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Effects of pomegranate peel extract on ruminal and post-ruminal *in vitro* degradation of rumen inoculum of the dairy cow

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ABSTRACT

This experiment was carried out to study the effect of water extracted pomegranate peel extract (PE) on ruminal protein degradation and post-ruminal digestion in the dairy cow. PE was added at six levels of total phenolics (g/kg of the basal diet); 3.75 (PE1); 4.4 (PE2); 5.05 (PE3); 5.70 (PE4); and 6.35 (PE5). Rumen degradable crude protein (rdCP) decreased with PE addition (L < 0.0001), but total CP degradability (tdCP) was not affected. Compared to PE0, PE2, and PE3 diets showed higher (L = 0.054, Q = 0.029) digestibility of bypass CP (dBCP). Increasing levels of PE resulted in a decrease in proteolytic bacteria numbers (p < 0.0001). At PE4 and PE5 levels, total VFA and acetate concentrations linearly decreased compared to PE0. PE inclusion lowered the acetate:propionate ratio (L = 0.0001) and Ammonia-N production after 24 h (L = 0.0008) of incubation. The total number of protozoa, genera *Dasytricha* and *Isotricha*, and subfamilies *Entodiniinae*, *Diplodiniinae*, and *Ophrioscolecinae* decreased with increasing dietary PE concentration (p < 0.0001). The results suggest that all levels of PE addition reduce the protozoal population and Ammonia-N concentration. All PE levels slowed down protein degradation in the rumen but PE2 and PE3 showed the greatest effect.

Introduction

The cost of animal feeds, particularly protein supplements, has increased mainly due to changing climatic conditions and a shortage of water resources; yet these supplements may be metabolized less efficiently (i.e., losses of NH₃-N) in the rumen resulting in a reduction in animal performance.¹ Consequently, many studies have been carried out to improve the efficiency of nitrogen utilization, including the use of aqueous extracts of plant secondary metabolites (AE).² Plant secondary metabolites (PSM) in tree leaves such as Salix babylonica and Leucaenaleucocephala^{3,4} were found to have a positive effect on ruminal fermentation parameters and to increase amino acid flow to the duodenum.⁵ This led to greater muscle deposition and milk production.⁶ PSM are a diverse group of molecules involved in the adaptation of plants to their environment but are not part of the primary biochemical pathways of cell growth and reproduction.² Other

KEYWORDS

Dairy cow; pomegranatepeel extract; protein; protozoa; rumen

studies have shown that plant secondary metabolites decreased rumen proteolysis.⁷ Research has shown that tannin extracts from *Cistus ladanifer* led to a reduction in effective rumen degradability of soybean protein and that linearly decreased with extract inclusion level.⁷ In other research, grape pomace extract slowed soybean meal protein degradation in the rumen and the post-ruminal digestibility of this undegraded protein increased in proportion to the concentration of the extract.⁸

PSM extraction is generally carried out using organic solvents,⁸ a relatively expensive procedure, but this work tests the effectiveness of a cheaper water extraction method (AE). Globally, the annual production of pomegranate peel (PP), a by-product of pomegranate juice extraction, is 15,000,000 t,⁹ and it contains high levels of PSM such as polyphenols, tannins, saponins, and punicalaginan.⁹ This study aims to determine the optimum levels of pomegranate peel

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Table 1. The economics of water and organic solvent (OS) extraction of Total Phenolics (TP) from Pomegranate Peel (PP).

| | Solvents Water | Ethanol + Methanol (50:50)(OS) | | |
|------------------------------|-------------------|-----------------------------------|--|--|
| Total Extract obtained (mL)* | 750 mL | 500 mL | | |
| mg TP/mL extract | 6.5 mg | 10 mg | | |
| Cost (\$)/total extract | 0.1 | 20 | | |
| Cost (\$)/mg total phenolic | 0.00002 | 0.004 | | |

*When mixing 1 kg of PP with either 1 L of water or 1 L OS, the total extract obtained was 750 mL and 500 mL respectively.

extract (PE) on the *in vitro* ruminal degradability and intestinal digestibility of a basal diet in the dairy cow.

Materials and methods

Pomegranate peel extract

The PP was sun-dried and then the AE was extracted using 1 g PP/mL of distilled water. This mixture was maintained at 40 °C for 72 h in a closed flask then the contents were strained and the extract kept at 4° C until used. Preliminary work had been done to extract total phenolics (TP) using an ethanol/methanol mixture (50:50, v/v) (OS) by the same method to enable a cost comparison (Table 1).

