

Assessing the accuracy of current near infra-red reflectance spectroscopy analysis for fresh grass-clover mixture silages and development of new equations for this purpose

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1	Assessing the accuracy of current near infra-red reflectance spectroscopy
2	analysis for fresh grass-clover mixture silages and development of new
3	equations for this purpose
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14	

16 Abstract

17 The purpose of this study was to ascertain whether Near Infra-Red Reflectance Spectroscopy (NIRS) prediction equations calibrated on grass silage samples, could 18 19 accurately predict the chemical composition of mixed grass-clover silage samples, and 20 furthermore, to develop and calibrate new grass-clover equations should the grass-21 based equations be insufficiently accurate for these silages. A set of 94 silage samples 22 from mixed grass-clover swards (clover concentration (CC) ranging from 4 to 1000 23 g/kg as fed; determined manually) were analysed for chemical composition using 24 reference laboratory techniques, in vivo digestible organic matter in the dry matter 25 (DOMD, in sheep), and *in situ* degradability of dry matter and crude protein (in cows). 26 The same samples were scanned fresh (undried and unmilled, as is standard practice 27 for silage analysis within UK laboratories) using NIRS (at AFBI, Northern Ireland) and grass-based prediction equations applied. Predicted and observed results were 28 29 compared. Of 15 chemical components that were tested for prediction accuracy, only 30 volatile-corrected dry matter and nitrogen were well predicted (RPD values of 4.9 and 31 2.4 respectively, with low root mean square errors of prediction (RMSEP)). Neutral 32 detergent fibre and DOMD showed low RPD values, however the predicted and 33 observed datasets had no significant bias between them and were therefore also 34 considered as fit for purpose. Variables with significant bias between predicted and 35 observed datasets that were not considered suitably accurate included crude protein, acid detergent fibre, microbial dry matter yield and the effective degradability of 36 protein. For many components, bias could be attributed at least in part to CC and 37 38 changes in the fractionation of nutrients present. For some variables such as crude 39 protein, grass-based equations were sufficiently accurate at low CCs but became 40 inaccurate as CC increased, as expected. In response to inadequate prediction

41 accuracy of certain nutrients, new grass-clover equations were calibrated using the 42 obtained spectra. These were validated and results indicated that the grass-clover-43 based equations outperformed their grass-based counterparts. The adoption of new 44 grass-clover equations, or alternatively, with further development, the use of a CC 45 correction factor to the existing grass-based equations, is recommended for 46 commercial laboratories offering undried and unmilled silage analysis on samples 47 containing clover.

48

49 Keywords: Grass, Clover, silage, mixtures, NIRS, calibration,

50

51 Abbreviations: ADF, acid detergent fibre; aNDF, neutral detergent fibre; CC, clover 52 concentration; CP, crude protein; VCODM, volatile corrected oven dry matter; EDN, 53 effective degradable nitrogen; EDDM, effective degradability of dry matter; EE, ether 54 extract; FiM, Feed into Milk; OM, organic matter; LA, lactic acid; MDM, microbial dry 55 matter; N, nitrogen; NH₃-N, ammonia nitrogen; NIRS, near infrared reflectance 56 spectroscopy; NMSC, normal multiplicative scatter correction; r^2 , coefficient of determination of cross validation; RMSEP, root mean standard error of prediction; 57 58 RPD, ratio of standard deviation of the measured population to the standard error of 59 prediction; SEC, standard error of calibration; SECV, standard error of cross 60 validation; SEP, standard error of prediction; SNVD, standard normal variate de-61 trending; TMR, total mixed ration; TVC, total volatile content; TVFA, total volatile fatty acids; WMSC, weighted multiplicative scatter correction; WSC, water soluble 62 63 carbohydrate.

