alpha\beta_{ij}$  = the interaction effect between dietary treatment and incubation time,  $\alpha\beta_{ij}$  = the interaction effect between dietary treatment and incubation time,  $\alpha\beta_{ij}$  = the interaction effect between dietary treatment and incubation time,  $\alpha\beta_{ij}$  = the interaction effect between dietary treatment and incubation time,  $\alpha\beta_{ij}$  = the interaction effect between dietary treatment and incubation time,  $\alpha\beta_{ij}$  = the interaction effect between dietary treatment and incubation time,  $\alpha\beta_{ij}$  = the interaction effect for significance using Tukey test at  $p \leq 0.05$ .

#### 3. Results

## 3.1. Fatty Acid Profiles before Incubation

At 0 h of ruminal incubation, dietary treatment did not affect total FA, but had effects on concentrations of individual FA (Table 3). The C15:0 concentration was higher (p = 0.047) in LSO than in Ca-salt. The Ca-salt had higher C18:1*cis*-9 concentration (p = 0.026) than CO and LSO. The C18:2n-6 concentration was lower (p = 0.013) in LSO than in CO and Ca-salt. The C18:3n-3 concentration was higher (p = 0.004) in LSO than in CO and Ca-salt. The Ca-salt had higher mono-unsaturated FA (MUFA) concentration (p = 0.029) than CO and LSO. The ratio of n-6 to n-3 was lower (p < 0.006) in LSO than in CO and Ca-salt. The concentration of other individual FAs, saturated FA (SFA), and PUFA were not affected by dietary treatments.

#### 3.2. Fermentation Characteristics and Greenhouse Gas Emission

The Ca-salt had higher pH (p = 0.024) and acetate concentration (p = 0.013) than CO, while dietary LSO was not different compared to other treatments (Table 4). The Ca-salt and LSO had a higher propionate concentration (p = 0.007) than CO. Moreover, Ca-salt had a higher acetate to propionate ratio (p = 0.015) than CON. Total gas volume, and concentrations of NH<sub>3</sub>-N, total VFA, butyrate, CH<sub>4</sub>, and CO<sub>2</sub> were not affected by the treatments.

	Treatments <sup>1</sup>			SEM	<i>p</i> -Value
	СО	LSO	Ca-Salt		p vulue
Total FA, mg/mL	3.05	3.19	3.62	0.069	0.604
C14:0, % total FA	2.65	2.52	1.96	0.393	0.316
C15:0, % total FA	11.4 <sup>ab</sup>	12.6 <sup>a</sup>	8.55 <sup>b</sup>	0.837	0.047
C16:0, % total FA	21.8	23.2	20.1	1.015	0.115
C16:1, % total FA	0.89	1.34	0.63	0.174	0.059
C17:0, % total FA	0.58	0.55	0.47	0.070	0.368
C18:0, % total FA	24.6	24.5	25.6	0.918	0.501
C18:1 <i>cis-</i> 9, % total FA	14.7 <sup>b</sup>	14.9 <sup>b</sup>	21.4 <sup>a</sup>	1.360	0.026
C18:2n-6, % total FA	20.0 <sup>a</sup>	9.33 <sup>b</sup>	18.7 <sup>a</sup>	1.643	0.013
C18:3n-3, % total FA	0.92 <sup>b</sup>	8.78 <sup>a</sup>	1.25 <sup>b</sup>	0.851	0.004
C20:0, % total FA	1.76	1.52	0.85	1.011	0.686
C22:0, % total FA	0.64	0.64	0.48	0.087	0.245
SFA <sup>2</sup> , % total FA	63.5	65.7	58.1	2.702	0.135
MUFA, % total FA	15.6 <sup>b</sup>	16.2 <sup>b</sup>	22.0 <sup>a</sup>	1.325	0.029
PUFA, % total FA	20.9	18.1	19.9	1.911	0.435
SFA: PUFA	3.07	3.65	2.92	0.436	0.347
n-6/n-3	21.7 <sup>a</sup>	1.06 <sup>b</sup>	15.0 <sup>a</sup>	2.216	0.006

**Table 3.** Effects of oil source supplementation containing essential fatty acid on fatty acid profile of rumen buffer just before incubation.