Experimental diets

Six levels of PE were used in this study. A basal diet was formulated (Table 2) according to NRC¹⁰ recommendations for dairy cows. 321 mg dry matter (DM) of diet was weighed into 120 mL serum bottles.¹¹ PE was added to the bottles at six levels of total phenolics (TP) (0, 0.65, 1.30, 1.95, 2.60, and 3.25 g TP/kg DM of basal diet). The basal diet (PE0) contains 3.1 g TP/kg DM, therefore, PE1 contains 3.75 g TP/kg DM with corresponding levels for PE2: 4.4 g, PE3: 5.05, PE4: 5.70, and PE5: 6.35. PE levels were selected for their secondary metabolite concentrations and impacts on protein degradability but kept within a range that would not negatively affect protein degradation and ruminal fermentation. Five sets of bottles were prepared for the determination of (i) ruminal in vitro DM and CP degradation, (ii) total in vitro DM and CP degradation, (iii) NH₃-N production and protozoa population after 12, 24, and 48 h of incubation, and VFA sampling and determination of proteolytic bacteria after 48 h of incubation (one set per incubation time).

In vitro gas production and related parameters

Nutrient degradation by ruminal microbes and by ruminal microbes plus HCl/pepsin were determined.¹¹

Table 2. Ingredients and chemical composition of the basal diet¹.

| Ingredients, g/kg DM | |
|---|-------------|
| Alfalfa hay | 148.4 |
| Corn silage | 229.0 |
| Wheat straw | 17.7 |
| Soybean meal, 44% CP | 119.2 |
| Canola meal | 34.6 |
| Cottonseed meal | 41.4 |
| Fish meal | 9.9 |
| Barley, rolled | 121.8 |
| Corn grain, ground, dry | 111.1 |
| Wheat grain, rolled | 17.1 |
| Wheat bran | 68.1 |
| Fat powder | 18.3 |
| Calcium carbonate | 5.7 |
| Sodium bicarbonate | 15.0 |
| Mineral and vitamin premix ² | 26.0 |
| Salt | 8.8 |
| Magnesium oxide | 8.8 |
| Chemical composition, g/kg DM | |
| DM | 955 |
| OM | 934 |
| CP | 160 |
| EE | 46 |
| ³ NE _L , Mcal/kg DM | 1.58 |
| NDFom ^a | 305 |
| ADFom ^b | 206 |
| ADL ^c | 57.60 |
| Decel dist Calculated from (NDC) Natio | wal Deserve |

¹Basal diet Calculated from (NRC) National Research Council (2012); ²Contained 196 g Ca, 96 g P, 71 g Na, 19 g Mg, 3 g Fe, 0.3 g Cu, 2 g Mn, 3 g Zn, 100 ppm Co, 100 ppm I, 0.1 ppm Se and 50×10^5 IU vitamin A, 10×10^5 IU vitamin D and 0.1 g vitamin E/kg; ³NE_L calculated from (NRC) National Research Council (2012): NE_L (Mcal/ kg) = 0.0245 × TDN (%) × 0.12; ^aNDFom: ash-free NDF; ^bADFom: ash-free ADF; ^cADL: lignin.

Extract effects were examined in three runs of *in vitro* gas production. The contents of one set of bottles were filtered after 48 h of incubation to measure rumen degradable dry matter and rumen degradable crude protein, and a second set was incubated for another 24 h in the presence of HCl/pepsin to measure the total degradability of dry matter and total degradability of crude protein.

Prior to their morning feed, rumen fluid was collected from three cannulated Holstein dairy cows fed three times daily on a TMR diet formulated according to NRC.¹⁰ The cows' diet consisted of 80% forage to 20% concentrate with 12% CP/kg DM and 1.5 Mcal/ kg DM of NEL (net energy of lactation). The rumen fluids were mixed and randomly assigned to each treatment group. The fluid was filtered through four layers of cheesecloth, transferred into a separator funnel, and gassed with CO₂. After standing for 15 min at 39 °C, the upper and lower parts of the fluid in the funnel were discarded. The residue was mixed with McDougall's buffer at a ratio 1:4 to obtain the inoculum. Fifty mL of the inoculum was added to the bottles with five bottles (n=5) prepared per diet. After gasifying with CO_2 , the bottles were closed with rubber stoppers and incubated at 39 °C.

