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Influence of formic acid and dry matter on protein degradation  
in the tanniniferous legume sainfoin

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## 23 **Abstract**

24 Improved protein utilization due to protein tannin binding has been observed in sainfoin and other  
25 tannin containing legumes. Since it is well known that binding of protein to tannins and dissociation of the  
26 resulting complexes are pH dependant, effects of low pH after treatment with unusually high amounts of  
27 acidic silage additives and different dry down stages on fresh herbage prior to ensiling was investigated.  
28 Criteria for tannin binding effects were based on N fractionation in an *in vitro* ruminal digestion system. A  
29 mixture of sainfoin varieties were wilted to different dry matter (DM) levels and treated with 0, 4 and 8 ml  
30 formic acid per kg fresh matter prior to ensiling for 60 days. Nitrogen fractionation included measurement  
31 of total N, buffer soluble N, non-protein N,  $\alpha$ -amino acid N and ammonia N. Protein degradation was  
32 measured by an *in vitro* ruminal protein digestion system. Results show beneficial effects of moderate and  
33 high acidification particularly for buffer soluble N at 180 and 400 g/kg DM ( $p < 0.0001$ ) while these effects  
34 level off at 500 and 600 g/kg DM ( $P < 0.852$ ). However, no negative effect related to impaired tannin  
35 protein binding in the silages or the *in vitro* protein degradation occurred.

36 *Keywords:* formic, pH, silage, inhibited, *in vitro*, additives

37 *Abbreviations:* AA-N, amino acid N; BSA, buffer soluble N; DM, dry matter; FA, formic acid; FD,  
38 fraction degraded protein N; FM, fresh matter; FUD, fraction undegraded protein NNPN, non protein N;  
39 RF, rumen fluid; TCA, trichloroacetic acid

## 41 **1. Introduction**

42 Forage conservation as hay or silage is accompanied by decreased nutritive value and loss of dry  
43 matter (DM) due to oxidation and fermentation of the forage. These losses occur during harvesting,  
44 ensiling and feed-out due to respiration, rain damage, mechanical losses and silage effluents. (McDonald  
45 et al., 1991). This is an economical loss to farmers and an environmental problem due to increased carbon  
46 emissions and pollution of ground water with nitrates (Woolford et al., 1983; Jones and Jones, 1995). To  
47 reduce these losses, a variety of ensiling techniques with various types of silage containments and silage

48 additives, have been implemented (Buxton et al., 2003). Wilting and application of acids are two  
49 conservation methods widely used in ensiling. Wilting decreases plant protease activity and clostridial  
50 growth, lowers the amount of silage effluent and increases the concentration of sugars in the crops thereby  
51 improving silage nutritional quality (Gross and Grub, 1968; McDonald et al., 1991; Henderson, 1993).  
52 This is particularly important when ensiling legumes, which generally have low sugar and high protein  
53 concentrations. Acidifiers, such as formic acid (FA), quickly lower pH at the beginning of ensiling.  
54 Depending on the crop ensiled and on the amount of FA used, conditions for bacterial activity and growth  
55 are altered (McDonald and Henderson, 1974; Randby, 2000). Trends to lower fermentation acid  
56 concentrations, decreased lipolysis and higher water soluble carbohydrate concentrations have been  
57 reported from FA treatment of legumes (Henderson et al., 1972; Adesogan and Salawu, 2004). Studies by  
58 Pursiainen and Tuori (2008) and Lorenz (2010) showed decreased protein degradation in acidified legume  
59 silages.

60 Drawbacks of using FA have been observed in studies where FA was applied to low DM grass  
61 silage, and include increased amounts of effluent (Henderson and McDonald, 1971) and higher DM losses  
62 due to yeast growth (Henderson et al., 1972). Very low concentrations of FA may also promote growth of  
63 clostridia (Beck, 1968). Forages which are difficult to ensile, such as cocksfoot or legumes, benefit from  
64 the addition of up to 6 ml FA per kg fresh matter (FM), as shown by Lancaster and Brunswick (1977).  
65 This study also showed that higher levels of FA addition, up to 8 ml FA/kg FM increased DM intake, daily  
66 weight gain and organic matter digestibility in lambs, although this was not considered to be economically  
67 feasible.