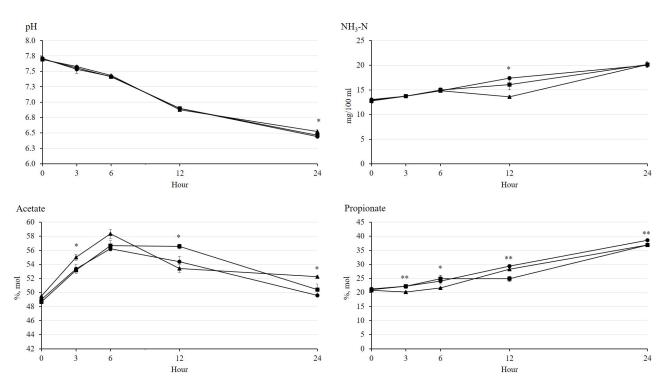
<sup>1</sup> CO, corn oil (C18:2n-6); LSO, linseed oil (C18:3n-3); Ca-salt, calcium salt of FA (protected C18:2n-6). <sup>2</sup> SFA, saturated fatty acid; MUFA, mono-unsaturated fatty acid; PUFA, poly-unsaturated fatty acid. <sup>a,b</sup> Means having different superscripts in the same row differ significantly (p < 0.05).

**Table 4.** Effects of oil source supplementation containing essential fatty acid on rumen fermentation characteristics and greenhouse gas emissions of rumen buffer after 24 h of incubation.

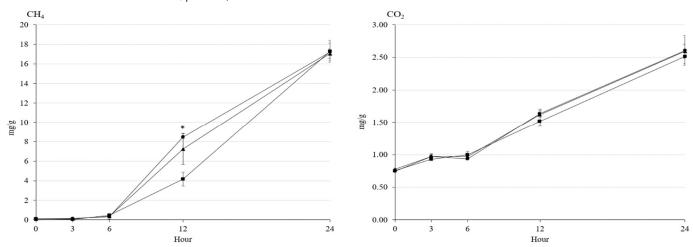
	Treatments <sup>1</sup>			SEM	<i>p</i> -Value
-	СО	LSO	Ca-Salt	SEIVI	p vulue
рН	6.44 <sup>b</sup>	6.46 <sup>ab</sup>	6.52 <sup>a</sup>	0.041	0.024
$NH_3$ -N, mg/100 mL	20.0	20.2	20.2	1.157	0.976
Total VFA, mmol/L	58.9	60.1	57.0	1.641	0.065
Acetate, % mol	49.6 <sup>b</sup>	50.4 <sup>ab</sup>	52.2 <sup>a</sup>	0.997	0.013
Propionate, % mol	38.5 <sup>a</sup>	36.9 <sup>b</sup>	36.9 <sup>b</sup>	0.630	0.007
Butyrate, % mol	11.0	11.9	10.6	0.685	0.060
Acetate: propionate	1.28 <sup>b</sup>	1.36 <sup>ab</sup>	1.41 <sup>a</sup>	0.048	0.015
Gas volume, mL/g DM	38.8	37.4	37.7	0.816	0.059
CH <sub>4</sub> , mg/g	17.2	15.2	17.1	1.171	0.209
$CO_2$ , mg/g	2.61	2.51	2.59	0.271	0.879

<sup>1</sup> CO, corn oil (C18:2n-6); LSO, linseed oil (C18:3n-3); Ca-salt, calcium salt of FA (protected C18:2n-6). <sup>a,b</sup> Means having different superscripts in the same row differ significantly (p < 0.05).