After 48 h, the incubation was stopped in one set of bottles by the addition of 1 mL of mercuric chloride solution (50 mg/mL). The second set of bottles had 6 mL of HCl solution (6.21 N) and 2 mL pepsin solution added (50 g/L; 2844-01, USA; 1:3000) and the incubation continued for 24 h, simulating postruminal digestion. After incubation, the contents of both sets of bottles were strained through N-free paper filters (no. 10300 012, S&S Whatman, Dassel, Germany). The residues were dried at 105 °C for 12 h, weighed and the nitrogen content measured using Kjeldahl analysis (method 984.13; AOAC 1990). For the first set of bottles, rdDM and rdCP were calculated by comparing DM and CP before and after incubation; the second set of bottles were similarly treated to calculate tdDM and tdCP. The difference between tdCP and rdCP was assumed to be equivalent to the degradable rumen escape CP (undegradable CP in the rumen but digested postruminally; pdBCP). From this value, the digestibility of bypass CP (dBCP) was calculated.

Total and individual VFA in samples were determined by gas chromatography. At the end of the incubation (48 h) 1 mL of the supernatant was collected in a microfuge tube containing 0.2 mL 20% orthophosphoric acid and 20 mM 2-ethyl butyric acid, as the internal standard, and then centrifuged at 15,000 g for 15 min at a temperature of 4 °C. The supernatant was collected and stored at -20 °C until analysis.¹² For VFA estimation, 1 ul supernatant was injected into a gas chromatograph (UNICAM 4600; SB Analytical, Cambridge, UK) equipped with a capillary column (Agilent J&W HP-FFAP, 10 m by 0.535 mm by 1.00 μ m, 19,095 F-121; Agilent, Santa Clara, CA)¹³

To measure NH₃-N production during fermentation, sub-samples of 5 mL were taken after 12, 24, and 48 h of incubation from the respective sets of bottles and mixed with 1 mL of 0.2 N HCl for NH₃-N analysis using the phenol-hypochlorite method.¹⁴ Subsamples were frozen at -20 °C until analysis.

Total number and subfamily counts of protozoa were determined.¹⁵ Two mL of rumen fluid was pipetted into a screw-capped test tube containing 5 mL of formalinized physiological saline (20 mL formaldehyde in 100 mL saline (0.85 g sodium chloride in 100 mL distilled water)). Total and differential counts of protozoa were made in 30 microscopic fields at a magnification of $200 \times$ in a Hemocytometer (Neubauer improved, Marienfeld, Germany).

A 5-mL sample from the content of each bottle was used for enumeration of proteolytic bacteria.

Modified anaerobic techniques¹⁶ were used in the preparation of the anaerobic culture media.

Analytical methods

Representative samples from basal diet and experimental diets were taken to carry out the following analysis: dry matter content was determined by oven drying at 105 °C for 16 h (AOAC, method 967.03).¹⁷ Ash content was determined by incineration at 550 °C for 6 h, and the OM content was calculated as the difference between 100 and the percentage of ash (AOAC, method 942.05).¹⁷ NDF was determined, with sodium sulfite in a neutral detergent solution (ND),¹⁸ and ADFom was determined according to AOAC, method 973.18¹⁷ and expressed exclusive of residual ash. The amount of ADL was determined by cellulose solubilization with sulfuric acid.¹⁹ Nitrogen content was determined by the Kjeldahl method.¹⁷

Total phenolics (TP), Non-tannin phenolics (NTP) were measured using the Folin-Ciocalteau method.²⁰ Total tannins (TT) were calculated as the difference between TP and NTP. Tannic acid (Merck GmbH, Darmstadt, Germany) was used as the standard to express the amount of TP and TT. Condensed tannins (CT) were measured by the HCl-butanol method. Hydrolyzable tannins were analyzed using the Rhodanine assay.²⁰ Ten mL of the extract were prepared after TP separation and double the volume of n-butanol was added to fractionate saponins.²¹ The GC-MS analysis of dihydromaltol and thymol isolated from PE was performed using a Perkin Elmer GC-MS (Model Perkin Elmer Clarus 500, USA) equipped with a VF-5 MS fused silica capillary column (30 m \times 0.25 mm id., film thickness 0.25 μ m). GC-MS spectroscopic detection, an electron ionization system with ionization energy of 70 eV was used. Pure helium gas (99.999%) was used as a carrier gas at a constant flow rate of ±1 mL/min. Mass transfer line and injector temperatures were set at 220 °C and 290 °C, respectively.

