

## Effect of polyethylene glycol on *in vitro* degradability of nitrogen and microbial protein synthesis from tannin-rich browse and herbaceous legumes

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Determination of microbial degradability of N is important in formulating a sound supplementation strategy for efficient utilisation of basal as well as supplementary diet components. *In vitro* degradability of N (IVDN) from tannin-containing browses (*Acacia cyanophylla*, *Acacia albida*, *Acacia barteri* and *Quercus ilex*) and two herbaceous legumes (*Desmodium intortum* and *Desmodium uncinatum*) was determined using the *in vitro* gas-production method coupled with NH<sub>3</sub>-N measurement in the presence and absence of a tannin-binding agent (polyethylene glycol (PEG), molecular mass 6000). Addition of PEG to tannin-containing feeds significantly ( $P < 0.05$ ) increased *in vitro* gas and short-chain fatty acid (SCFA) production, and IVDN. The use of PEG as a tannin-binding agent increased IVDN from 28 to 59, 32 to 72, 19 to 40, 32 to 73, 40 to 80, and 26 to 77% in *A. cyanophylla*, *A. albida*, *A. barteri*, *D. intortum*, *D. uncinatum* and *Q. ilex* respectively. There was significant correlation between total phenolic compounds (total phenol, TP; total tannin, TT) in leguminous forages and percentage increase in IVDN on addition of PEG ( $P < 0.05$ ;  $R^2$  0.70 and 0.82 for TP and TT respectively). The difference in IVDN observed in the absence and presence of PEG indicates the amount of protein protected from degradation in the rumen by tannins. When measured after 24 h incubation, tannin-containing feeds incubated in absence of PEG resulted in higher microbial protein synthesis than in the presence of PEG. Addition of PEG significantly ( $P < 0.05$ ) reduced the efficiency of microbial protein synthesis expressed as  $\mu\text{mol}$  purine/mmol SCFA.

### Polyethylene glycol: Protein degradability: Tannins: Microbial protein synthesis

Herbaceous and tree legumes are rich in N, and these form an integral part of livestock feed in dry seasons in many parts of tropical and subtropical regions where the major feed supply is based on crop residues. Results from *in vivo* (Campling *et al.* 1962; Ørskov & Grubb, 1978), nylon bag (Negi *et al.* 1988) and *in vitro* studies (Getachew *et al.* 1998c) indicated that the rumen-degradable N supply from crop residues was not sufficient to meet the maintenance requirement of animals. The nutritive value of crop residues can be enhanced by N supplementation in the form of leguminous forages. However, the concentration of phenolic compounds (particularly tannin) in multipurpose tree leaves and some herbaceous legumes is generally high; they may bind to protein thus rendering the protein undegradable by rumen microbes. The extent to which tannins in tropical browses and herbaceous legumes bind to proteins and make them undegradable by rumen microbes is not known.

Phenolics are compounds that possess one or more

hydroxyl (OH) substituents bonded onto an aromatic ring (Waterman & Mole, 1994). Tannins are naturally occurring polyphenolic compounds of high molecular mass (500 to 3000 Da) which form complexes with proteins. Tannins are grouped in two types: (1) the hydrolysable tannins consist of a carbohydrate moiety in which the hydroxyl groups are esterified with gallic acid or ellagic acid, and (2) the condensed tannins consist of oligomers of the flavan-3-ols (the catechins) and related flavanol residues which produce anthocyanidins on acid degradation (Mangan, 1988).

Tannins have both beneficial and adverse effects. Beneficial effects of tannin in ruminants include bloat suppression (Jones *et al.* 1973) and protection of dietary protein in the rumen and subsequently enhancement of amino acid absorption and utilisation by the ruminant animal (Waghorn *et al.* 1994). When tannin-containing herbage is masticated, tannin–protein complexes are formed; these are stable over the pH range 3.5–7.0 but

**Abbreviations:** IVDN, *in vitro* degradability of nitrogen; NDF, neutral-detergent fibre; NDS, neutral-detergent solution; PEG, polyethylene glycol; SCFA, short-chain fatty acids.

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dissociate in the abomasum and anterior duodenum. This protects the protein from microbial hydrolysis and deamination in the rumen and increases the proportion of feed proteins available for digestion and absorption post-rumen. Adverse effects of tannins include: (1) lower intake and digestibility; (2) inhibition of digestive enzymes; (3) loss of endogenous proteins; (4) systemic effects as a result of uptake of degraded products of hydrolysable tannins from digestive tract. Condensed tannins do not seem to be degraded by rumen microbes (Makkar *et al.* 1995c) and there is no absorption of condensed tannins from the gastrointestinal tract (Terrill *et al.* 1994). Polyethylene glycol (PEG), with a molecular mass of 4000, is a non-ionic detergent which forms complexes with hydrolysable and condensed tannins over a wide pH range (2–8.5) (Jones, 1965; Silanikove *et al.* 1996). PEG of various molecular masses (2000–35 000) have been used to prevent binding between tannins and proteins (Silanikove *et al.* 1994; Makkar *et al.* 1995b; Khazaal *et al.* 1996). PEG can also displace protein from a pre-formed tannin–protein complex (Barry & Manley, 1986).

Many workers have used chemical assays (Weiss *et al.* 1992) and nylon-bag techniques (Aufrere *et al.* 1991; Boila & Ingalls, 1992, 1995) to assess the protein degradability of conventional feeds. However, these methods are not applicable for tanniferous forages due to the binding of tannins to proteins. Ørskov & Ryle (1990) mentioned the possible underestimation of DM loss from the nylon-bag due to adherence of microbes and this effect could be much higher for tannin-containing feeds as tannins have high affinity to microbial proteins. Khazaal *et al.* (1994) compared the *in vitro* gas and nylon-bag techniques to evaluate nutritive value of feeds and concluded that the nylon-bag technique may not be suitable for evaluating feeds with anti-nutritive effects. Broderick (1987) developed an *in vitro* inhibitor technique to measure protein degradability from the appearance of breakdown products such as amino acids and NH<sub>3</sub> and the method has been used to study the protein degradability of conventional feeds (Broderick *et al.* 1992; Neutze *et al.* 1993). However Broderick & Albrecht (1997) using feeds containing phenolic compounds reported the binding of these compounds to free amino acids or peptides resulting in a negative rate of protein degradation. The *in vitro* gas production and NH<sub>3</sub>-N measurement have been used to assess the *in vitro* degradability of N (IVDN) in protein-rich diets (Raab *et al.* 1983) and in poor quality roughages (Getachew *et al.* 1998c).