The rumen pH decreased by an increase of incubation time without any effect by treatments (Figure 1). The NH<sub>3</sub>-N concentration increased by incubation time, and Ca-salt presented the lowest (p < 0.05) NH<sub>3</sub>-N concentration among the treatments at 12 h. The acetate concentration increased by 6 h, but then decreased by 24 h of incubation. The highest acetate concentration was shown in Ca-salt among the treatments at 3 h (p < 0.05) and in LSO at 12 h (p < 0.05) of incubation. The propionate concentration gradually increased with the increase in incubation time. At 3 and 6 h of incubation, the propionate concentration was higher (p < 0.01) either in CO or LSO than in Ca-salt, but higher (p < 0.01) either in CO or Ca-salt than in LSO at 12 h of incubation. There was an interaction between dietary treatment and incubation time (p < 0.001) on NH<sub>3</sub>-N, acetate, and propionate concentrations. In addition, concentrations of CH<sub>4</sub> and CO<sub>2</sub> gases gradually increased with incubation time (Figure 2). The LSO resulted in the lowest (p < 0.05) CH<sub>4</sub> emission at 12 h of incubation among the treatments, while the emission of CO<sub>2</sub> was not affected by dietary treatment during incubation.



**Figure 1.** Changes in rumen pH and concentrations of NH<sub>3</sub>-N, acetate, and propionate during rumen incubation with oil sources containing essential fatty acid for 24 h. Diet with corn oil, C18:2n-6 (•); diet with linseed oil, C18:3n-3 (**I**); and diet with Ca-salt, protected C18:2n-6 (**A**). The respective significance levels of dietary treatment, incubation time, dietary treatment × incubation time, and SEM for pH, NH<sub>3</sub>-N, acetate, and propionate are p < 0.001, p = 0.310, p = 0.376, and SEM=0.048; p < 0.001, p = 0.001, p < 0.001, and SEM = 0.864; p < 0.001, p = 0.002, p < 0.001, and SEM = 0.982; and p < 0.001, p < 0.001, p < 0.001, and SEM = 1.007. Different symbols at the same time differ (\*, p < 0.05; \*\*, p < 0.01).



**Figure 2.** Changes in concentrations of CH<sub>4</sub> and CO<sub>2</sub> during rumen incubation with oil sources containing essential fatty acid for 24 h of incubation. Diet with corn oil, C18:2n-6 (•); diet with linseed oil, C18:3n-3 (**■**); and diet with Ca-salt, protected C18:2n-6 (**▲**). The respective significance levels of dietary treatment, incubation time, dietary treatment × incubation time, and SEM for CH<sub>4</sub> and CO<sub>2</sub> are p < 0.001, p = 0.272, p = 0.052, and SEM = 1.120; and p < 0.001, p = 0.616, p = 0.930, and SEM = 128.76. Different symbols at the same time differ (\*, p < 0.05).

## 3.3. Rumen Microbes

In the present study, methanogenic archaea were lower (p = 0.003) in CO and Ca-salt than in LSO (Table 5). Rumen ciliates were lower (p = 0.018) in LSO than in CO, but

not different in Ca-salt compared to other treatments. On the other hand, Ca-salt had higher abundance of *F. succinogens* (p < 0.001) and *R. flavefaciens* (p = 0.005) than CO and LSO. The abundance of *R. albus* was highest with Ca-salt, followed by CO, and the lowest was in LSO (p < 0.001). The abundance of *S. bovis* was not affected by treatment in the present study. In general, all rumen microbes studied in the present study decreased with longer incubation (Figure 3). Abundance of methanogenic archaea was higher (p < 0.01) in dietary LSO than in CO and Ca-salt at 3 h and 24 h of incubations. Ca-salt had the highest (p < 0.05) abundance of *F. succinogens* at 3, 12, and 24 h, *R. albus* at 12 h, and *R. flavefaciens* at 12 h and 24 h among the treatments. The abundance of *S. bovis* in LSO was highest (p < 0.01) among the treatments at 12 h. The interaction effects between dietary treatment and incubation time (p < 0.05) were reported in the present study, which was founded in all rumen microbes, except rumen ciliates. The abundance of rumen ciliates was not affected by dietary treatments during 12 h of incubation.