Statistical analysis

Incubation was done in three separate *in vitro* runs with five replicate test feed samples. Data on *in vitro* runinal fermentation parameters of each of the three runs were averaged and used as the mean value for each individual sample within the diets. Data were subjected to one-way analysis of variance using the MIXED procedure of SAS version 9.0 (SAS Institute, Cary, NC, USA)²² and the effects of the different

Table 3. Secondary metabolites levels (g/kg of DM) of the experimental diets (n = 20).

| Diets ¹ | | | | | | | | |
|--------------------|---|--|--|--|--|--|--|--|
| PE0 | PE1 | PE2 | PE3 | PE4 | PE5 | | | |
| 3.10 ± 0.05 | 3.75 ± 0.06 | 4.40 ± 0.08 | 5.05 ± 0.03 | 5.70 ± 0.05 | 6.35 ± 0.09 | | | |
| 0.80 ± 0.06 | 1.40 ± 0.09 | 2.00 ± 0.05 | 2.60 ± 0.08 | 3.20 ± 0.05 | 3.80 ± 0.07 | | | |
| 2.30 ± 0.05 | 2.35 ± 0.04 | 2.40 ± 0.08 | 2.45 ± 0.05 | 2.50 ± 0.07 | 2.55 ± 0.07 | | | |
| - | 0.4 ± 0.02 | 0.8 ± 0.03 | 1.2 ± 0.03 | 1.6 ± 0.05 | 2.0 ± 0.04 | | | |
| - | 0.045 ± 0.005 | 0.090 ± 0.004 | 0.135 ± 0.005 | 0.180 ± 0.002 | 0.225 ± 0.003 | | | |
| 11.00 ± 0.10 | 11.25 ± 0.100 | 11.50 ± 0.09 | 11.75 ± 0.10 | 12.00 ± 0.12 | 12.25 ± 0.12 | | | |
| - | 0.03 ± 0.002 | 0.06 ± 0.002 | 0.09 ± 0.003 | 0.12 ± 0.001 | 0.15 ± 0.003 | | | |
| - | 0.02 ± 0.001 | 0.04 ± 0.002 | 0.06 ± 0.001 | 0.08 ± 0.040 | 0.10 ± 0.030 | | | |
| | $3.10 \pm 0.05 \\ 0.80 \pm 0.06 \\ 2.30 \pm 0.05 \\ - \\ - \\ 11.00 \pm 0.10 \\ - \\ -$ | $\begin{array}{ccccc} 3.10\pm 0.05 & 3.75\pm 0.06 \\ 0.80\pm 0.06 & 1.40\pm 0.09 \\ 2.30\pm 0.05 & 2.35\pm 0.04 \\ - & 0.4\pm 0.02 \\ - & 0.045\pm 0.005 \\ 11.00\pm 0.10 & 11.25\pm 0.100 \\ - & 0.03\pm 0.002 \end{array}$ | $\begin{tabular}{ c c c c c c c } \hline PE0 & PE1 & PE2 \\ \hline $3.10 \pm 0.05 & $3.75 \pm 0.06 & 4.40 ± 0.08 \\ $0.80 \pm 0.06 & $1.40 \pm 0.09 & 2.00 ± 0.05 \\ $2.30 \pm 0.05 & $2.35 \pm 0.04 & 2.40 ± 0.08 \\ $-$ & $0.4 \pm 0.02 & 0.8 ± 0.03 \\ $-$ & $0.045 \pm 0.005 & 0.090 ± 0.004 \\ $11.00 \pm 0.10 & $11.25 \pm 0.100 & 11.50 ± 0.09 \\ $-$ & $0.03 \pm 0.002 & 0.06 ± 0.002 \\ \hline \end{tabular}$ | $\begin{tabular}{ c c c c c c c c c c c c c c c c c c c$ | $\begin{tabular}{ c c c c c c c c c c c c c c c c c c c$ | | | |

¹Diets: PE0, control, no additive = 3.1 g TP/kg DM of basal diet; PE1, control + 0.65 = 3.75 g TP/kg DM of basal diet; PE2, control + 1.30 = 4.4 g TP/kg DM of basal diet; PE3, control + 1.95 = 5.05 g TP/kg DM of basal diet; PE4, control + 2.60 = 5.70 g TP/kg DM of basal diet; PE5, control + 3.25 = 6.35 g TP/kg DM of basal diet.