The objectives of this present study were to evaluate the effect of PEG as a tannin-binding agent on microbial fermentation, efficiency of rumen NH<sub>3</sub>-N uptake and rumen microbial growth when tannin-rich feeds were incubated with buffered rumen fluid in the presence and absence of PEG, and to quantify the extent of tannin-mediated protection of protein from microbial degradation.

## Materials and methods

### *In vitro* gas production, true and apparent digestibility

Tannin-containing plants were selected to study the effect of addition of polyethylene glycol (molecular mass 6000) as a

tannin-binding agent since it was found to be the best amongst a number of tannin-binding agents (Makkar *et al.* 1995b). Leaves of two browse (*Acacia cyanophylla*, *Acacia albida*) two herbaceous legumes (*Desmodium intortum* and *Desmodium uncinatum*) were obtained from the International Livestock Research Institute (ILRI, Ethiopia) and *Acioa barteri* and *Quercus ilex* were sampled from Benin (West Africa) and Landesarboretum Hohenheim (Germany) respectively. The plants with known high-tannin content (Makkar *et al.* 1988, 1995b,d) were taken for this study. Samples were air-dried and ground to pass through 1 mm screen and used for the IVDN. Feed sample (0.5 g), starch (0.1 g) and PEG (1 g) were weighed into glass syringes in their respective order before the buffered rumen fluid was added. Rumen fluid was taken before morning feeding from a rumen-cannulated dairy cow receiving approximately 3 kg hay/d and straw *ad libitum*. Rumen fluid was collected into a pre-warmed thermos bottle, homogenised in a laboratory blender, strained using a nylon cloth of 100 µm and then filtered through glass wool. All handling was carried out under continuous flushing with CO<sub>2</sub>. The incubation volume was 40 ml. The N content of the buffered rumen fluid was 7.3 (SD 0.26) mg (*n* 28) in the 40 ml system. The details of procedures and composition of incubation medium are given in Getachew *et al.* (1998c) and Makkar *et al.* (1995b). For IVDN determination, two sets of incubations were carried out all in triplicate. First, tannin-containing feeds were incubated with and without starch as described by Raab *et al.* (1983), and second, the method described earlier was combined with the method outlined by Makkar *et al.* (1995b); the incubations were as in the first set except that PEG was included. This was repeated in four incubations. Raab *et al.* (1983) found no significant difference in protein degradability using one level and five levels of starch and therefore one level of starch was used in this present experiment.

After 24 h incubation, gas production was recorded and for determination of apparent digestibility, the contents of the syringes were transferred into centrifuge tubes and centrifuged at 23 000 g for 15 min at 4°C. The supernatant was pipetted into a 50 ml plastic bottle and stored in a freezer until it was analysed for NH<sub>3</sub>-N and SCFA. The syringes were washed twice with a total of 50 ml of 0.2% NaCl solution in order to quantitatively transfer the syringe contents into centrifuge tube. The washing solution was emptied each time from the syringes into the same centrifuge tube containing the pellet of the first centrifugation. The tubes were centrifuged again at 23 000 g for 15 min. The residual pellet was frozen and lyophilised in the tubes overnight. The residual moisture, if any, was removed by drying the tubes overnight at 60°C and then the tubes were weighed. Apparent digestibility was calculated from the weight of the centrifuge tube containing the pellet minus that of the empty tube and corrected for the pellet weight of the blank incubation. N in these pellets was determined and termed as apparent undegraded residue-N (APUR-N).

For true digestibility after 24 h incubation, the syringe contents from parallel sets of incubations were transferred into a 600 ml beaker and the syringes were washed twice with a total of 50 ml neutral-detergent solution (NDS; Van Soest & Robertson, 1985) and emptied into the beaker. The

contents were refluxed for 1 h, and then filtered through pre-tared filter crucibles. The crucibles were dried overnight at 100°C and weighed (Van Soest & Robertson, 1985). True degradability was calculated as the weight of substrate incubated minus the weight of the residue after NDS treatment. The residue after NDS treatment (neutral-detergent residue) was used for NDF-N determination. The ratio substrate truly degraded (mg) : gas (ml) volume, termed as partitioning factor (Blümmel *et al.* 1997b), was calculated using measured gas volume and truly degraded substrate after 24 h incubation of the feeds.

IVDN was calculated from the linear regressions of NH<sub>3</sub>-N concentration (*y*; mg) *v.* gas production (*x*; ml) observed on incubation of feed for 24 h with and without exogenous energy source (starch), as described in Raab *et al.* (1983). The intercept at the Y-axis represents the amount of NH<sub>3</sub>-N which is released when no gas was produced. The difference between the Y-intercept and NH<sub>3</sub>-N content in 0 h blank indicated the amount of NH<sub>3</sub>-N liberated from the degradation of protein and other N-containing compounds of the feedstuff incubated. The NH<sub>3</sub>-N in the 24 h incubated blank was higher than the 0 h blank values due to microbial lysis that occurred during incubation leading to under-estimation of IVDN (Getachew *et al.* 1998b). Therefore 0 h blank NH<sub>3</sub>-N values were used in the calculation of IVDN. The IVDN at 24 h was calculated as a proportion of total N incubated by the equation:

IVDN =

$$\frac{\text{NH}_3\text{-N at zero gas production} - \text{NH}_3\text{-N in 0 h blank}}{\text{Total N of feed incubated}}$$

NH<sub>3</sub>-N disappearance with increasing energy (starch) availability was assumed to be due to its incorporation into microbial protein; therefore, the decline in NH<sub>3</sub>-N/ml gas produced (slope; rate of NH<sub>3</sub>-N uptake) is defined as the efficiency of microbial protein synthesis (Raab *et al.* 1983).

#### Microbial nitrogen

Microbial N was measured using three different methods: gravimetric (the difference in weight between truly and apparently degraded feed) (Blümmel *et al.* 1997b), N

balance, and purine determination. In the Nitrogen balance method two approaches were used. The first approach was:

$$\text{MN} = \text{TN} - (\text{NDF-N} + \text{NH}_3\text{-N}),$$

where MN is the microbial N, TN is total N (feed N + N in buffered rumen fluid) in the syringe before incubation, and the NDF-N and NH<sub>3</sub>-N are NDF-N and NH<sub>3</sub>-N in the supernatant after 24 h incubation.

In a closed system, the N present before incubation (total N) will be in the form of microbial mass, NDF-N, amino acids and NH<sub>4</sub><sup>+</sup> after fermentation. Negligible amounts of amino acids and peptides are present in the supernatant during fermentation (Krishnamoorthy *et al.* 1990) and therefore these can be ignored in calculation of microbial N.