after 24 h of incubation (fold change unit).
Treatments <sup>1</sup>

Table 5. Effects of oil sources supplementation containing essential fatty acid on rumen microbes

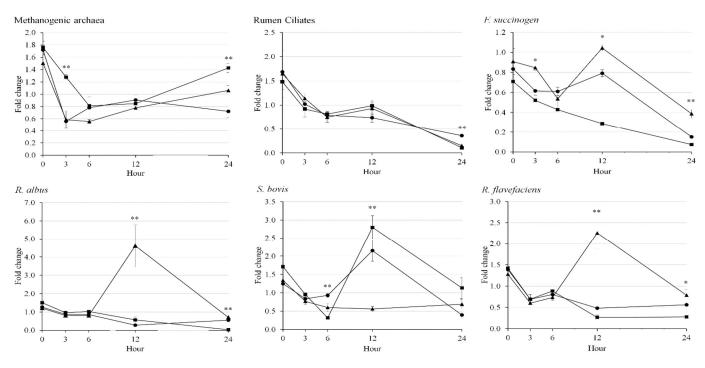
	Treatments <sup>1</sup>			SEM	<i>p</i> -Value
	СО	LSO	Ca-Salt	0 Em	,
Methanogenic archaea	0.72 <sup>b</sup>	1.43 <sup>a</sup>	1.06 <sup>b</sup>	0.145	0.003
Ciliates	0.36 <sup>a</sup>	0.13 <sup>b</sup>	0.22 <sup>ab</sup>	0.111	0.018
Fibrolytic bacteria					
F. succinogens	0.15 <sup>b</sup>	0.06 <sup>b</sup>	0.45 <sup>a</sup>	0.067	< 0.001
R. albus	0.43 <sup>b</sup>	0.04 <sup>c</sup>	0.82 <sup>a</sup>	0.146	< 0.001
S. bovis	0.39	1.13	0.68	0.324	0.083
R. flavefaciens	0.48 <sup>b</sup>	0.34 <sup>b</sup>	1.02 <sup>a</sup>	0.276	0.005

 $\overline{1}$  CO, corn oil (C18:2n-6); LSO, linseed oil (C18:3n-3); Ca-salt, calcium salt of FA (protected C18:2n-6). <sup>a-c</sup> Means having different superscripts in the same row differ significantly (p < 0.05).

#### 3.4. Fatty Acid Profiles after Incubation

After 24 h of ruminal incubation, total FA was not affected by treatment in the present study but had effects on concentrations of individual FAs (Table 6).

CO had higher concentrations of C14:0 (p = 0.007), C15:0 (p = 0.035), C16:0 (p = 0.005), and C16:1 (p = 0.003) than LSO and Ca-salt. LSO had the highest C18:0 concentration (p = 0.002), followed by CO, and then by Ca-salt. Ca-salt had higher C18:1*cis*-9 concentration (p < 0.004) than CO and LSO. The C18:2n-6 concentration (p = 0.002) was higher in CO and Ca-salt than LSO. The C18:3n-3 concentration (p < 0.001) was highest in LSO, followed by Ca-salt, and then by CO. The concentrations of C20:0 (p = 0.020) were higher in LSO than in Ca-salt, while CO was not different from other treatments. The SFA concentration (p < 0.004) was higher in CON and LSO than in Ca-salt, but the MUFA concentration (p < 0.004) was higher in Ca-salt than in CO and LSO. The concentration of PUFA was higher (p = 0.048) in LSO and Ca-salt than in CO in the present study. The SFA to PUFA ratio was higher in CO than in LSO and Ca-salt (p = 0.016), while the n-6 to n-3 FA ratio was higher in CO and Ca-salt than in LSO (p < 0.001).