Table 4. In vitro dry matter degradation (mg/g incubated DM) of the experimental diets incubated with pomegranate peel extract.

| | Diets ¹ | | | | | | | p-V | alue |
|-------------------|--------------------|--------|--------|--------|--------|--------|------------------|----------------|-------|
| | PE0 | PE1 | PE2 | PE3 | PE4 | PE5 | SEM ² | L ³ | Q^4 |
| rdDM⁵ | 681.17 | 681.39 | 691.34 | 686.28 | 560.68 | 550.37 | 34.253 | 0.039 | 0.048 |
| tdDM ⁶ | 716.91 | 716.45 | 714.63 | 712.32 | 700.28 | 691.42 | 6.450 | 0.079 | 0.337 |

¹Diets: PE0, control, no additive = 3.1 g TP/kg DM of basal diet; PE1, control + 0.65 = 3.75 g TP/kg DM of basal diet; PE2, control + 1.30 = 4.4 g TP/kg DM of basal diet; PE3, control + 1.95 = 5.05 g TP/kg DM of basal diet; PE4, control + 2.60 = 5.70 g TP/kg DM of basal diet; PE5, control + 3.25 = 6.35 g TP/kg DM of basal diet.

²SEM: standard error of the mean; ³L: linear; ⁴Q: quadratic; ⁵rdDM: Degradation of DM incubated in ruminal fluid.

⁶tdDM: Degradation of DM when incubated with ruminal fluid + HCl/pepsin.

levels of PE were partitioned into linear and quadratic components by orthogonal polynomials. Mean comparisons were performed using Duncan's test at p < 0.05.

$$Yij = \mu + Si + eij,$$

Where *Yij* is the general observation, μij the general mean, *Si* the ith effect of extracts on the observed parameters and *eij* the standard error term.

Results

Ingredients and chemical composition

Ingredients, chemical composition and net energy of lactation (NEL) in the basal diet, and secondary metabolites of the experimental diets are given in Tables 2 and 3, respectively.

Dry matter and crude protein degradability

The amount of rdDM did not differ between PE0, PE1, PE2, and PE3 diets but was lower in PE4 and PE5 compared to PE0 (Q = 0.048) (Table 4). The rdCP decreased linearly with increased addition of PE (L < 0.0001, Q < 0.0001), but tdDM and tdCP values were not affected by PE inclusion.

Addition of PE increased (L = 0.001, Q = 0.002) the proportion of ruminally un-degraded CP (rumen escape CP- Table 5). The amount of dBCP in PE2

and PE3 diets was higher compared to the PE0 diet (L=0.054, Q=0.029), but did not differ between PE1, PE4, and PE5 diets. The population of proteolytic bacteria decreased with increasing PE levels (L < 0.0001, Q < 0.0001).

The effect of pomegranate peel extract on volatile fatty acids and Ammonia-N production

The profile of VFA production is shown in Table 6. Total VFA and acetate concentrations did not differ between PE0, PE1, PE2, and PE3 diets but declined in PE4 and PE5 compared to PE0 (L = 0.029 and L = 0.008 respectively). Propionate, butyrate, isobutyrate and valerate concentrations were not influenced by the diets. The acetate to propionate ratio was lower (L = 0.0001) in PE3, PE4, and PE5 diets compared to the PE0 diet. The addition of PE decreased NH₃-N production (L = 0.0038 and Q = 0.0113) compared to the PE0 diet (Table 6).

Enumeration of rumen protozoa

Total numbers of protozoa (L < 0.0001, Q < 0.0001), genus Isotricha (L < 0.0001, Q < 0.0001), and Dasytricha (L < 0.0001, Q < 0.0001), subfamily of Entodiniinae (L = 0.0003, Q = 0.0083), Diplodiniinae (L < 0.0001, Q < 0.0001), and Ophrioscolecinae (L < 0.0001, Q < 0.0001) decreased with increasing levels of PE in the diet.

| | Diets ¹ | | | | | | | p-V | alue |
|---|--------------------|--------|--------|--------|--------|--------|------------------|----------------|----------------|
| | PE0 | PE1 | PE2 | PE3 | PE4 | PE5 | SEM ² | L ³ | Q ⁴ |
| rdCP⁵ | 132.77 | 119.23 | 120.80 | 119.40 | 119.10 | 119.60 | 0.171 | <0.0001 | < 0.0001 |
| tdCP ⁶ | 136.81 | 133.07 | 135.42 | 133.89 | 132.91 | 133.47 | 1.142 | 0.077 | 0.704 |
| pdBCP ⁷ | 6.79 | 13.83 | 14.61 | 14.49 | 13.81 | 13.87 | 1.33 | 0.001 | 0.002 |
| pdBCP ⁷ dBCP ⁸ | 0.25 | 0.34 | 0.37 | 0.36 | 0.34 | 0.34 | 0.029 | 0.054 | 0.029 |
| Proteolytic bacteria | 7.42 | 6.42 | 6.40 | 6.38 | 6.28 | 6.26 | 0.014 | < 0.0001 | <.0001 |

Table 5. *In vitro* crude protein degradation of the experimental diets incubated with pomegranate peel extract either in ruminal fluid or ruminal fluid followed by HCI/pepsin, and the proteolytic bacterial count (log₁₀/g digesta).