68 Effects of tannins in legumes on protein degradation, resulting from tannin-protein binding and  
69 precipitation, has been described in many publications (Karnezos et al., 1994; Koivisto and Lane, 2001)  
70 and were also reviewed by Mueller-Harvey (2006) and Rochfort et al. (2008). Tannin-protein binding may  
71 increase the nutritive value of the forage for ruminants by an increase in rumen escape proteins which can

72 subsequently be released at the low abomasal pH (Oh and Hoff, 1987) for absorption in the small intestine  
73 (Thomson et al., 1970; Jones and Mangan, 1977).

74 Tannin-protein complexes in silage have been exposed to a low pH before entering the rumen.  
75 Hence, if pH is too low, the positive protein sparing effect could be impaired if the complexes have  
76 dissociated. However, numerous reports have shown reduced ruminal protein degradation in tannin-rich  
77 legume silages such as sainfoin (*Onobrychis viciifolia*; Wang et al. (2007) and Lorenz et al. (2010)),  
78 birdsfoot trefoil (*Lotus corniculatus*; Albrecht and Muck (1991)) and tannin supplemented perennial  
79 ryegrass (*Lolium perenne*) silage (Salawu et al., 1999).

80 In light of these known dissociative effects on the protein-tannin complex associated with low pH,  
81 coupled with current recommendations for increased FA application to improve silage storage quality,  
82 investigations into FA application limits are appropriate. Therefore, our objectives were to assess effects  
83 of acidification levels, in combination with different DM contents, on the protein protecting effect of  
84 tannins in sainfoin during silage fermentation and digestion on the basis of N-fractionation and *in vitro*  
85 ruminal degradation of sainfoin silages.

86

## 87 **2. Material and methods**

### 88 *2.1. Plant material*

89 Thirteen sainfoin varieties, grown at the National Institute of Agricultural Botany, Cambridge,  
90 Great Britain (NIAB) were harvested in August 2008, frozen and transported to the Kungsängen Research  
91 Center (Uppsala, Sweden). The varieties were: *Onobrychis viciifolia* SCOP var. CPI63761, var. CPI63815,  
92 var. CPI63826, var. Sparceto, var. CPI63825, var. CPI63752, var. CPI63808, var. Dnepropetrovsk, var.  
93 Fizes, var. Palio, var. AR-111 and *Onobrychis antasiatica* var. Sisiani Local and var. Akhurian-107.

### 94 *2.2. Ensiling*

95 All samples were chopped (~3 cm), pooled and mixed together thoroughly to make one large  
96 forage mass. The mixture was ground in a meat grinder (Bankeryd A90B, Bankeryd, Sweden) and divided

97 into four batches which were either not dried (180 g/kg DM) or dried in a force draft oven at 35°C to a  
98 DM content of 400 g/kg, 500 g/kg or 600 g/kg. Each batch was further divided into three portions and put  
99 into separate plastic bags. Formic acid was added at levels of either 0, 4 (i. e. moderate acidification) or 8  
100 ml (i. e. high acidification) FA/kg FM by spraying acid on the chopped plant material followed by  
101 thoroughly shaking the bags. The plant material was packed in glass mini-silos (20 \* 3 (inner diameter)  
102 cm) in duplicate. After filling the silos with 80 to 90 g FM, leaving a 1 cm free headspace, the tubes were  
103 closed with a rubber stopper and a water lock. Silos were incubated in a dark 20°C room for 60 days, after  
104 which the content of each silo was divided into two batches, one of which was freeze dried and ground on  
105 a cutter mill (Brabender, Type 880803, Brabender OHG, Duisburg, Germany) to pass a 1 mm screen  
106 awaiting further analysis and the other batch was frozen for analysis of  $\alpha$ -amino acid N (AA-N) and  
107 ammonia NH<sub>3</sub>) N.