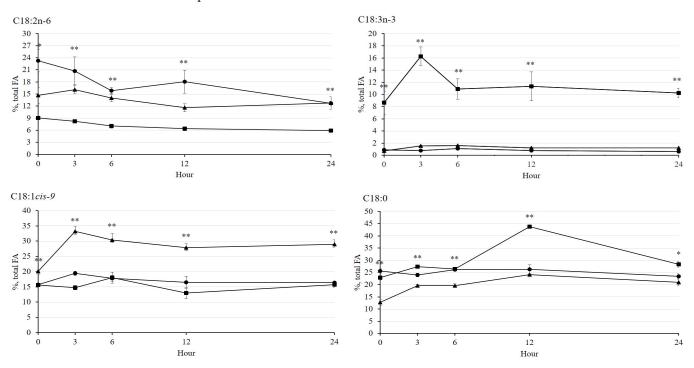
**Figure 3.** Fold changes in rumen microbes compared to that during rumen incubation with oil sources containing essential fatty acid for 24 h of incubation. Diet with corn oil, C18:2n-6 (•); diet with linseed oil, C18:3n-3 (**■**); and diet with Ca-salt, protected C18:2n-6 (**▲**). The respective significance levels of dietary treatment, incubation time, dietary treatment × incubation time, and SEM for Methanogenic archaea, rumen ciliates, *F. succionogens*, *R. albus*, *S. bovis*, and *R. flavefaciens* are p < 0.001, p < 0.001, p < 0.001, and SEM = 0.157; p < 0.001, p = 0.803, p = 0.838, and SEM = 0.283; p < 0.001, p < 0.001, p < 0.001, p < 0.001, and SEM = 0.138; p < 0.001, p < 0.001, p < 0.001, and SEM = 0.263; and p < 0.001, p < 0.001, and SEM = 0.198. Different symbol at the same time differ (\*, p < 0.05; \*\*, p < 0.01).

**Table 6.** Supplementation effects of oil sources containing essential fatty acid on fatty acid profiles of rumen buffer after 24 h of incubation.

	Treatments <sup>1</sup>			SEM	<i>p</i> -Value
-	CO	LSO	Ca-salt	JEIVI	p value
Total FA, mg/mL	4.04	5.02	5.51	0.076	0.086
C14:0, % total FA	8.27 <sup>a</sup>	6.28 <sup>b</sup>	5.24 <sup>b</sup>	0.494	0.007
C15:0, % total FA	8.00 <sup>a</sup>	5.02 <sup>b</sup>	5.46 <sup>b</sup>	0.785	0.035
C16:0, % total FA	27.8 <sup>a</sup>	22.2 <sup>b</sup>	23.0 <sup>b</sup>	0.876	0.005
C16:1, % total FA	3.04 <sup>a</sup>	1.75 <sup>b</sup>	2.06 <sup>b</sup>	0.161	0.003
C17:0, % total FA	1.06	0.80	0.81	0.082	0.058
C18:0, % total FA	25.4 <sup>b</sup>	33.8 <sup>a</sup>	20.1 <sup>c</sup>	0.938	0.002
C18:1 <i>cis-</i> 9, % total FA	14.2 <sup>b</sup>	15.0 <sup>b</sup>	28.2 <sup>a</sup>	2.339	0.004
C18:2n-6, % total FA	11.0 <sup>a</sup>	5.41 <sup>b</sup>	13.1 <sup>a</sup>	0.887	0.002
C18:3n-3, % total FA	0.65 <sup>c</sup>	9.24 <sup>a</sup>	1.24 <sup>b</sup>	0.066	< 0.001
C20:0, % total FA	0.62 <sup>ab</sup>	0.53 <sup>b</sup>	0.68 <sup>a</sup>	0.034	0.020
SFA, % total FA	71.2 <sup>a</sup>	68.6 <sup>a</sup>	55.3 <sup>b</sup>	3.053	0.008
MUFA, % total FA	17.2 <sup>b</sup>	16.8 <sup>b</sup>	30.3 <sup>a</sup>	2.204	0.004
PUFA, % total FA	11.6 <sup>b</sup>	14.6 <sup>a</sup>	14.4 <sup>a</sup>	0.910	0.048
SFA: PUFA	6.14 <sup>a</sup>	4.69 <sup>b</sup>	3.88 <sup>b</sup>	0.626	0.016
n-6/n-3	16.9 <sup>a</sup>	0.58 <sup>b</sup>	10.6 <sup>a</sup>	1.045	< 0.001