¹Diets: PE0, control, no additive = 3.1 g TP/kg DM of basal diet; PE1, control + 0.65 = 3.75 g TP/kg DM of basal diet; PE2, control + 1.30 = 4.4 g TP/kg DM of basal diet; PE3, control + 1.95 = 5.05 g TP/kg DM of basal diet; PE4, control + 2.60 = 5.70 g TP/kg DM of basal diet; PE5, control + 3.25 = 6.35 g TP/kg DM of basal diet.

²SEM: standard error of the mean; ³L: linear; ⁴Q: quadratic; ⁵rdCP: Degradation of CP incubated with ruminal fluid (mg/g incubated CP); ⁶tdCP: Degradation of CP incubated with ruminal fluid + HCI/pepsin (mg/g incubated CP); ⁷pdBCP: Protein un-degraded in ruminal fluid (bypass CP) but degraded with HCI/pepsin (mg/g incubated CP); ⁸dBCP: Degradability of bypass CP in HCI/pepsin (mg/g rumen escape CP).

Table 6. Effect of pomegranate peel extract level on the total VFA (mmol), individual VFA (mol/100 mol) and NH₃–N production (mg/dL).

| ltem | | | | <i>p</i> -Value | | | | | |
|--|-------|-------|-------|-----------------|-------|-------|------------------|----------------|--------|
| | PE0 | PE1 | PE2 | PE3 | PE4 | PE5 | SEM ² | L ³ | Q^4 |
| Total VFA ⁵ Individual VFA | 74.00 | 73.50 | 74.40 | 70.43 | 59.46 | 59.00 | 4.249 | 0.029 | 0.068 |
| Acetate | 46.58 | 47.18 | 45.45 | 42.72 | 34.52 | 31.69 | 3.122 | 0.008 | 0.104 |
| Propionate | 15.51 | 14.66 | 16.68 | 15.62 | 14.14 | 14.50 | 1.105 | 0.580 | 0.264 |
| Butyrate | 7.01 | 6.99 | 7.99 | 6.75 | 7.23 | 7.07 | 0.499 | 0.902 | 0.522 |
| lsobutyrate | 2.54 | 3.53 | 2.96 | 4.08 | 2.62 | 4.27 | 0.426 | 0.604 | 0.067 |
| Isovalerate | 0.84 | 0.88 | 1.02 | 1.00 | 0.71 | 1.17 | 0.090 | 0.649 | 0.032 |
| Valerate | 0.25 | 0.25 | 0.27 | 0.26 | 0.23 | 0.29 | 0.012 | 0.409 | 0.061 |
| Acetate:Propionate | 3.12 | 3.22 | 2.72 | 2.66 | 2.27 | 2.20 | 0.132 | 0.0001 | 0.301 |
| NH ₃ –N ⁶ | 77.66 | 52.00 | 50.52 | 49.49 | 51.58 | 47.39 | 5.676 | 0.0038 | 0.0113 |

¹Diets: PE0, control, no additive = 3.1 g TP/kg DM of basal diet; PE1, control + 0.65 = 3.75 g TP/kg DM of basal diet; PE2, control + 1.30 = 4.4 g TP/kg DM of basal diet; PE3, control + 1.95 = 5.05 g TP/kg DM of basal diet; PE4, control + 2.60 = 5.70 g TP/kg DM of basal diet; PE5, control + 3.25 = 6.35 g TP/kg DM of basal diet.

²SEM: standard error of the mean; ³L: linear; ⁴Q: quadratic; ⁵VFA: Volatile fatty acids; ⁶NH₃–N: ammonia-N.

