A note on the effect of rape seed oil supplementation on microbial protein synthesis in sheep

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ABSTRACT

The effect of rape seed oil on net microbial protein synthesis and other metabolic parameters was estimated on four rams fitted with rumen cannulas in a 4 x 4 Latin square design consisting of four diets differing in the percentage of rape seed oil supplement. The basal diet was composed of 50% concentrate and 50% meadow hay and was supplemented with 4, 8 or 10% of rape seed oil. Supplementation of the diet with 4 and 8% of rape seed oil caused a drop in net microbial protein production, determined by allantoin excretion in urine, but the differences were not significant. The highest dose of rape seed oil had no effect on the intestinal flow of microbial protein. Fat supplement had no influence on pH and ammonia-N or on total VFA, acetate and propionate concentrations. The butyrate concentration was, however, lower (P<0.05) when rape seed oil was added to the basal diet.

KEY WORDS: rumen, fat, microbial protein, allantoin, sheep

INTRODUCTION

It is difficult to meet energy requirements with forages or grain in animals already receiving maximum dry matter intake, hence the numerous attempts to increase the energy content of feeds for ruminants by increasing fat concentrations in diets (Sutton et al., 1983; Murphy et al., 1987). The main problem with a high level of fat is its effect on rumen microflora, which results in decreased cellulose digestion and reduced acetate/propionate ratio (Garnsworthy, 1997). On the other
hand, diets supplemented with fat improve the efficiency of microbial protein synthesis (Tesfa, 1993). The effect of fat added to ruminant rations has been the subject of considerable research, but there are many contradictory results. Furthermore, there is also a close relationship between microbial protein production measured by urinary allantoin excretion and diet composition (Puchala and Kulasek, 1992). Ben Salem et al. (1993) suggested that the negative effects of lipids on rumen digestion was less important when fibre intake was high.

The objective of this study was to evaluate the effect of added fat and concentrate/hay diet on basic ruminal parameters in sheep, mainly on net microbial protein synthesis.

**MATERIAL AND METHODS**

**Animals, feeds and feeding**

Four rumen cannulated rams of mean body weight of $50 \pm 3$ kg were fed four diets in a Latin square design. The diets were formulated on the basis of rape seed oil (RSO) content (Table 1). The basal diet was 1.2 kg/d of concentrate and hay (50:50 % w/w). RSO was mixed with concentrate while hay was chaffed into particles of 2-5 cm length. The energy value of the ration was 5.75 MJ NE, and the crude protein content was 144 g per kg. The animals were kept in individual pens and fed two equal portions of feed at 08.00 and 16.00. They had free access to fresh water.

<table>
<thead>
<tr>
<th>TABLE 1</th>
<th>Diets composition, % of DM</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>I</td>
</tr>
<tr>
<td>Rapeseed oilmeal</td>
<td>3.0</td>
</tr>
<tr>
<td>Wheat, ground</td>
<td>12.0</td>
</tr>
<tr>
<td>Rye, ground</td>
<td>10.0</td>
</tr>
<tr>
<td>Triticale, ground</td>
<td>20.0</td>
</tr>
<tr>
<td>Meadow hay</td>
<td>53.0</td>
</tr>
<tr>
<td>Rape seed oil</td>
<td>0.0</td>
</tr>
<tr>
<td>Mineral-vitamin mixture</td>
<td>1.5</td>
</tr>
<tr>
<td>Trace mineral salt*</td>
<td>0.5</td>
</tr>
</tbody>
</table>

* contains NaCl – 95.8%

**Sampling**

Rumen fluid. Samples from ventral sac of rumen fluid were taken at 0 (pre-feeding), 3 and 6 h post-feeding during 2 consecutive days after an adaptation
modified rumen fistula cap with polypropylene plate

sampling syringe

nylon tube

pneumating fitting

sockets

Figure 1. Sampling device

period to the diet (12 days). A sampling device which fits into a cannula cap (Figure 1) was used to collect the rumen fluid without access to oxygen. The sieving probe of the device is prepared from a solid block of polypropylene, 55 mm long with a diameter of 18 mm. The block was hollowed out from the top to give a cylinder with about 2 mm walls (the base remains solid). Small holes are drilled into the walls of the cylinder. A pneumatic fitting joins the probe with the tubing (nylon tubing 6 mm diameter). The tubing is approximately 140 mm long. The second pneumatic fitting connects the tube to the sampling syringe. To take samples the blanking plug is removed from the tube and the end of a syringe is pushed into the push-in fitting giving a good connection. The modified rumen fistula cap is pushed over the polypropylene plate of the sampling device which then forms a secure closure. The polypropylene plate has the same diameter as the inside of the rumen fistula cap. It has three holes in it. The middle one is the same diameter as the tubing. The other two can be used for infusion if the dual marker system is being used, or can be blanked off by inserting a short piece of tubing in "U" shape through the two holes. The sampled rumen fluid was subsampled for ammonia-N (20 ml) and VFAs (20 ml) determination.