<sup>1</sup> CO, corn oil (C18:2n-6); LSO, linseed oil (C18:3n-3); Ca-salt, calcium salt of FA (protected C18:2n-6). SFA, saturated fatty acid; MUFA, mono-unsaturated fatty acid; PUFA, poly-unsaturated fatty acid. <sup>a-c</sup> Means having different superscripts in the same row differ significantly (p < 0.05).

In the present study, it was observed that the higher (p < 0.05) concentration of C18:2n-6 was either in CO or Ca-salt rather than in LSO during incubation (Figure 4). On the other hand, the C18:3n-3 concentration in LSO was highest (p < 0.01) among the treatments in all incubation time periods, while Ca-salt showed the highest (p < 0.01) concentration of C18:1*cis*-9. In general, LSO showed the highest (p < 0.01) C18:0 concentration among the treatments throughout the incubation, especially at 3, 12, and 24 h of incubation. These results could be influenced by interaction effects between dietary treatment and incubation time (p < 0.05) in the present study, which were reported in all major 18-carbon fatty acids, except C18:2n-6.



**Figure 4.** Changes in major 18-carbon fatty acid concentrations (%, FA) during rumen incubation with oil sources containing essential fatty acid for 24 h of incubation. Diet with corn oil, C18:2n-6 (•); diet with linseed oil, C18:3n-3 (**■**); and diet with Ca-salt, protected C18:2n-6 (**▲**). The respective significance levels of dietary treatment, incubation time, dietary treatment × incubation time, and SEM for C18:2n-6, C18:3n-6, C18:1*cis*-9, and C18:0 FAs are p < 0.001, p < 0.001, p = 0.478, and SEM = 2.442; p < 0.001, p < 0.001, p < 0.001, and SEM = 0.967; p < 0.001, p < 0.001, p = 0.040, and SEM = 2.098; p < 0.001, p < 0.001, p < 0.001, and SEM = 2.199. Different symbols at the same time differ (\*, p < 0.05; \*\*, p < 0.01).

# 4. Discussion

### 4.1. Rumen Fermentation Characteristics, Microbial Population, and Methanogenesis

The rumen pH gradually decreased with dietary FA with the advancement of time during ruminal incubation in the present study, but it still remained in a normal range [37]. Another study also observed that dietary linoleic and linolenic acids presented rumen pH within the range of 6.46 to 6.56 after 24 h of incubation [10], which was similar to the present study. On the other hand, CO showed the lowest acetate concentration, but the highest propionate concentration after 24 h of incubation in the present study, which might be caused by a lower population of methanogenesis archaea than other treatments. Decreased methanogens in the rumen usually shift hydrogen utilization from methanogenesis towards propionate synthesis [9,10]. Moreover, dietary C18:2n-6 reduced the methanogenic archaea during incubation [9,10]. Similar to our previous study [10], the present study reported that C18:3n-3 in LSO inhibits the CH<sub>4</sub> emission more effectively than the other dietary treatments, especially at 12 h of incubation.

Unlike our previous study, rumen ciliates decreased significantly with LSO as the C18:3n-3 source compared to CO and Ca-salt as the C18:2n-6 sources in the present study. Usually, the ciliate population in rumen decreased with fat supplementation, even though effects may have varied [14]. In agreement with the present study, linseed oil was reported to have more negative effects on ciliates (protozoa) than other sources of lipids [38]. Doreau and Ferlay [14] suggested that C18:3n-3 content of linseed might be responsible for negative effects on ciliates. A marked reduction in rumen ciliates (protozoa) with the increasing level of stearic (C18:0) and oleic acid (C18:1*cis*-9) was reported earlier [10]. In the present study, together with linolenic acid (C18:3n-3), stearic acid concentrations were also highest in LSO at 24 h of incubation. Therefore, these two FAs together might affect the rumen ciliate concentration in LSO.