Discussion

Dry matter and crude protein degradability

Dietary supplementation with PE decreased rdDM in PE4 and PE5 compared to PE0. Similarly, it was reported that purified condensed tannins extracted from *Calliandra*, *Flemingia*, and *Leucaena* decreased rumen degradability of dry matter in soybean meal.¹¹ Other work indicated that the addition of chestnut (*Castanea spp.*) tannins to soybean meal decreased *in vitro* dry matter degradability.²³

The tdDM was not influenced by the addition of PE, as in a previous study that found a reduction in DM digestibility of soybean meal at pH 7.0 with the addition of tannic acid whereas abomasal digestibility was not affected.²³ Other studies have shown a decrease in *in vitro* DM degradation by rumen bacteria plus pepsin with the addition of tannins to soybean meal,²⁴ suggesting that some tannins remain bound and other *in vivo* studies have shown that feeding tannins to sheep increased fecal nitrogen excretion.²⁵

The effect of PE was greater for CP than DM, because of the strong hydrogen bond affinity of the

phenolic groups for the carbonyl oxygen of the peptide group.²⁶ All PE levels decreased the extent of CP degradation. Other studies^{11,27} have shown that quebracho-treated soybean meal decreased CP degradation but grape pomace tannins had no impact on ruminal CP degradability.⁸ Total degradability of CP did not differ between diets, suggesting that tannins did not reduce proteolysis, degradation of peptides or deamination of amino acids.⁸ PE-related effects were more noticeable in rumen fluid than under conditions simulating abomasal degradation due the stronger interaction between PE and proteins at a neutral pH, as the most stable tannin-protein complexes are formed between pH 4.0 and 7.0.26 Inclusion of PE increased pdBCP due to its interaction with protein thus affecting ruminal protein digestibility. Studies have shown that 300, 600 and 900 mg condensed tannins per g soybean meal protein increased bypass CP.¹¹ Other work reported that the addition of PE in in vitro conditions decreased ammonia, large and small peptides and amino acid levels.²⁸ Inhibition of amino acid deamination is a significant effect as it increases dietary protein bypass improving nitrogen efficiency.²⁹

Diets¹ p-Value L^3 Q^4 Protozoa PE0 PE1 PE2 PE3 PE4 PE5 SEM² Total 5.84 5.40 5.18 5.02 5.03 5.18 0.058 < 0.0001 < 0.0001 Isotricha 3.52 0.19 0.00 0.00 0.00 0.00 0.318 < 0.0001 < 0.0001 < 0.0001 Dasytricha 5.00 1.67 0.18 0.18 0.00 0.37 0.370 < 0.0001 Entodiniinae < 0.0001 0.0083 5.63 5.37 5.17 5.01 5.03 5.13 0.057 < 0.0001 < 0.0001 Diplodiniinae 2.22 0.19 0.00 0.00 0.00 0.37 0.259 Ophrioscolecinae 3.33 0.00 0.00 0.00 0.00 0.00 0.340 < 0.0001 < 0.0001

Table 7. Effects of PE level on protozoal concentration (log₁₀/g digesta).

¹Diets: PE0, control, no additive = 3.1 g TP/kg DM of basal diet; PE1, control + 0.65 = 3.75 g TP/kg DM of basal diet; PE2, control + 1.30 = 4.4 g TP/kg DM of basal diet; PE3, control + 1.95 = 5.05 g TP/kg DM of basal diet; PE4, control + 2.60 = 5.70 g TP/kg DM of basal diet; PE5, control + 3.25 = 6.35 g TP/kg DM of basal diet; ²SEM: standard error of the mean; ³L: linear; ⁴Q: quadratic.

Inclusion of PE increased dBCP compared to the control diet. Other research showed that increasing levels of tannin extract increased protein bypass degradability⁸ whereas another study¹¹ reported that increasing dietary levels of condensed tannin could inhibit postruminal CP degradation partly explaining the increase in fecal nitrogen. Another study²⁷ found that soybean meal treated with annins 10-250 g/kg of quebracho tannins, reduced intestinal crude protein digestibility of un-degraded protein but only at the highest level. In the present study, intestinal crude protein digestibility of un-degraded protein decreased at the highest PE level which was lower than other studies.²⁷ The apparent differences in intestinal digestibility of protein treated with PE may be associated with differences in its chemical structure and how this affects biological activity.¹¹ Differences in tannins and their ability to bind with proteins suggests that the post-ruminal reversibility of the process could also vary¹¹ (i.e., variability in the tannin's ability to decrease the ruminal CP degradation leading to differences in bypass CP reaching the duodenum).