Urine. Urine was collected for 48 h according to Gonda and Lindberg (1997), and the collection was performed using a urine collection device (Kowalczyk et al., 1996). The collected urine was weighed and sampled every 24 h. The samples were stored at -20°C until analysis.

Analytical methods

Rumen pH was measured immediately after collection using a Mera Tronic N 517 pH-Meter. Ammonia was measured by the Conway method (1962). Volatile fatty acids were determined qualitatively and quantitatively by gas chromatogra-
Microbial protein synthesis in sheep fed rapeseed oil

**Phy** (Ziolecki and Kwiatkowska, 1973) using a Chrom 5 chromatograph. Allantoin was determined by the modified HPLC method of Balcells et al. (1992) using a Hewlett-Packard HPLC system with UV detector. Separations were achieved by using two Adsorbsphere C18 (150 x 4.6 mm; Alltech) columns and ODS Hypersil column (200 x 4.6mm; Hewlett-Packard) connected in series.

*Calculation of microbial protein flow*

Amount of allantoin excreted in the urine was used for calculation of microbial N flow to the small intestine according to the equation: $N_M = (N_A - N_{AF}) \times 4 \times (100/18)$ where $N_M$ is microbial N (g d\(^{-1}\)), $N_A$ is allantoin N excreted during 24 h and $N_{AF}$ is allantoin N excreted during 24 h of fasting (Puchala and Kulaszk, 1992).

*Statistical calculations*

The results were subjected to statistical analysis of variance using SAS software program (User’s Guide, 1988).

**RESULTS AND DISCUSSION**

**Intestinal flow of microbial protein**

Studies on the urinary excretion of purine derivatives by ruminants have been stimulated by the possible use of their excretion as evidence of the rumen microbial protein supplied to the host animal. This is because nucleic acids flowing to the small intestines in ruminants are essentially of microbial origin. Absorbed purines are degraded to hypoxanthine, xanthine, uric acid and allantoin, which are excreted in urine, and should relate quantitatively to the amount of purines and, hence, microbial protein absorbed (Chen et al., 1992).

Dewhurst et al. (1988) assumed that allantoin-N represents 86% of purine N, which is absorbed from the gut with 86% efficiency. Some authors report that allantoin accounts for 85-90% of the total excretion of purine derivatives (Verbic et al., 1990; Vagnoni and Broderick, 1997). In fact some other authors suggested that when all purine derivatives are taken into consideration, quantitative assessment of microbial protein reaching the small intestine can be more accurate than that using allantoin alone (Czauderna and Kowalczyk, 1995).

In our experiment, allantoin was used for calculation of net microbial protein synthesis. Urinary excretion of allantoin was not significantly affected by the experimental treatments. The amounts of microbial nitrogen ($N_M$) calculated from urinary allantoin excretion were 12.57, 10.21, 10.53 and 12.05 g N d\(^{-1}\), in groups I,
II, III and IV, respectively (Table 2). According to Elliot et al. (1995) replacement of nonstructural carbohydrates such as starch, decreases the amount of energy that is available for microorganisms and may decrease microbial protein synthesis. We also observed some decrease in $N_M$ when the diet was supplemented with 4 and 8% RSO (see Tables 1 and 2). The differences, however, were statistically insignificant ($P > 0.05$). On the other hand, Murphy et al. (1987) found a positive correlation between the amount of fat added and net microbial protein synthesis when milking cows were fed rape seeds. A similar relationship was observed by Sutton et al. (1983) in relation to sheep fed a ration with linseed oil. If fat in the diet is to be fed successfully, it is recommended that ruminants consume the proper amount of fibre. Fibre minimizes the potential for negative effects of fat on rumen microbes (Grant and Weidner, 1992). In our experiment, sheep were fed equal portions of concentrate and roughage containing structural carbohydrates so none of the basic constituents was neglected. Further experiments with various levels of fibre in diets seem to be necessary.

**Ruminal fermentation characteristics**

The effect of added fat on pH and ammonia-$N$ content in rumen fluid are shown in Table 2. The values were similar in all groups and ranged from 6.4 to 6.5 for pH and 4.2 to 5.8 mmol/l for ammonia-$N$, respectively. The level of ammonia-$N$ was low, especially in the control group, and tended to increase as the amount of rape seed oil increased, but the differences were not statistically significant. In contrast with this, Tesfa (1993) reported a reduction in ammonia-$N$ concentration when Fresian bulls were fed a basal diet supplemented with 0.5 kg RSO from 9.61 mmol to 6.26 mmol in the basal and RSO groups, respectively. Similarly, feeding supplementary fat has been observed by Hall et al. (1990) to depress rumen ammonia concentration. Also, in our earlier experiment, the addition of 3 and 6% of RSO to the diet of sheep caused a significant depression in ammonia-$N$ concentration from 14.8 (control group) to 11.0 mmol and 8.5 mmol, respectively (Szumacher-Strabel and Potkański, 1997).