The second approach was:

$$\text{MN} = \text{APUR-N} - \text{NDF-N},$$

where MN is the microbial N, APUR-N is apparent undegraded residue-N after incubation. The basis of this approach was similar to that of Blümmel *et al.* (1997b). The rumen microbial sample was prepared using the method described by Makkar & Becker (1998) to assess the relationship of purine:N. The efficiency of microbial protein synthesis was defined as μmol purine/mmol SCFA.

#### Chemical analysis

DM and ash were determined according to the method of the Association of Official Analytical Chemists (1990). NDF and acid-detergent fibre were determined following the method of Van Soest & Robertson (1985). N in the feeds, apparent undegraded residues and lyophilised rumen microbes was determined by using Kjeldahl procedure. NH<sub>3</sub>-N in the supernatant was also determined using Kjeldahl procedure by distilling 15 ml of the supernatant under alkaline condition (Makkar & Becker, 1996). Total phenol was determined by Folin–Ciocalteu reagents, total tannins as the difference of phenolics before and after tannin removal with and without the use of insoluble polyvinylpyrrolidone (Makkar *et al.* 1993), and condensed tannin using the butanol–HCl–Fe reagent (Porter *et al.* 1986). Total phenols and tannins were expressed as tannic acid

**Table 1.** Chemical composition (% dry matter) of browse and herbaceous legume species used in the present experiment

Species	Components						
	CP	NDF	ADF	Ash	TP*	TT*	CT†
<i>Acacia saligna</i>	14.9	40.0	46.1	10.4	13.4	10.3	10.0
<i>Acacia albida</i>	9.0	35.4	43.8	9.0	12.0	11.1	7.9
<i>Acacia barberi</i>	8.6	64.2	64.6	10.9	9.7	8.5	14.8
<i>Desmodium intortum cv greenleaf</i>	19.9	44.7	53.2	9.8	8.5	7.6	10.7
<i>Desmodium uncinatum cv silverleaf</i>	19.7	47.5	55.0	10.0	8.6	7.7	8.9
<i>Quercus ilex</i>	10.0	50.3	50.0	5.0	18.5	16.8	7.1

CP, crude protein; NDF, neutral-detergent fibre; ADF, acid-detergent fibre; TP, total phenol; TT, total tannin; CT, condensed tannin.

\* Tannic acid equivalent.

† Leucocyanidin equivalent.

**Table 2.** *In vitro* gas production, ammonia-nitrogen (NH<sub>3</sub>-N) in 40 ml supernatant†, nitrogen in apparent undegraded residue (APUR-N), *in vitro* dry matter degradability, and partitioning factor (PF) from different browse and herbaceous legumes after 24 h incubation with and without polyethylene glycol (PEG) and/or starch in buffered rumen fluid  
(Mean values and standard deviations)

Feeds	24 h gas (ml)		APUR-N (mg)		NH <sub>3</sub> -N (mg)		Dry matter degradability (%)				PF (mg/ml gas)	
	Mean	SD	Mean	SD	Mean	SD	Apparent		True		Mean	SD
							Mean	SD	Mean	SD		
<i>Acacia cyanophylla</i>	34.6	1.34	12.0	0.23	6.9	0.20	21.7	1.14	68.2	0.49	9.2	0.07
<i>Acacia</i> + PEG	66.1	1.72	8.1	0.12	10.5	0.98	23.0	2.27	59.5	1.33	4.2	0.09
<i>Acacia</i> + starch	72.1	2.51	14.9	0.38	4.4	0.38	29.7	0.48	73.1	2.28	5.6	0.17
<i>Acacia</i> + PEG + starch	105.6	3.37	9.5	0.26	8.9	0.94	32.3	1.13	63.9	0.88	3.4	0.04
ANOVA												
PEG	****		****		****		*		****		****	
Starch	****		****		****		****		***		****	
PEG × Starch	NS		****		*		NS		NS		****	
<i>Acacia albida</i>	54.2	0.77	18.5	0.36	8.6	0.59	32.4	1.63	75.2	2.40	6.5	0.21
<i>Acacia</i> + PEG	71.1	1.59	12.3	0.22	14.9	1.10	35.9	0.70	74.4	3.55	4.9	0.23
<i>Acacia</i> + starch	92.5	3.15	21.4	0.31	5.7	0.58	38.8	1.46	79.5	0.83	4.8	0.05
<i>Acacia</i> + PEG + starch	109.8	2.37	15.1	0.26	11.9	0.87	42.1	1.23	77.2	1.66	3.9	0.09
ANOVA												
PEG	****		****		****		***		NS		****	
Starch	****		****		****		****		*		****	
PEG × Starch	NS		NS		NS		NS		NS		*	
<i>Acacia barteri</i>	12.7	0.16	8.4	0.47	7.3	0.22	6.0	2.27	45.0	1.45	16.1	0.53
<i>Acacia</i> + PEG	27.1	0.91	6.9	0.46	8.1	0.64	-6.4	2.00	33.5	2.64	5.6	0.44
<i>Acacia</i> + starch	51.8	1.56	9.9	0.50	5.1	0.51	17.1	0.63	52.9	0.13	5.6	0.02
<i>Acacia</i> + PEG + starch	68.9	0.87	8.6	0.47	6.1	0.56	7.0	0.85	42.5	1.44	3.4	0.11
ANOVA												
PEG	****		****		****		****		****		****	
Starch	****		****		****		****		****		****	
PEG × Starch	*		NS		NS		NS		NS		****	
<i>Desmodium introtum</i>	70.3	0.59	14.3	0.39	8.6	0.12	34.8	3.19	70.9	2.38	4.7	0.16
<i>Desmodium</i> + PEG	84.3	1.99	9.3	0.18	13.5	0.14	32.9	0.42	68.9	2.73	3.8	0.15
<i>Desmodium</i> + starch	111.5	1.57	16.1	0.43	6.7	0.45	39.5	1.77	75.7	2.86	3.8	0.14
<i>Desmodium</i> + PEG + starch	125.5	2.24	10.9	0.86	11.4	0.43	40.6	3.37	71.2	1.25	3.2	0.05
ANOVA												
PEG	****		****		****		NS		*		****	
Starch	****		****		****		***		*		****	
PEG × Starch	NS		NS		NS		NS		NS		NS	
<i>Desmodium uncinatum</i>	59.4	2.05	14.6	1.06	8.3	0.24	30.1	0.72	70.8	1.02	5.6	0.08
<i>Desmodium</i> + PEG	80.2	0.68	8.5	0.13	13.9	0.23	30.8	0.60	71.2	5.06	4.2	0.30
<i>Desmodium</i> + starch	86.5	3.59	15.8	0.61	6.2	0.25	38.6	0.53	75.5	1.34	4.9	0.09
<i>Desmodium</i> + PEG + starch	120.8	1.69	10.6	0.34	11.3	0.37	37.4	0.48	67.7	5.36	3.1	0.25
ANOVA												
PEG	****		****		****		NS		NS		****	
Starch	****		****		****		****		NS		****	
PEG × Starch	****		*		*		*		NS		NS	
<i>Quercus ilex</i>	49.6	1.63	9.9	0.22	5.6	0.08	25.9	0.54	58.6	2.16	5.6	0.20
<i>Quercus</i> + PEG	65.3	0.69	7.6	0.24	7.6	0.16	25.8	0.89	59.6	2.30	4.3	0.17
<i>Quercus</i> + starch	88.4	1.42	11.6	0.30	3.1	0.09	33.7	0.88	62.0	3.50	4.0	0.22
<i>Quercus</i> + PEG + starch	103.5	1.84	9.9	0.47	4.6	0.13	32.9	0.74	67.1	1.13	3.7	0.06
ANOVA												
PEG	****		****		****		NS		NS		****	
Starch	****		****		****		****		**		****	
PEG × Starch	NS		*		****		NS		NS		*	