### 108 *2.3. Analytical methods*

109 Representative 5 g subsamples of each silo were collected immediately after application of FA and  
110 after opening the silos after 60 d and mixed with 10 ml of de-ionized water for pH measurement.

111 Dry matter of fresh samples was determined by drying at 105°C to constant weight in a forced  
112 draught oven and for ensiled material by freeze drying. Dry matter contents were corrected for remaining  
113 water and volatile losses by multiplying by a correction factor of 0.94. Nitrogen analyses were in duplicate  
114 on the freeze dried material by a Kjeldahl procedure using a Kjeltac Analyzer 2400 and a 2020 Digester  
115 (Foss, Hillrød, Denmark) with Cu as a catalyst. Buffer soluble N (BSN) was determined by extracting  
116 freeze dried samples with a borate phosphate buffer, pH 6.75, at 39°C for 1 h according to a modified  
117 method by Licitra et al. (1996): 1.5 g of the freeze dried plant material was weighed into a 50 ml tube  
118 (Sarstedt, Nümbrecht, Germany) and mixed with 50 ml of borate phosphate buffer. Tubes were sealed,  
119 shaken and incubated for 1 h in a 39°C water bath with an additional thorough shaking every 15 min.  
120 Thereafter, tubes were centrifuged at 3000 x g for 10 min at 20°C on a swing-out rotor centrifuge (G4.11,  
121 Jouan, Saint Herbain, France). Twenty ml of the supernatant was transferred to a Kjeldahl tube and

122 analyzed for BSN. To avoid floating particles, a polyester cloth with 20- $\mu$ m openings was wrapped around  
123 the tip of the pipette and the liquid was pipetted slowly. A stepwise increase of the temperature of the  
124 Kjeltex digestion block was needed when analyzing the aqueous samples to prevent extensive foaming  
125 during digestion. Initially, temperature was slowly increased to evaporate excess water. In the last step, Cu  
126 containing  $K_2SO_4$  tablets were added and the standard procedure described above for solid samples was  
127 completed. For analysis of non-protein N (NPN), another 15 ml of the BSN extract was transferred to a 30  
128 ml polypropylene centrifuge tube and 1.5 ml of trichloroacetic acid (TCA; 200 g/L) was added and  
129 incubated for 1 h in ice water to precipitate polypeptides and proteins. After incubation, the sample was  
130 centrifuged at 27000 x g for 15 min at 20°C in a Suprafuge 22 centrifuge with fixed-angle rotor (Heraeus  
131 Sepatech GmbH, Osterode, Germany) and the aqueous supernatant was analyzed for NPN according to the  
132 procedure for aqueous samples described above.

133 Aqueous extracts from fresh-frozen samples were analyzed for  $NH_3$ -N and AA-N using phenol-  
134 hypochlorite and ninhydrin (Broderick and Kang, 1980), respectively, on a Technicon Auto Analyser  
135 (Dublin, Ireland). Leucine was used as a standard for amino acids and ammonium sulphate (Merck,  
136 Darmstadt, Germany) for  $NH_3$ -N. Both were expressed as a proportion of BSN.

137 Tannins were measured by a HCl/butanol method: Triplicates of 50 mg freeze dried sample were  
138 weighed into 30 ml polypropylene test tube. Five ml of HCl/butanol reagent (HCl:butanol=1:20) were  
139 added to the samples and they were incubated in a water bath set at 100°C for 60 min. Tubes were cooled  
140 to 20°C and absorbance read on a spectrophotometer (Pharmacia LKB, Uppsala, Sweden) at 550 nm.  
141 Tannin values were reported on the basis of absorbance units at 550 nm per 50 mg DM ( $AU_{(550)}$ ).