64

65 **1. Introduction**

66 Near Infra-Red Reflectance Spectroscopy (NIRS) is a relatively rapid and inexpensive 67 technique, routinely used to provide nutritional analysis of silage and other livestock feeds in the dairy and beef industries. However, obtaining accurate results requires 68 69 robust prediction equations. This is particularly relevant to the UK where most silages 70 are analysed 'fresh' (i.e. undried and unmilled) for rapid through-put in comparison to 71 Europe where analysis of dried, ground samples is more common. Dry analysis 72 requires lengthy sample preparation but has the benefit of increased precision of NIRS 73 prediction, partly explained by the increased homogeneity of ground samples as well 74 as the stability of the feedstuff after the removal of water (Sorensen, 2004). Currently 75 UK laboratories do not offer NIRS equations for grass-legume mixtures, instead, a 76 prediction equation with a monoculture grass-based calibration is used for a number 77 of different grass and legume-based forages.

78 This study focusses on NIRS analysis for grass-clover silages, since clover is 79 thought to be present within grass swards on 70% of UK dairy farms, and therefore is 80 likely to be the most widely-grown forage legume in the UK (DEFRA, 2015). 81 Furthermore, clover-containing forages are thought to be a promising feed to increase 82 sustainability on farms due to reduced inorganic fertilser required for growth in 83 comparison to ryegrasses (Elgersma *et al.*, 2000), while maintaining high yields of milk 84 or meat due to a fast rate of passage promoting intake (Dewhurst et al., 2009; Copani 85 et al., 2016). A preliminary study has shown that the current NIRS analysis available 86 for use on grass silages in the UK has poor prediction accuracy of crude protein, pH and lactic acid when used on mixtures containing both clover and grass (Davies et al., 87 88 2012). However, Davies et al. (2012) did not evaluate the degradability of dry matter 89 (DM), nitrogen (N), or the apparent total tract digestibility of organic matter (OM; from 90 which metabolisable energy (ME) is calculated) for prediction accuracy, despite these

91 nutrient fractions being very important for diet formulation when balancing the ratio of 92 metabolisable protein to metabolisable energy supply. Imbalances in the degradable 93 protein to fermentable energy ratio will result in poor N use efficiency. Creating 94 calibration equations for grass-clover silages poses a challenge because these silages are a mixture of two (or more) forage species, meaning that any resulting equation 95 96 must be able to deal with a broad spectrum of sample composition. To date, the majority of forage-based NIRS calibrations have focussed on predicting the nutritional 97 98 composition of just one species, and moreover, in a few instances where mixtures 99 were analysed using NIRS, typically the focus of the study was on predicting botanical 100 composition rather than chemical composition (Wachendorf et al., 1999; Cougnon et 101 al., 2014; Karayilanli et al., 2016).

The objective of this study was primarily to assess the adequacy of a grass silage-based prediction equation, commonly used in the UK for predicting chemical composition, when it was applied to grass silage samples that contained clover in varying concentrations. Subsequently, a secondary objective was to investigate whether using grass-clover based prediction equations could improve accuracy of predicted chemical composition.

108

109 **2. Material and methods**

110 2.1 Experimental design

In total, 94 grass-clover silages were sourced from commercial farms and transported to the Centre for Dairy Research (CEDAR), (Arborfield, Reading, UK) for processing. Samples were acquired from a diverse range of UK farms to ensure maximum variation within the sample set, in line with the findings of Cougnon *et al.* (2014) for sourcing robust calibration data. Silage was collected over three consecutive years

(2012/13, 2013/14, and 2014/15). The clover content range of greatest importance was deemed to be 300 - 600 g/kg DM as a more even distribution of grass-clover within a ley has been shown to create the most advantageous conditions for growth and promote symbiotic N fixation (Nyfeler *et al.*, 2011; Luescher *et al.*, 2014); although samples containing < 300 and > 600 g/kg DM clover were also included to provide sufficient range for statistical analysis and equation evaluation.