PF, partitioning factor (substrate truly degraded).

\*  $P < 0.05$ , \*\*  $P < 0.01$ , \*\*\*  $P < 0.001$ , \*\*\*\*  $P < 0.0001$ .

† After centrifugation of syringe contents (see p. 74).

equivalent and condensed tannins as leucocyanidin equivalent. SCFA in the supernatant were determined using GC (Hewlett Packard, 5880A, Avondale, PA, USA) fitted with a flame-ionisation detector. Purine determination in apparent undegraded residues and rumen microbes was carried out using the HPLC method described by Makkar & Becker (1999).

### Statistical analysis

The data were subjected to ANOVA for two factor design to assess the effect of PEG and starch and their interactions using the General Linear Model available in Statistical Analysis Systems (SAS/STAT program, SAS Institute Inc., Cary, NC, USA). The differences between means

were compared using Duncan's multiple range test. The IVDN and N uptake in the presence and absence of PEG was compared using Student's *t* test.

## Results

### *Chemical composition of the feeds, in vitro gas production, rumen ammonia concentration, and true and apparent degradability*

The chemical composition of the feeds used in the experiment is given in Table 1. Acid-detergent fibre values were higher than the NDF values. The contents of total phenols, total tannins and condensed tannins ranged from 8.5 to 18.5%, 7.6 to 16.8%, and 7.1 to 14.8% respectively. There was a large variation in crude protein content of feeds.

*In vitro* gas production, apparent undegraded residue-N, NH<sub>3</sub>-N in the supernatant, and apparent and true degradability of feeds are shown in Table 2. Addition of PEG to tannin-containing feeds increased *in vitro* gas production in all feeds. There was significant interaction ( $P < 0.05$ ) for *in vitro* gas production between PEG and starch for *A. barteri* and *D. uncinatum*. The increases ranged between 20% for *D. intortum* to 113% *A. barteri*. Addition of readily-soluble carbohydrate (starch) increased true digestibility. The addition of PEG reduced N in apparent undegraded residues in all feeds. In *A. albida*, the interaction between PEG and starch was not significant ( $P > 0.05$ ) for all variables except the partitioning factor. NH<sub>3</sub>-N in the supernatant was higher ( $P < 0.05$ ) for samples incubated in presence of PEG as compared to its absence. Inclusion of starch significantly ( $P < 0.05$ ) depressed NH<sub>3</sub>-N in the supernatant. Despite considerable increase in gas production in presence of PEG, the apparent and true degradability were not significantly different in the presence and absence of PEG. The partitioning factor ranged from 3.1 to 16.1.

### *In vitro degradability of nitrogen*

The IVDN ranges were 19–40% in tannin-containing feed alone and 40–80% in tannin-containing feed + PEG (Table 3). Addition of PEG increased the availability of protein to rumen microbes by more than 100%. The extent

**Table 3.** *In vitro* degradability of nitrogen (IVDN) after 24 h incubation in the presence and absence of polyethylene glycol (PEG) (Mean values and standard deviations)

Species	IVDN†				<i>n</i>
	–PEG		+PEG		
	Mean	SD	Mean	SD	
<i>Acacia cyanophylla</i>	0.28	0.08	0.59*	0.17	4
<i>Acacia albida</i>	0.32	0.07	0.72***	0.07	4
<i>Acacia barteri</i>	0.19	0.05	0.40	0.17	4
<i>Desmodium intortum</i>	0.32	0.08	0.73***	0.09	4
<i>Desmodium uncinatum</i>	0.40	0.04	0.80***	0.04	4
<i>Quercus ilex</i>	0.26	0.08	0.77***	0.05	4

Mean values were significantly different from control group (–PEG), \*  $P < 0.05$ , \*\*\*  $P < 0.001$  (Student's *t* test).

† Proportion of feed nitrogen incubated.

**Table 4.** Relationship between increase in *in vitro* degradability of nitrogen (IVDN) and *in vitro* gas production in presence of polyethylene glycol (PEG), with contents of phenolic compounds

Phenolic compounds	Increase in IVDN (%)		Increase in <i>in vitro</i> gas (%)	
	<i>R</i> <sup>2</sup>	<i>n</i>	<i>R</i> <sup>2</sup>	<i>n</i>
Total phenols	0.698*	6	0.040	6
Total tannins	0.817*	6	0.004	6
Condensed tannins	0.268	6	0.558	6

Mean values were significantly different from control group, \*  $P < 0.05$ .

of increase in the microbial degradability of N on addition of PEG indicates the amount of protein protected by tannins from microbial degradation. There was significant correlation between the content of phenolic compound and percentage increase in IVDN (Table 4). There was no significant correlation ( $P > 0.05$ ;  $R^2$  0.01) between the combined effect of phenolic content and *in vitro* gas production. The relationship between the joint effect of phenolic content and IVDN was weak but significant ( $P < 0.05$ ;  $R^2$  0.22). Although the overall N degradability in the medium was higher in presence of PEG, there was no significant difference ( $P > 0.05$ ) in the rate of NH<sub>3</sub>-N uptake by rumen microbes (Table 5). When individual feed was considered, there was significant difference ( $P < 0.05$ ) in the rate of NH<sub>3</sub>-N uptake in the presence and absence of PEG for *A. cyanophylla*, *D. intortum* and *Q. ilex*. The rate of NH<sub>3</sub>-N uptake tended to be higher when PEG was absent (0.0683 v. 0.0606 mg NH<sub>3</sub>-N uptake/ml gas in absence and presence of PEG respectively; Table 5).