#### 142 *2.4. Inhibited in vitro (IIV) protein degradation*

143 Protein degradation was measured by a modified IIV method from Broderick (1987). An  
144 equivalent of 4 mg N of freeze dried, not ensiled material or freeze dried silage was weighed in a 30 ml  
145 polypropylene centrifuge tube in duplicate. The material was wetted for 60 min at 20°C in 10 ml  
146 McDougal's buffer. Rumen fluid was collected from two fistulated cows in the morning after feeding and

147 strained through four layers of cheese cloth. Cows were fed daily 3.4 kg DM grass hay with a crude  
148 protein content (CP) of 155 g/kg DM and 1.6 kg concentrates (CP: 200 g/kg DM). Strained rumen fluid  
149 was pre-incubated with 8 g/L maltose, 4 g/L soluble starch, 4 g/L sucrose, 4 g/L xylose, 4 g/L pectin and 5  
150 g/L NaHCO<sub>3</sub> in a 39°C water bath for 3 h during constant stirring. After pre-incubation, pH was adjusted  
151 to 6.7 with 3 N NaOH. Bacterial growth was inhibited by adding 41.67 ml hydrazine sulphate solution (72  
152 mM/L), 41.67 ml chloramphenicol solution (10 mM/L) and 0.234 g mercaptoethanol per liter pre-  
153 incubated rumen fluid. After additional 30 min, the sample fermentation was initiated by adding 20 ml of  
154 rumen fluid to the sample tubes. Final concentrations of hydrazine sulphate and chloramphenicol were 30  
155 and 1.3 µM/L in each incubation vessel. Shaking and a constant temperature of 39°C were maintained  
156 under anaerobic conditions. Samples of 1 ml for NH<sub>3</sub>-N and AA-N were collected every hour for 4 h and  
157 mixed with 0.1 ml TCA solution (550 g/L) to stop fermentation.

158 The fraction of undegraded protein N at the respective time point  $i$  ( $FUD_{(i)}$ ) was calculated as:

159 
$$FUD_{(i)} = 1 - FD_{(i)}$$

160 
$$FD_{(i)} = (\text{NH}_3\text{-N}_{(i)} + (\text{AA-N}_{(i)}/0.7003))/N_{(\text{sample})}$$

161 0.7003 is the average proportion of  $\alpha$ -amino N in total N determined for a number of protein  
162 sources, which derives from an average of 50 µmol total amino acids in 1 mg total N for most feed  
163 proteins after acid hydrolysis, and

164  $N_{(\text{sample})}$  is the amount of N in mg weighed into the incubation flask, and

165  $\text{NH}_3\text{-N}_{(i)}$  is in mg present in the sample, and

166  $\text{AA-N}_{(i)}$  in mg present in the sample corrected for the blank at time=0.

## 167 2. 5. *Biometric analysis*

168 Silage data was analyzed using Proc Mixed of SAS (SAS, 2003). The experiment had a factorial  
169 design with four DM levels (i.e. 180, 400, 500, 600 g/kg) and three FA levels (i. e. 0, 4, 8 ml FA/kg FM).  
170 Dependent variables were N, BSN, NPN, AA-N, NH<sub>3</sub>-N and pH. Analysis of variance with means  
171 separations by pairwise comparisons using Tukey's test (not shown in table form) was completed.



172 Proc Mixed in SAS was used for repeated measurements with unstructured covariance matrix used  
173 to test for effect of DM and FA in the IIV digestion. Class variables were time, DM and FA. Response  
174 variable was FUD at respective time points. According to the technique of Dhanoa (1988), the MLP  
175 program (Ross, 1987) was used to fit all curves of FUD *versus* time to the equation:

$$176 \quad \text{FUD} = a + b * \text{Exp}(c * t)$$

177 where: 'a' = undegradable protein fraction; 'b' = potentially degradable protein fraction, 'c' = rate  
178 (/h) and t = time (h).

179 One curve out of 12 was excluded due to poor fit. To test for differences in parameter values, the  
180 approximate analysis of variance provided by the MLP program (i. e. parallel curve analysis option; CPCA  
181 = 2) was applied. Four residual sums of squares were estimated being i) within individual curves, ii) within  
182 curves fitted using a common non-linear parameters ('c'), iii) within curves fitted using a common 'c' and  
183 'b'-value and iv) for the combined data set. Tests for curve displacement ('a'), common linear ('b') and  
184 common non-linear ('c') parameters among the curves used an F-test and the mean residual sums of  
185 squares were calculated according to Ross (1987).