122

123 2.2 The silage sample set

124 2.2.1 Sample description. The set of 94 silage samples consisted of 58 bales and 36 125 samples from clamps which were collected from 50 different locations distributed 126 across the UK. Of the samples where the clover variety was known (n=65) 66 % were 127 red clover, 20 % were white clover and 14 % were a mixture of both. Different cuts 128 were also represented within the set with 36 first, 20 second, 16 third and 4 fourth cut 129 silages (harvest number not reported for 22 samples). The mean CC within the set 130 was 310 g/kg DM (Table 1). The sample containing the least clover contained 4 g/kg 131 DM clover and two samples contained 1000 g/kg DM clover, however all samples 132 originated from swards that were grass-clover mixtures. Twenty-three of the 94 133 samples contained < 70 g/kg DM CC and were conosidered a 'minimal' clover group 134 for which we hypothesised prediction accuracy would be similar to that of a pure grass 135 silage. The measured concentration of weed species within samples (any species 136 other than grass or clover) ranged from 0 - 380 g/kg DM with a mean of 50 g/kg DM.

2.2.2. Sample processing. Samples sourced were either unchopped bales or chopped
clamped material. If in the form of an unchopped bale, it was mixed and chopped in a
feeder wagon (Hi-Spec Mix Max, Hi-Spec Engineering, Co. Carlow, Ireland) for 45
minutes to minimize variability in chop length. Clamp silages that were already

141 chopped, were mixed in a DataRanger diet mixer which did not contain knives 142 (American Calan, Northwood, NH, USA). The DM content of the silage was estimated 143 from the loss in weight of a subsample after it has been repeatedly placed in a 144 microwave oven (Belling 384TC, 850 Watts) until a constant weight was achieved. 145 From this determination, the amount of silage (fresh weight) required for feeding an 146 individual sheep for 63 days was calculated. This amount was then weighed into 147 polythene bags with one days' feed per bag, the air was removed under vacuum, and 148 the bags were sealed and stored frozen (-20°C) until required. Frozen subsamples of 149 each silage were stored separately for future analysis of chemical and botanical 150 composition.

151

152 2.3 Nutritional analysis

153 2.3.1 NIRS analysis A 2 kg frozen subsample of each silage was sent to the Agri-Food 154 and Biosciences Institute (AFBI; Hillsborough, Northern Ireland) where the reference 155 chemical composition of the silages was determined using UKAS accredited methods 156 and NIRS spectra were obtained. Before scanning, all samples were further chopped 157 by hand to approximately 2.5 cm lengths and then thoroughly mixed. Two separate 158 packages were prepared by wrapping approximately 100 g of fresh sample in non-159 PVC cling film (Park et al., 1999). These packages were then placed in a rectangular 160 coarse transport cell and scanned through a Foss NIRSystems 6500 instrument (Foss, 161 Hillerød, Denmark). The optical values for each scan were recorded as Log 162 1/Reflectance over the range 400-2498 nm at 2 nm gaps using the ISI v3.10 (Infrasoft 163 International, Port Matilda, PA, USA) software.

164 2.3.2 Laboratory reference analyses Dry matter was determined in a forced-air oven
165 and corrected for the loss of VFAs, lactic acid (LA), alcohols and ammonia (Porter and

166 Murray, 2001) and reported as volatile-corrected oven dry matter (VCODM). Ash was 167 measured through combustion in a muffle oven at 550°C for 18 h. Lactic acid and other 168 volatile compound measurements (total volatile fatty acids (TVFA) were determined 169 using gas chromatography following extraction of representative samples in distilled 170 water (Erwin et al., 1961; Givens et al., 2009). Nitrogen (N) was measured using the 171 macro Kjeldahl method 954.01 (AOAC, 2000) and Ammonia-N (NH₃-N) was 172 determined using a calibrated ammonia ion selective electrode, which required 30 g 173 silage soaked in 150 ml of purified water for 18 h at 4°C. (McDonald et al., 1981; Orion 174 Research, 1990). Both ether extract (EE) and water soluble carbohydrate (WSC) were 175 measured on dried and ground samples: EE according to AOAC method 920.29 176 (AOAC, 1990), and WSC as described previously (Fuller, 1967). Dried and ground 177 samples were subsequently passed on to Trouw Nutrition (Ashbourne, Derbyshire) 178 who performed analyses for neutral detergent fibre (aNDF) and acid detergent fibre 179 (ADF) both inclusive of residual ash using Fibrecap equipment (Foss, Hillerod, 180 Denmark) (Robertson and Van Soest, 1981; Kitcherside et al., 2000; Mertens et al., 181 2002). A further 200 g of silage was manually separated into clover, grass and other 182 species to determine the CC of the silage. This procedure was predominantly 183 performed by the same individual to minimise human error. Resulting fractions were 184 then dried to determine species composition on a DM basis. In vivo reference methods 185 were performed at CEDAR to determine silage digestibility and degradability.