### *Short-chain fatty acid production*

The net total and individual SCFA production values on incubation of tannin-containing feeds with and without PEG are shown in Table 6. Addition of PEG significantly ( $P < 0.001$ ) increased production of total SCFA (from 38% in *D. intortum* and to 540% in *A. barteri*). Incubation of tannin-containing feed with PEG tended to decrease the molar proportion of propionate. Addition of PEG also

**Table 5.** Rate of nitrogen uptake (mg N uptake/ml gas) by rumen microbes on addition of starch when tannin-rich browse and herbaceous legumes were incubated with and without polyethylene glycol (PEG) (Mean values and standard deviations)

Feeds	N uptake (mg/ml gas)			
	–PEG		+PEG	
	Mean	SD	Mean	SD
<i>Acacia cyanophylla</i>	0.0878	0.0258	0.0462*	0.0159
<i>Acacia albida</i>	0.0762	0.0159	0.0759	0.0086
<i>Acacia barteri</i>	0.0543	0.0103	0.0459	0.0035
<i>Desmodium intortum</i>	0.0476	0.0142	0.0540	0.0139
<i>Desmodium uncinatum</i>	0.0781	0.0082	0.0621*	0.0072
<i>Quercus ilex</i>	0.0658	0.0070	0.0794*	0.0048

Mean values were significantly different from control group (–PEG), \*  $P < 0.05$  (Student's *t* test).

**Table 6.** Net total and individual short-chain fatty acid (SCFA) production ( $\mu\text{mol}/40\text{ ml}$  incubation medium) on incubation (24 h) of browse and herbaceous legumes with and without polyethylene glycol (PEG) and/or starch  
(Mean values and standard deviations)

Feeds	Total SCFA		C2		C3		C4		C4i		C5i		C5	
	Mean	SD	Mean	SD	Mean	SD	Mean	SD	Mean	SD	Mean	SD	Mean	SD
<i>Acacia cyanophylla</i>	856	25.4	609	20.0	284	6.1	-30.7	4.0	-3.7	0.46	-10.9	0.69	8.0	0.92
<i>Acacia</i> + PEG	1610	128.7	1089	91.7	439	30.7	21.3	4.0	15.5	0.46	24.9	2.01	20.5	1.06
<i>Acacia</i> + starch	1374	12.9	883	23.1	503	20.1	-4.7	2.8	-3.9	1.29	-11.2	1.67	7.3	1.83
<i>Acacia</i> + starch + PEG	2135	127.0	1369	84.9	551	31.1	145.3	5.7	19.4	0.85	27.9	2.83	21.7	1.70
ANOVA														
PEG	****		****		****		****							
Starch	****		****		****		****							
PEG $\times$ Starch	NS		NS		*		****							
<i>Acacia albida</i>	1103	28.6	849	28.3	237	0.0	9.3	0.0	1.4	0.28	-1.3	0.0	7.3	0.00
<i>Acacia</i> + PEG	1648	43.0	1183	30.6	328	10.1	46.6	2.3	22.5	0.46	40.5	2.31	28.0	0.61
<i>Acacia</i> + starch	1676	31.2	1169	20.0	393	13.9	105.3	4.0	2.1	0.23	-3.2	1.22	9.1	0.23
<i>Acacia</i> + starch + PEG	2410	24.8	1643	23.1	489	6.9	184.0	6.1	25.3	0.23	41.9	0.00	27.1	0.23
ANOVA														
PEG	****		****		****		****							
Starch	****		****		****		****							
PEG $\times$ Starch	*		*		NS		****							
<i>Acacia barteri</i>	62	5.7	87	2.8	15	2.8	-30.7	0.0	-2.2	0.28	-4.5	1.13	-3.7	0.85
<i>Acacia</i> + PEG	395	45.2	316	30.6	84	8.3	-12.0	4.6	2.9	0.61	5.7	1.22	-1.7	0.46
<i>Acacia</i> + starch	720	34.4	429	34.6	291	8.3	-2.7	0.0	-1.5	0.83	-2.9	1.20	-1.7	0.46
<i>Acacia</i> + starch + PEG	1224	59.6	803	46.2	265	4.0	136.0	10.1	5.4	3.11	8.1	3.68	1.1	0.85
ANOVA														
PEG	****		****		*		****							
Starch	****		****		****		****							
PEG $\times$ Starch	*		*		****		****							
<i>Desmodium intortum</i>	1646	2.5	1181	0.0	342	2.8	76.0	0.0	10.6	0.28	14.2	0.28	21.8	0.28
<i>Desmodium</i> + PEG	2268	18.9	1588	11.5	442	6.1	126.7	2.3	28.9	0.61	44.0	0.00	38.0	0.00
<i>Desmodium</i> + starch	2456	42.0	1635	23.1	552	8.0	205.3	9.2	15.1	0.83	18.9	1.40	29.9	0.61
<i>Desmodium</i> + starch + PEG	2815	112.1	1888	75.7	576	17.4	236.0	14.4	31.5	0.92	44.0	4.00	39.3	2.31
ANOVA														
PEG	****		****		****		****							
Starch	****		****		****		****							
PEG $\times$ Starch	*		*		****		NS							
<i>Desmodium uncinatum</i>	1413	29.6	1008	23.1	305	6.1	68.0	0.0	7.7	0.23	6.3	1.29	17.7	0.92
<i>Desmodium</i> + PEG	2054	68.5	1435	46.2	399	12.9	124.0	6.9	26.3	1.01	40.7	1.15	29.6	1.20
<i>Desmodium</i> + starch	2178	11.3	1421	0.0	531	8.3	181.3	2.3	10.9	0.23	11.6	0.69	21.9	0.46
<i>Desmodium</i> + starch + PEG	2869	163.0	1915	100.7	596	36.7	248.0	14.4	31.5	2.27	44.7	6.43	34.7	2.95
ANOVA														
PEG	****		****		****		****							
Starch	****		****		****		****							
PEG $\times$ Starch	NS		NS		NS		NS							
<i>Quercus ilex</i>	1120	13.1	835	11.5	221	2.3	56.0	0.0	1.6	0.0	1.3	0.61	5.6	0.00
<i>Quercus</i> + PEG	1660	54.1	1215	41.6	300	10.6	109.3	2.3	10.0	0.69	14.5	1.22	11.6	0.00
<i>Quercus</i> + starch	1847	48.8	1221	40.0	531	6.1	82.7	2.3	2.9	0.23	2.4	1.39	7.5	0.61
<i>Quercus</i> + starch + PEG	2249	42.4	1535	30.6	529	8.3	146.7	4.6	10.7	0.46	14.0	1.39	13.9	0.23
ANOVA														
PEG	****		****		****		****							
Starch	****		****		****		****							
PEG $\times$ Starch	*		NS		****		*							