186

### 187 **3. Results**

188 Good nutritional silage quality was anticipated since there were no visible moulds or yeast growth  
189 and NH<sub>3</sub>-N ranged from 4.67 to 12.14 g/kg N. A photometric value of 1.244 AU<sub>(550)</sub> confirmed the  
190 presence of tannins in sainfoin.

#### 191 *3.1. pH*

192 Formic acid addition resulted in pH values from 4.05 to 5.48 prior to ensiling and values from 3.67  
193 to 4.76 for the 60-d silage. Silage pH was increased by higher DM (P<0.001) and increased FA level  
194 lowered pH (P<0.001; Table 1). Lowest silage pH was at the highest acidification level and lowest DM  
195 content, whereas the highest pH was measured in the non-acidified, high DM silages.

196 The interaction between DM stage and FA addition ( $P<0.001$ ) indicates that increasing FA addition  
197 decreased the effect of DM.

### 198 3.2. *N and BSN*

199 Data on the distribution of N in different fractions are in Table 1. Total N content was lower in  
200 acidified compared to non-acidified silages ( $P<0.008$ ). The increase from moderate to high FA addition  
201 did not further decrease total N content ( $P=0.916$ ). The only reduction in total N content as an effect of  
202 DM content was seen when low DM (180 g/kg) silage was compared to high DM (600 g/kg) silage  
203 ( $P<0.045$ ).

204 The proportion of BSN to total N was influenced by both FA ( $P<0.01$ ) and DM level ( $P<0.001$ ).  
205 The response of FA addition was dependant on DM, as effects of FA tended to be lower at high DM stages  
206 (Interaction FA\*DM  $P=0.001$ ). Wilting from 180 g/kg DM to 400 g/kg DM in non-acidified silage had the  
207 highest impact on BSN (-33%) followed by FA addition to 180 g/kg DM silages (-20%). Overall,  
208 moderate and high FA addition decreased BSN ( $P<0.01$ ) in silages whereas there was no difference  
209 between moderate and high FA addition ( $P<0.224$ ). Lowest levels of BSN were in samples dried to 500  
210 and 600 g/kg DM ( $P<0.002$ ; Figure 1)

### 211 3.3. *NPN, AA-N and NH<sub>3</sub>-N*

212 The NPN proportion, expressed relative to total N was altered by DM ( $P<0.001$ ) and FA addition  
213 ( $P<0.015$ ; Table 1). Increasing FA from medium to high had no effect on NPN, AA-N and NH<sub>3</sub>-N. Wilting  
214 was effective for AA-N and NH<sub>3</sub>-N up to 500 g/kg and 400 g/kg DM, respectively.

### 215 3.4. *In vitro protein degradation*

216 Dry matter content ( $P<0.001$ ), FA addition ( $P<0.001$ ) and their interaction (FD\*DM,  $P<0.001$ )  
217 affected protein degradation expressed as FUD (Table 2).

218 The highest protein degradation was for silages with low DM and no FA addition. Overall, FUD  
219 was high for all treatments and ranged from 0.56 to 0.63 (Figure 2). A pairwise comparison of treatments  
220 show differences in FUD between silages at all DM ( $P<0.001$ ) except for silages of 400 and 500 g/kg DM.

221 Parameter estimates from fitting the degradation data are also in Table 2. Rates ('c') did not differ among  
222 treatments in spite of a wide range in rate from 0.12 to 1.2 /h and neither did the potentially degradable  
223 fractions ('b') but there was curve displacement ( $P < 0.001$ ), signifying differences in amounts of  
224 undegradable protein ('a'). The curve fitted to the combined data sets gave estimates for 'a', 'b' and 'c' of  
225 0.59, 0.21 and 0.40 /h, respectively (not shown in table).