186

187 2.4. In vivo analyses.

2.4.1 In vivo Digestibility Eighteen Mule x Texel wether sheep originating from a local
breeder were used to measure *in vivo* silage digestibility using a series of 3 x 3 Latin
square design experiments so that the final digestibility values comprised the mean of

191 measurements from three different animals. Each sheep was fed a silage sample *ad* 192 *libitum* (with 10% refusals) for 16 d adaption followed by a 5 d sampling period during 193 which sheep were placed in a metabolism crate for faeces and urine collection as 194 described previously (Givens *et al.*, 1989; Bratzler, 1951). All *in vivo* procedures were 195 licensed and monitored by the UK government Home Office under the Animal 196 (Scientific Procedures) Act 1986.

197 Sheep were enrolled on the study when they reached adult weight at > 30 kg. 198 Their diet was supplemented with 20 g/d of a general purpose vitamin/mineral mixture 199 for sheep (Countrywide, Evesham, Worcestershire, UK) and the weights of feed 200 offered and refused was recorded each day during the collection period. A subsample 201 of feed was taken and analysed for DM and ash to calculate OM content. Refused 202 feed was also corrected for DM. Out of the 94 samples, 4 were excluded from in vivo 203 analysis as there was insufficient material for the 9 week feeding schedule, but were 204 still used for all other analyses. Complete collections of faeces were taken for each 205 sheep. Each days' faecal material from the 5 d collection period was refrigerated at < 4°C until bulked together on d 5, thoroughly mixed and three 200 g subsamples 206 207 obtained. These subsamples were immediately placed in a forced air oven at 60°C for 208 72 h to determine DM content. Dried samples were then bulked, ground and a further 209 subsample was placed in a muffle oven for combustion at 500°C for 16 h for 210 determination of OM content. Digestibility results have been presented as digestible 211 organic matter in total dry matter (DOMD, g/kg DM).

212 *2.4.2 In situ degradability.* Degradability values were obtained using an *in situ* method 213 with rumen cannulated Holstein-Friesian dairy cattle. These cattle were housed in a 214 dedicated metabolism unit, fed a commercial grass-maize based total mixed ration 215 (**TMR**) diet once daily and milked twice daily at 0600 h and 1600 h approximately.

216 Fresh samples of each silage were placed in porous (43 µm pore size) bags that were 217 sequentially incubated in the rumen for six time intervals (3, 6, 12, 24, 48, and 72 h) 218 using a complete exchange method as described previously (Lovett et al. 2004). 219 Replicates were obtained by repeating the procedure with three different animals. To 220 quantify '0' hour washing loss, three further bags per silage were placed in a tub of 221 cold tap water and swirled for 5 minutes. All bags were washed (Zanussi SupeLluxe, 222 Electrolux plc, Luton, UK) on a 53 min cold wash cycle, dried (at 60°C), and weighed 223 for the determination of DM degradability, then further analysed for N (as described 224 previously). The solubility (S) of DM and N was determined by adding 1 g of DM to 30 225 ml of water and stirring for 5 minutes every half hour for a period of 2 h, the insoluble 226 material was then filtered (Whatman filter paper grade 4, Sigma-Aldrich, MO, USA) 227 (Hvelplund and Weisbjerg, 2000). The filter paper and substrate was then dried and 228 weighed to determine DM solubility by difference and residual N was measured as 229 described previously.