C2, acetate; C3, propionate; C4, butyrate; C4i, isobutyrate; C5i, isovalerate; C5, valerate.  
\*  $P < 0.05$ , \*\*\*  $P < 0.001$ , \*\*\*\*  $P < 0.0001$ .

increased the concentration of isoacids. In the absence of PEG there was net uptake of isoacids which is the result of protection of proteins from rumen deamination by tannins. Addition of PEG also increased production of butyrate.

#### Microbial protein synthesis

The relationship of purine ( $\mu\text{mol}$ ): N (mg) was found to be 1.4 (SD 0.021) ( $n$  6). Microbial N measured using the different methods is presented in Table 7. The gravimetric method was poorly correlated with the N balance method.

There was strong correlation ( $R^2$  0.98) between microbial protein synthesis measured using the two N balance methods. Addition of PEG to tannin-containing feeds increased rumen  $\text{NH}_3$ -N concentration but reduced microbial protein synthesis. Incubation of starch with tannin-containing feeds significantly increased microbial protein synthesis. Incubation of tannin-containing feed in combination with PEG and starch yielded higher microbial protein synthesis than with PEG alone. The efficiency of microbial protein synthesis was significantly higher ( $P < 0.05$ ) in tannin-containing feed than tannin-containing feed + PEG.

**Table 7.** Microbial protein synthesis on incubation of tannin-rich browses and herbaceous legumes (24 h incubation) in the presence and absence of polyethylene glycol (PEG) and/or starch

Feeds	Microbial N measured (mg)									
	Nitrogen balance†								EMPS ( $\mu$ mol purines/ mmol SCFA)	
	Method 1		Method 2		Gravimetric‡		Purines§			
	Mean	SD	Mean	SD	Mean	SD	Mean	SD	Mean	SD
<i>Acacia cyanophylla</i>	11.1	0.12	10.3	0.19	16.3	0.71	5.6	0.15	6.6	0.03
<i>Acacia</i> + PEG	8.5	0.73	6.8	0.09	12.7	0.27	4.2	0.17	2.6	0.30
<i>Acacia</i> + starch	13.9	0.13	13.1	0.33	18.2	1.16	9.4	0.31	6.8	0.26
<i>Acacia</i> + PEG + starch	10.5	0.27	8.1	0.50	13.1	0.10	6.7	0.23	3.2	0.19
ANOVA										
PEG	****		****				****		****	
Starch	****		****				****		*	
PEG $\times$ Starch	NS		*				*		NS	
<i>Acacia albida</i>	16.4	0.54	14.9	0.27	15.6	1.05	7.0	0.36	6.4	0.61
<i>Acacia</i> + PEG	10.7	0.47	9.6	0.30	13.9	1.27	6.0	0.06	3.7	0.10
<i>Acacia</i> + starch	18.8	0.80	17.8	0.18	17.6	0.18	11.0	0.50	6.6	0.37
<i>Acacia</i> + PEG + starch	13.7	1.87	12.6	1.50	15.1	0.46	9.7	0.40	4.0	0.16
ANOVA										
PEG	****		****				**		****	
Starch	*		****				****		NS	
PEG $\times$ Starch	NS		NS				NS		NS	
<i>Acacia barteri</i>	4.0	0.04	5.0	0.07	13.9	0.40	3.0	0.27	48.8	0.01
<i>Acacia</i> + PEG	2.6	0.23	3.4	0.25	15.1	0.55	3.0	0.06	7.7	0.88
<i>Acacia</i> + starch	6.2	0.21	7.0	0.27	15.0	0.23	8.9	0.32	12.6	1.08
<i>Acacia</i> + PEG + starch	4.3	0.24	4.9	0.20	14.9	0.59	5.7	0.20	4.7	0.38
ANOVA										
PEG	****		****				****		****	
Starch	****		****				****		****	
PEG $\times$ Starch	NS		NS				****		***	
<i>Desmodium intortum</i>	14.3	0.08	13.0	0.30	13.1	0.94	5.9	0.36	3.5	0.09
<i>Desmodium</i> + PEG	9.6	0.08	8.7	0.12	13.0	0.84	5.4	0.10	2.4	0.02
<i>Desmodium</i> + starch	15.6	0.14	14.6	0.16	15.6	1.22	8.9	0.46	3.6	0.17
<i>Desmodium</i> + PEG + starch	11.3	0.23	9.2	0.70	13.2	1.03	7.9	0.44	2.8	0.12
ANOVA										
PEG	****		****				*		****	
Starch	****		****				****		*	
PEG $\times$ Starch	*		NS				NS		NS	
<i>Desmodium uncinatum</i>	14.0	0.10	13.8	0.88	13.5	0.48	6.5	0.30	4.6	0.25
<i>Desmodium</i> + PEG	8.9	0.30	7.7	0.10	15.1	0.13	5.2	0.12	2.5	0.14
<i>Desmodium</i> + starch	16.4	0.17	14.6	0.46	17.6	0.68	9.8	0.29	4.5	0.16
<i>Desmodium</i> + PEG + starch	10.9	0.36	9.4	0.28	15.0	1.72	8.2	0.18	2.9	0.22
ANOVA										
PEG	****		****				****		****	
Starch	****		*				****		NS	
PEG $\times$ Starch	NS		NS				NS		NS	
<i>Quercus ilex</i>	9.8	0.10	9.0	0.13	12.0	0.65	5.0	0.17	4.5	0.19
<i>Quercus</i> + PEG	7.9	0.10	6.7	0.37	12.5	0.80	4.4	0.59	2.6	0.43
<i>Quercus</i> + starch	12.3	0.10	10.6	0.38	11.4	0.31	9.2	0.06	5.0	0.16
<i>Quercus</i> + PEG + starch	10.8	0.03	9.1	0.24	14.9	0.33	9.1	0.06	4.0	0.10
ANOVA										
PEG	****		****				*		****	
Starch	****		****				****		***	
PEG $\times$ Starch	*		*				NS		*	

EMPS, efficiency of microbial protein synthesis.