226

## 227 **4. Discussion**

### 228 *4.1. Effect of treatments on pH*

229 Wilting and FA addition both altered pH in the silage as expected (Table 1).  
230 Addition of 4 or 8 ml FA/kg FM represents a moderate and a high level of acid application compared to  
231 other studies on forages using 2.5 to 4.5 ml FA/kg FM (Adesogan and Salawu, (2004); Lorenzo and  
232 O'Kiely, (2008); Mann and McDonald, (1976). Surprisingly, silage pH was not decreased in our study by  
233 addition of 4 ml FA/kg FM on the low-DM forage, probably as a result of compensatory inhibition of  
234 lactic acid fermentation as previously suggested (Henderson and McDonald, 1971; McDonald and  
235 Henderson, 1974). This explains the strong interaction between DM and FA level, and the influence of  
236 DM on the efficacy of FA to decrease pH level.

### 237 *4.2. Effect of treatments on N fractions*

238 A treatment effect of DM and FA addition on all N-fractions occurred, in accordance with other  
239 studies (Valentine and Brown, 1973; Charmley, 2001). Higher DM resulted in higher total N probably due  
240 to loss of carbohydrates from plant respiration. The change in total N by FA addition cannot be explained,  
241 but changing total N values have been observed in studies by Salawu (2001), Adesogan and Salawu (2004)  
242 or Lorenzo and O'Kiely (2008).

243 The most pronounced effect of FA addition and DM was on the soluble N fraction, which  
244 decreased with increasing FA addition and DM up to 500 g/kg. No effects of FA addition on BSN in silage  
245 with 500 or 600 g/kg DM occurred. The reducing effect of FA on  $\text{NH}_3\text{-N}$  has been previously described,

246 reviewed by McDonald et al. (1991), and was also observed in our study with a 50% NH<sub>3</sub>-N decrease in  
247 comparison with non-acidified silages.

248 That AA-N continued to decrease with higher DM content from 400 to 500 g/kg, while NH<sub>3</sub>-N did  
249 not decrease further, indicates that NH<sub>3</sub>-N was not a good indicator of inhibition of protein degradation, as  
250 suggested by Davis (1998). Ammonia N would overestimate protein degradation whereas AA-N seemed  
251 to be more indicative of protein degradation. A previous study (Hristov and Sandev, 1998)) showed that  
252 AA-N seems to be a good indicator of protein degradation.

### 253 4.3. *In vitro* silage protein degradation

254 Results from the IIV incubations resulted in an estimated degradation rate of 0.4/h for the  
255 combined data sets and an undegradable fraction of 0.6. This may have been the result of the method with  
256 a slowly dying microflora and a short incubation time. However, it seems clear that the ensiling treatments  
257 had little effect on protein degradation rate. Protein degradation rates in tanniniferous plants such as  
258 lespedeza or sainfoin, calculated by linear regression after <sup>c</sup>log transformation, have been found to be very  
259 low compared to non-tannin containing plants, ranging from less than 0.01 to 0.06/h in un-ensiled material  
260 (Broderick and Albrecht, 1997). As rate estimates may differ, depending on calculation method, we also  
261 calculated them according to Broderick and Albrecht (1997). This resulted in degradation rates from 0.017  
262 to 0.056/h (not shown in tables). These similarly low rates might be attributed to tannin binding and to  
263 inhibitory tannin effects on the proteolytic bacteria and their enzymes (Jones et al., 1994). However, this  
264 method estimates rate, assuming complete degradation of the protein by fermentation, something which  
265 was not evident from curve analysis.

266 Acidification and DM had an impact on FUD (i. e. DM \* FA effect) as shown earlier in  
267 degradation studies by McDonald et al. (1991). This caused curves displacement as curves shifted upwards  
268 in the low DM acidified silages relative to the non-acidified silages, which means that the protein fraction  
269 'a' (i. e. undegradable protein fraction) increased (Fig. 2 a and b). However, no change in rate of  
270 degradation or the potentially degradable fraction occurred (Table 2).