Proteolytic bacteria numbers decreased with the addition of PE probably due to direct inhibition of rumen microbial function caused by interaction with the bacterial cell wall, or indirectly by decreasing protein availability (i.e., reducing ammonia (Table 6)). Other researchers³⁰ reported that tanniniferous feed lowered proteolytic bacterial numbers. Other work reported that *Sesbania sesban* tannins³¹ had no effect on proteolytic, peptidolytic or deaminative activity. Other studies indicate that the addition of concentrated pomegranate-residue extract increased total rumen bacteria population.³²

The effect of pomegranate peel extract on volatile fatty acids and ammonia-N production

VFAs are the final products of rumen microbial fermentation, and the main supply of energy for the ruminant.¹⁵ The PE3 level of 5.05 g TP/kg DM had no effect on total VFA and acetate concentrations suggesting that additives up to this level do not affect diet fermentability or energy availability, but higher levels decreased these parameters due to a decrease in DM degradation (Table 4). The major fermentation end products of protozoa are acetate and propionate¹⁵ so a reduction in acetate concentration may be due to a decrease in protozoa numbers (Table 7). Studies using other plant metabolites (50 and 100 g/kg DM of quebracho tannin or tannic acid) had no effect on total VFA in *in vitro* fermentations.³³

Results vary, the addition of *Leucaena leucocephala* and *Salix babylonica* extracts (0.6, 1.2, 1.8 mL extract/ g DM) increased VFA levels³ but Sainfoin hay extract (4.1 g catechin/kg DM substrate) and PE (2.8 g tannic acid/kg DM substrate) decreased total VFA production.²⁶

PE inclusion linearly decreased the acetate to propionate ratio and other studies using *P. kurroa* aqueous extract³⁴ and purified hydrolyzable (Chestnut and Sumach) and condensed tannins (Mimosa and Quebracho)³⁵ also showed a lower ratio. In contrast, the ratio increased with the addition of 50 g gallic acid and 100 g tannic acid per kg DM,³³ but Sainfoin hay extract (4.1 g catechin/kg DM substrate) and PE (2.8 g tannic acid/kg DM substrate) had no effect.²⁸ These inconsistencies can be attributed to an adaptation of the bacterial population and experimental conditions of studies, including diet type, animal and plant species, dose levels and the chemical structure of extract and the rumen fluid pH values.²⁶

The lower acetate to propionate ratio observed in this study was similar to that found with methane inhibitors.²⁹ Methane production was not directly measured but changes in fermentation end products suggest a reduction in methane production.

A reduction in NH_3-N concentration with increased levels of PE suggest an inhibitory effect on proteolytic activity (Table 5).³⁵ Additionally, decreased

ruminal NH₃–N concentrations are associated with the inhibition of protozoa,³⁶ probably as a consequence of reduced bacterial lysis.³⁷ *Entodinium* spp. protozoa are responsible for most ruminal bacterial breakdown³⁸ and in this study PE addition lowered *Entodinium* numbers (Table 7) leading to a decrease in NH₃–N concentration. NH₃–N concentration is also affected by ammonia binding to saponin-like compounds.²¹ Other research showed that 15, 30 g TP/Kg DM of PE decreased NH₃–N concentration.¹²

Enumeration of rumen protozoa

Total protozoa numbers, genera Dasytricha and Isotricha and subfamilies Entodiniinae, Diplodiniinae, and Ophrioscolecinae decreased with increasing PE levels in the diet. The antiprotozoal effect of PE is probably due to the phenolic structure of active metabolites (i.e., tannins and saponins) that interrupt protozoal membranes and inactivate enzymes depriving protozoa of substrates and metal ions that are vital for cell metabolism.³⁹ Data on the effects of PE on protozoa populations are not consistent. Castanea sativa wood extract containing hydrolyzable tannins (0.5 and 2.5 g tannin/kg DM) in rumen simulation experiments had no effect on total protozoa populations, Holotrichs or Entodiniomorphs.40 Another study12 showed a decrease in total protozoa but in other research⁴¹ total protozoa numbers increased with the addition of 200 mg/g of Enterolobium cyclocarpum (crude saponins, 19 mg/g) or 200 mg/g of Pithecellobium saman (crude saponins, 17 mg/g). Sanguisorba officinalis tannin extract (3 g tannin/kg DM substrate) increased Entodiniomorph sp. but did not affect Holotricha.42 These inconsistencies could be explained by experimental differences such as in vitro vs. in vivo, level and bioactivity of tannins used and genetic variation in ruminal protozoa (i.e., ruminal protozoal species differ in their sensitivity to tannins).

Conclusion

All levels of PE in the diet decreased ruminal degradation of protein without detrimentally affecting its intestinal digestion. Post-ruminal digestibility of ruminally un-degraded protein increased with the addition of PE with PE2 and PE3 levels conferring the best results. This study showed that water extraction of PE is as effective as the more expensive solvent extraction (Table 7). However, *in vivo* studies are needed to confirm the influence of PE on the efficiency of nitrogen utilization in ruminant animals.

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