\*  $P < 0.05$ , \*\*  $P < 0.01$ , \*\*\*  $P < 0.001$ , \*\*\*\*  $P < 0.0001$ .† Method 1, total N in the system – neutral-detergent fibre-N +  $\text{NH}_3$ -N in the supernatant; method 2, apparent undegraded residue-N – neutral-detergent fibre-N (see p. 78).

‡ Converted to microbial N assuming an average N content in microbial dry matter to be 7.7% (Ørskov &amp; Ryle, 1990).

§  $\mu$ mol in apparent residues.

## Discussion

### Chemical composition of the feed

The crude protein content of feeds was within the range reported by Rittner & Reed (1992). Rittner & Reed (1992) mentioned high variation in crude protein content of tropical

browse species (4.4–38%). The condensed tannin content was above the level suggested (Barry *et al.* 1986) to yield positive effects by improving the efficiency of N digestion. The sugar content of the feeds used in the present study was not analysed. However, Ford (1978) reported < 5% sugars in *Desmodium intortum*. *Leucaena leucocephala* and

*Gliricidia sepium* had total sugar contents of 12 and 14 % respectively (Vadiveloo & Fadel, 1992).

#### *In vitro* degradability of nitrogen

Incubation was carried out in N-rich medium (approximately 182 mg N/l). This value was within the level suggested for optimum rumen fermentation (Ørskov, 1982). Using similar incubation medium, the IVDN values observed from browse and herbaceous legumes in the absence of PEG were lower than those reported for low quality roughages (Getachew *et al.* 1998c). However, due to their relatively high crude protein content, browses and herbaceous legumes could play a significant role in supplying rumen degradable N. In the presence of PEG, the IVDN values were higher. The difference between IVDN values observed in presence and absence of PEG indicates the amount of protein protected by tannins from degradation in the rumen. Whether this amount of protein protected by tannins from microbial degradation is fully available to animals post-ruminally requires further research. In the current experiment the IVDN was measured after 24 h incubation. Raab *et al.* (1983) reported a close relationship between *in vivo* and *in vitro* values when incubation was terminated after 17 h. When conventional-protein feed was used, about 80 % of the 24 h value was degraded in the first 8 h incubation whereas in protected protein feed only 60 % of the value measured after 24 h incubation were degraded (Raab *et al.* 1983). The appropriate incubation time for *in vitro* degradability studies may depend on the rate of degradability of protein. The presence of phenolic compounds reduces the rate of degradation of feeds (Khazaal *et al.* 1994; Broderick & Albrecht, 1997) making the choice of incubation time more complicated. Kaitho *et al.* (1998) used a range of browse species with different level of phenolic compounds and reported a degradation rate of -0.1–19.0 %/h. Broderick & Albrecht (1997) using the *in vitro* inhibitor system for tannin-containing feeds reported low rates of degradability of protein. The rate of protein degradability can be affected not only by the amount of phenolic compounds but also by their biological activity (Broderick & Albrecht, 1997). Condensed tannins from different plant species have been reported to show different physical and chemical properties (Mangan, 1988).

#### *In vitro* gas production, apparent and true degradability, and partitioning of nutrients

Phenolic compounds (particularly tannins) bind to macromolecules, in particular to proteins, forming tannin-protein complexes and creating difficulties in the analysis of fibre using the detergent method (Makkar *et al.* 1995d). This was also evident from the higher value of acid-detergent fibre as compared to NDF (Table 1). Since tannins are heterogeneous in nature their biological activity may depend on the type of tannins and the degree of polymerisation (Makkar *et al.* 1988). Despite the large increase in gas production, apparent digestibility in presence of PEG was not significantly ( $P > 0.05$ ) different from that observed in

absence of PEG mainly due the tannin-PEG complexes which became insoluble in the NDS, distorting the weight of degraded sample (Makkar *et al.* 1995b,d). For conventional feeds (roughages), the ratio substrate truly degraded: gas volume, termed as the partition factor, ranges from 2.74–4.65 mg/ml (Blümmel *et al.* 1997a). A partitioning factor as high as 16.1 has been obtained for *A. barteri* (Table 2) which is well beyond the theoretical (2.75–4.41) or observed (2.74–4.65) ranges of partitioning factors (Blümmel *et al.* 1997a). This result also reinforces the observation of Makkar *et al.* (1997) that due to various artefacts the partitioning factors can not be determined for tannin-rich forages by the approach of residue determination using NDS. This high partitioning factor could be due to: (1) leaching of tannins from the feed during fermentation contributing to the DM loss but without contributing to the gas production; (2) inhibition of cell solubles by tannins; or a combination of (1) and (2) (Makkar *et al.* 1997; Getachew *et al.* 1998a). All these results suggest again that the detergent system of fibre analysis should be used with caution for tannin-containing feeds.

Large increases in *in vitro* gas and SCFA production due to the addition of PEG in this present study confirms that the effect of tannin is on depression of DM degradation. Norton (1994) summarised data on legume supplementation to poor quality roughage and reported that the DM digestibility in legumes ranged from 36–60 % and these values were not significantly different from the values for basal diet. This indicates that the effect of tannins in legumes is not only in reducing N availability but also to reduce digestibility of other components.

#### *Microbial protein synthesis*

Microbial N determined by both N balance methods (see p. 75) take into account the NDF-N. Our previous study (Makkar *et al.* 1995d) showed that NDF or NDF-N values are distorted to a greater extent if the apparent undegraded residue is subjected to NDS treatment rather than subjecting the syringe contents after fermentation directly to the NDS treatment. In the present study, NDF-N was determined in the NDF residue obtained on subjecting the syringe contents after fermentation directly to the NDS treatment. In addition, had the NDF-N values been affected by artefacts arising from the presence of tannin-microbial protein complexes, the observed partitioning factor should not have been higher than the theoretical range. Therefore, an error, if any, in the NDF-N values is not expected to be large. The higher microbial N by the gravimetric method than with the N balance methods can be explained by the presence of tannins and/or tannin-protein complexes in the apparently undegraded residue (Makkar *et al.* 1995d) leading to an overestimation of microbial mass and hence of microbial N. The conclusions drawn from the microbial N estimated by these methods are not affected by the factors mentioned earlier since the values obtained under different conditions (with or without PEG and starch) for a tannin-rich feed had a similar pattern using any of the three approaches. In addition, the pattern observed for microbial N was similar to that of purines (Table 7); purines content is an indicator of microbial protein.