<table>
<thead>
<tr>
<th>Group</th>
<th>$N_M$, g/d</th>
<th>$N_3$-N, mmol</th>
<th>pH</th>
</tr>
</thead>
<tbody>
<tr>
<td>I</td>
<td>12.57 ± 3.42</td>
<td>4.29 ± 2.43</td>
<td>6.46 ± 0.61</td>
</tr>
<tr>
<td>II</td>
<td>10.21 ± 2.54</td>
<td>5.08 ± 2.30</td>
<td></td>
</tr>
<tr>
<td>III</td>
<td>10.53 ± 3.27</td>
<td>5.80 ± 3.83</td>
<td></td>
</tr>
<tr>
<td>IV</td>
<td>12.05 ± 4.03</td>
<td>5.87 ± 4.59</td>
<td></td>
</tr>
</tbody>
</table>

**TABLE 2**

Effect diets of on net microbial protein synthesis, rumen pH and the concentration of NH$_3$-N in the rumen of sheep
TABLE 3

<table>
<thead>
<tr>
<th></th>
<th>Group</th>
<th></th>
<th></th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>I</td>
<td>II</td>
<td>III</td>
<td>IV</td>
</tr>
<tr>
<td>Total VFA</td>
<td>74.82 ±16.99</td>
<td>79.39 ± 17.13</td>
<td>78.36 ±16.25</td>
<td>82.74 ±15.20</td>
</tr>
<tr>
<td>Acetone</td>
<td>44.71 ± 7.17</td>
<td>38.68 ± 10.21</td>
<td>44.09 ± 7.07</td>
<td>36.73 ±12.76</td>
</tr>
<tr>
<td>Propionic</td>
<td>23.06 ± 3.71</td>
<td>20.28 ± 4.03</td>
<td>21.61 ± 5.04</td>
<td>22.99 ± 7.16</td>
</tr>
<tr>
<td>Butyric</td>
<td>11.94 ±2.83</td>
<td>9.26 ± 1.97</td>
<td>9.84 ± 3.80</td>
<td>8.16 ± 2.46</td>
</tr>
<tr>
<td>Isobutyric</td>
<td>2.36 ± 0.43</td>
<td>2.17 ± 0.64</td>
<td>2.27 ± 0.61</td>
<td>2.50 ± 0.92</td>
</tr>
<tr>
<td>Valeric</td>
<td>2.00 ± 0.76</td>
<td>1.59 ± 0.59</td>
<td>2.07 ± 0.92</td>
<td>1.55 ± 0.74</td>
</tr>
<tr>
<td>Isovaleric</td>
<td>1.83 ± 0.62</td>
<td>1.72 ± 0.37</td>
<td>2.18 ± 0.82</td>
<td>1.66 ± 2.46</td>
</tr>
</tbody>
</table>

A, B – P<0.01

pH values were not significantly affected by treatment. These observations are consistent with those reported by Murphy et al. (1987) in lactating cows fed diets containing full-fat rape seeds. In an experiment by Kowalczyk et al. (1977) rumen pH tended to increase with as the amount of tallow in diets for sheep rose (0, 50, 100, 150 g of tallow/kg grass).

There were no significant effects of RSO on the VFA level. The butyric acid concentration, however, was significantly reduced (P < 0.05) in animals fed diets supplemented with RSO (Table 3). According to Tesfa (1993), who obtained similar results, the decrease in butyrate concentration was accompanied by a decrease in the number of ciliates. This suggests that RSO could influence the fermentation pattern in the rumen affecting the composition of the rumen microbial population.

According to Tackett et al. (1996) supplementation of unprotected fats generally has a negative influence on ruminal fermentation. In our study, there was no negative effect of rape seed oil in the range from 4 to 10% on rumen parameters. We observed a positive effect up to 8% and a small depression at 10%.

In conclusion, the experiment supports the assumption that feeding RSO to ruminants does not necessarily depress net microbial protein synthesis, but this depends on the quantity of added fat.

REFERENCES


STRESZCZENIE

Wpływ dodatku oleju rzepakowego na syntezę białka mikroorganizmów w żwaczu u owiec

Na czterech owcach, z trwałymi przetokami żwacza, badano wpływ zróżnicowanego dodatku oleju rzepakowego (4, 8 i 10%) do diety, składającej się z 50% paszy treściwej i 50% siana, na produkcję białka drobnoustrojowego, stosując alantoinę wydalaną w moczu jako wskaźnik. Oznaczono pH, N-amonowy i LKT w płynie żwaczowym.

Dodatek 4 i 8% oleju spowodował obniżenie produkcji białka drobnoustrojowego, jednakże w stopniu statystycznie nieistotnym. Dodatek tłuszczu nie miał także wpływu na pH i poziom N-amonowego. Tłuszcz dodany do diety obniżył poziom kwasu masłowego (P<0.05) we wszystkich grupach w porównaniu z grupą kontrolną.