In the present study, the negative effects of FA on the growth of most fibrolytic bacteria examined were lower in Ca-salt compared to CO and LSO. It is well recognized that calcium salts of FAs affect rumen fermentation to a lesser extent [39,40]. These salts are dissociated slowly in rumen [20] and hence produce lower toxicity to microbes compared to unprotected oil sources. The negative effects of FA on bacterial growth depend on their adsorption on the substrate cell wall [20]. Adsorption of FA decreases when the concentration of divalent cations is high [41] and its effect seems more dominant for Ca<sup>2+</sup> than others cations [42]. On the other hand, ionized Ca enhances the adhesion of bacteria to cellulose [43]. Application of LSO has shown greater toxicity to rumen ciliates as well as fibrolytic bacteria (except *S. bovis*) compared to CO. It was shown in a previous study that C18:3n-3 was comparatively more lethal to microbes, especially fibrolytic bacteria compared to C18:2n-6 [9,10]. The result of the present study regarding fibrolytic rumen bacteria is therefore in agreement with previous studies.

### 4.2. Rumen Fatty Acid and Profiles

After 24 h of incubation, along with C18:2n-6 and C18:3n-3, most other FAs were also found to be affected by dietary treatments in the present study. Honkanen et al. [25] and Amanullah et al. [10] had similar results, and reported that the results might be caused by a different rate of lipolysis and/or biohydrogenation among the treatments. The lowest total SFA, especially C18:0 along with the highest C18:1*cis*-9, and total MUFA in dietary Ca-salt indicated a lower degree of biohydrogenation with this treatment [12].

In the process of biohydrogenation of C18:2 and C18:3, different cis- and/or transisomers of C18:1 intermediates are produced, and then converted into C18:0 as the end product [12,25]. In the present study, a Ca-salt diet had the lowest C18:0 concentration, but highest total C18:1*cis*-9 concentration from 0 to 24 h of incubation. The C18:1 *cis*-9 concentration was increased rapidly at 3 h, which resulted from biohydrogenation mainly of C18:2n-6 and the higher level continued till the end. Results indicated calcium salts of FA can resist biohydrogenation of UFA by rumen microbes to some extent, as also reported by Wu et al. [13]. In particular, the release of free FA (lipolysis) in the rumen is markedly reduced when lipids are supplied in the form of calcium salts [44].

Higher n-6 to n-3 FA ratio in CO and Ca-salt than LSO at 24 h of incubation is an important indication for dietary supplementation strategy of n-6 and n-3 FAs in ruminants. For a long time, efforts have been made to incorporate these FAs or change their ratio in the products considering the health benefits of consumers. However, extensive biohydrogenation of those FAs acts as a bottleneck concerning the use of dietary supplementation strategies. The increased concentrations of n-6 and n-3 FA in the bottle content of respective treatments after 24 h of incubation suggest the potential incorporation of these FAs at a physiological level by dietary supplementation. By fitting data with single-pool, first-order kinetic models, Honkanen et al. [25] showed that the rate of linoleic acid disappearance decreased with increases in substrate availability.

# 5. Conclusions

Based on the present study with a high-energy diet for beef cattle, the supplementation of CO increased the molar proportion of propionate concentrations after 24 h of in vitro incubation compared to other FAs. Supplementation of CO and Ca-salt decreased methanogenic archaea, while supplementation of LSO decreased rumen ciliates efficiently. As a recommendation, the supplementation of Ca-salt was found to be less toxic to rumen fibrolytic bacteria and resistant against FA biohydrogenation indicated by lower total SFA, especially C18:0 along with the highest C18:1*cis*-9, and total MUFA compared to other treatments.

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