271 Addition of FA resulted in lower FUD values compared to non-acidified silages, with no signs of  
272 possible dissociation of tannin bound proteins, either during ensiling or *in vitro* degradation.

273

## 274 **5. Conclusions**

275 Effects of formic acid on N fractions were more pronounced at low than at higher dry down stages. The  
276 hypothesis that a dissociative effect of tannin protein binding and subsequent loss of dietary protein due to  
277 increased proteolysis occurs, could not be validated. A pH value as low as 3.67, which could result from  
278 use of high application levels of formic acid, is therefore unlikely to jeopardize the beneficial effects of  
279 sainfoin tannins either by increasing protein degradation in the silage or in the rumen.

280

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287

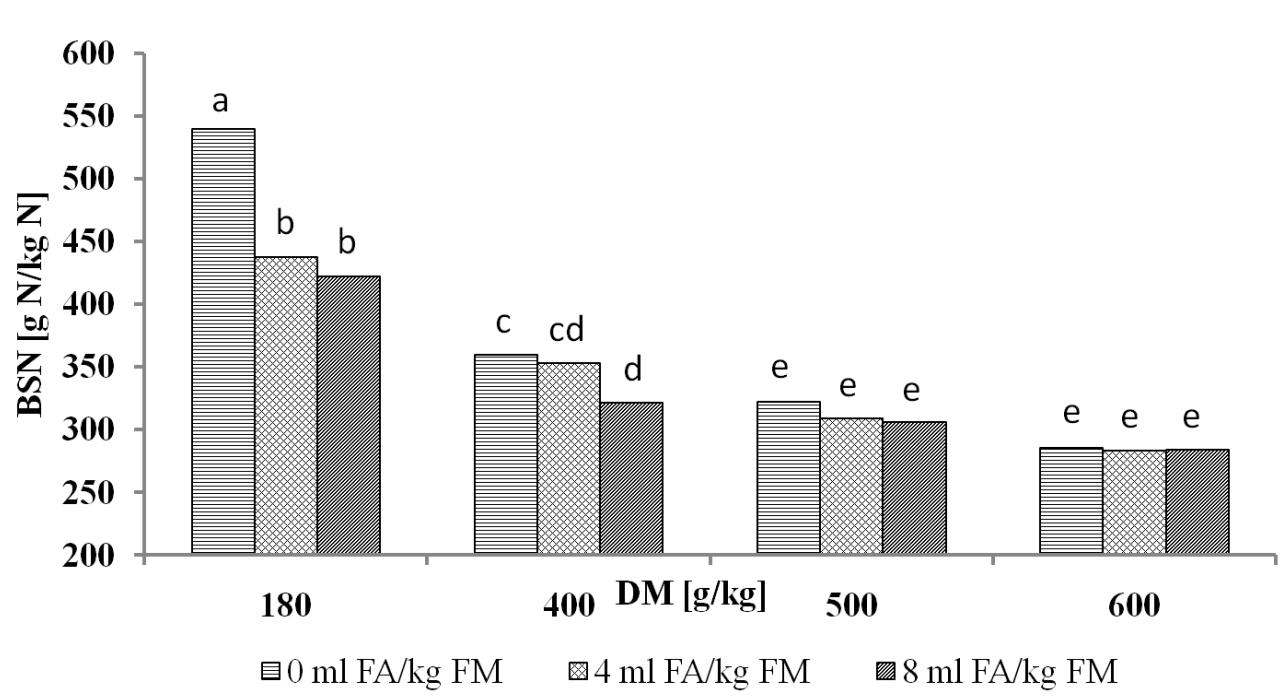
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383 Figure 1. Buffer soluble N (BSN) in acidified and wilted silages (BSN values in Table 1). Values with  
384 different superscripts differ (P<0.05).



385  
386 Figure 2. Fraction of undegraded protein (FUD) after *in vitro* degradation at: a) 180 g/kg DM; b) 400 g/kg  
387 DM; c) 500 g/kg DM; d) 600 g/kg DM (FUD values in Table 2).