### *Ammonia-nitrogen concentration*

Addition of PEG increased the  $\text{NH}_3\text{-N}$  concentration in the supernatant. In tannin-containing feeds, degradability of proteins was markedly depressed by tannins resulting in low  $\text{NH}_3\text{-N}$  concentration. The increased  $\text{NH}_3\text{-N}$  in presence of PEG did not increase microbial protein synthesis, probably due to poor synchronisation between N release and carbohydrate fermentation. An increase in microbial protein synthesis when tannin-containing feeds were incubated with starch indicated that uncoupled fermentation occurred when tannin-containing feed was incubated with PEG leading to accumulation of  $\text{NH}_3\text{-N}$ . A rapid degradation of N not matched to the availability of energy could lead to accumulation of  $\text{NH}_3\text{-N}$  in the *in vitro* system or to a high absorption of  $\text{NH}_3\text{-N}$  from the rumen *in vivo*. *In vivo*, the  $\text{NH}_3\text{-N}$  not captured in the rumen is absorbed and converted into urea, and the synthesis of urea in the liver requires expenditure of energy, each mole of urea requiring four moles of ATP (Van Soest, 1994). Synchronisation of the rate of degradation of N and carbohydrate components in the rumen is extremely important for efficient utilisation of rumen  $\text{NH}_3\text{-N}$  for synthesis of microbial protein. Therefore, it appears that utilisation of browse and other legume species with high tannin levels can be improved by inclusion of tannin-binding agents such as PEG and an additional energy source to trap the N resulting from the fermentation.

### *Efficiency of microbial protein synthesis*

Incubation of tannin-containing feeds alone resulted in a higher efficiency of microbial protein synthesis as compared to the values measured in presence of PEG (24 h incubation). Norton & Ahn (1997) evaluated the effect of PEG infusion to sheep fed on *Calliandra calothyrsus* and reported higher microbial efficiency in the non-PEG group compared to those infused with PEG. Addition of PEG to starch-containing feeds decreased the molar proportion of propionate and hence the efficiency of microbial protein synthesis compared to that in absence of PEG. Higher propionate production in absence of PEG was also reported by Nunez-Hernandez *et al.* (1991). The amount of  $\text{CH}_4$  produced was not measured in the current experiment, but *in vitro* studies indicated that the amount of  $\text{CH}_4$  production per unit of degraded substrate was lower as the level of tannin-containing feed in the mixture was increased (Hayler *et al.* 1998). The use of an anti-methanogenic compound in the diet also increased the propionate:acetate ratio and improved the efficiency of feed utilisation (McCraab *et al.* 1997). Lower  $\text{CH}_4$  production and a higher proportion of propionate are consistent with a higher efficiency of microbial protein synthesis in presence of tannins. In the presence of PEG, not only the amount, but also the composition of microbes was affected (Getachew *et al.* 1998d). Other explanations for the higher efficiency of microbial yield in the absence of PEG could be that tannins may act as an antilytic agent in the rumen and also lead to better synchronisation of nutrients released during fermentation in presence of tannins. Recently, it has been postulated that tannins at a lower concentration reduce bacteriophages (bacterial viruses) which cause the reduction in the efficiency of

feed utilisation in the rumen through the non-specific lysis of bacteria (Klieve *et al.* 1996). Tannins have also been shown to have anti-protozoal activity (Makkar *et al.* 1995a) which will also increase the efficiency of microbial protein synthesis in the rumen.

The net microbial mass production would depend on the balance between decreased degradable DM and higher microbial mass production per unit DM digested. The beneficial effect of tannins when forages containing low level of tannins are fed (Barry *et al.* 1986; Waghorn *et al.* 1994) could be due to the protection of protein from microbial degradation by tannins; this increases the amount of by-pass protein to the lower gut as well as causing a higher flow of microbial proteins to the intestine as a result of higher efficiency of microbial protein synthesis. To our knowledge, this is the first report which demonstrates unequivocally a lower efficiency of microbial protein synthesis on incubation of tannin-containing feeds with PEG or conversely, higher efficiency of microbial protein synthesis in presence of tannins.

The beneficial effects of tannins include protection of dietary proteins in the rumen and the prevention of bloat. Montossi *et al.* (1997) reported that addition of PEG to tannin-containing diets reduced live-weight gain. Similar results were reported by Terrill *et al.* (1992) and Barry *et al.* (1986). Inclusion of PEG in the diet resulted in reduced non- $\text{NH}_3\text{-N}$  digested post-rumen, increased N excreted in urine, and increased rumen  $\text{NH}_3\text{-N}$  (Barry *et al.* 1986). These authors also reported an increase in apparent digestibility in the PEG-treated group. Degen *et al.* (1998) used PEG with *Acacia saligna*, diets containing Quebracho tannin, and tannic acid, and found that addition of PEG increased DM intake and body-weight gain not only for *Acacia* and Quebracho (which contain mainly condensed tannins) diets but also for tannic acid diets (hydrolysable tannin). This suggests that the effect of PEG *in vivo* is not only restricted to condensed tannins but also to the hydrolysable tannins in feeds.

The extent of positive or negative effects of tannins may vary depending on the type and level of tannins in feeds and their biological activity, level of tannin intake, and quality of basal diets. Addition of PEG is advantageous if the tannin content of the feed is high to the extent that it depresses microbial activity and digestibility of feeds drastically. On the other hand, addition of PEG to low-tannin feeds may result in negative effects by reducing the amount of by-pass protein and also by decreasing the efficiency of microbial protein synthesis.

### **Conclusion**

Browse and herbaceous legumes are rich in N content, and these form an integral part of livestock feed in the dry season in many parts of tropical regions. The consumption of a feedstuff containing high levels of tannins produces adverse effects on nutritive value of feed and animal performance, and therefore there is a tendency to select forages for low tannin content. Screening of germplasm against tannin content may result in disappearance of the important genes responsible for the agronomic, ecological and nutritional advantages of the plants. Addition of PEG

to tannin-containing feed increased *in vitro* gas and SCFA production, and *in vitro* degradation of N. Therefore, there appears to be a potential to improve the utilisation of tannin-containing feeds without altering the genetic pool of tannin-containing plants with the use of a tannin-binding agent such as PEG. Inclusion of energy sources with the aim of synchronising N degradability and availability of energy could increase the efficiency of microbial protein synthesis in the presence of PEG. Provision of additional energy sources in PEG- and tannin-containing feed would also increase the efficiency of utilisation of resources and ensure environmentally sound feeding systems. The results obtained in the present study indicate the possibility of improving the utilisation of tannin-containing plants, which have an agronomic advantage over the non tannin-containing plants in their adaptation to biotic and environmental stresses.

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