230 The percentage of material degraded at each time-point was used to plot a 231 degradation curve as described by Ørskov and McDonald (1979). Degradability 232 fractions termed 'a', 'b' and 'c' were obtained from the intercept, asimptote and slope 233 of the curve. Fraction 'a' contained material that is apparently degraded almost 234 immediately upon ingestion and 'b' contained the remaining insoluble but degradable 235 material with 'c' being the rate of degradation of 'b'. Two different approaches were 236 used to calculate effective degradabilitiy (ED) based on the above fractions. To ensure 237 the best comparison with predicted data, the ED of nitrogen (EDNFIM) and of dry matter 238 (EDDM_{FIM}) were calculated using the 'Feed into Milk' (FiM) rationing software 239 equations (Equation 1). In this equation the outflow rate of small (k_{liq}) and large (k_f) 240 particles was standardised at 0.075 and 0.045 respectively to fairly compare against

- 241 predicted data. EDDM_{FIM} was converted to microbial dry matter (**MDM**_{FIM}, g/kg DM)
- 242 using standard equations to convert EDDM into ATP supply as described previously
- 243 (FiM consortium, 2004).
- 244
- 245 Equation 1. $ED_{FIM} = (0.9s/(0.9+k_{Iiq}))+(b_D c/(c+k_{Iiq}))+(bc/(c+k_f))$
- Where s is the soluble proportion, k_{liq} is the fractional outflow rate of the liquid pool (0.075), b_D is the degradable small particle proportion, b is the degradable large particle proportion, c is the fractional degradation rate of b, and k_f is the fractional outflow rate of the large particle pool (0.045).
- 250

A second, simpler, approach was also tested simultaneously to calculate the ED of N and DM using 0.08 as the standard outflow rate (k) of all particles (**EDN**_{0.08},

- and **EDDM**_{0.08}) (Equation 2; Ørskov and McDonald, 1979).
- 254
- 255 Equation 2. ED = a+bc/(c+k)

Where a is the rapidly degraded, b is the slowly, potentially degradable proportion, c is the fractional rate of degradation of b, and k is the fractional outflow rate of material (0.08 h^{-1}).

259 2.5 Statistical analysis

260 2.5.1 Tests of relationships and trends within the measured dataset. Statistical 261 analysis was conducted using Genstat 16th Edition (VSNI, Hemel Hempstead, UK). 262 Composition of the silages was predicted from NIRS spectra using equations 263 developed for the UK Forage Analysis Assurance (FAA) group (www.faagroup.co.uk) 264 initially using 136 grass silage calibration samples from the studies reported by Park 265 et al. (1997, 1998) which were regularly updated with new spectra over time for most 266 chemical component variables other than those requiring in vivo reference analyses. 267 The measured dataset has been presented as maximum, minimum, mean and 268 coefficient of variation (CV%) values for each measured variable. The effect of CC on 269 each of the other variables was tested by grouping samples into minimal, low, medium 270 and high groups (which are equal quartiles of the dataset; representing samples within 271 the ranges of < 70, 70 - 250, 250 - 500 and > 500 g/kg DM CC respectively) which 272 were compared using analysis of variance (ANOVA). A post hoc Tukey test was 273 performed to determine whether there were significant differences between the means 274 of the 4 groups. The means of the observed and NIRS predicted datasets were 275 compared using a student's t-test to determine significance. Crude protein (CP) was 276 not directly measured or predicted but calculated using either measured or predicted 277 N and VCODM values (6.25 x Total N on a DM basis). For all dry matter values 278 throughout this study, VCODM has been used rather than DM, in accordance with the 279 industry standard used by UK laboratories. For ash, EE, WSC, ADF, and aNDF 280 (variables where the measured concentration is produced from a dry sample) 281 equations were produced that predicted concentrations on both a fresh basis and 282 directly on a DM basis.

283 2.5.2 Tests of prediction accuracy during validation. For the grass-based prediction 284 equation results, the difference between laboratory assays and NIRS predicted values 285 was calculated using measured minus predicted values and is henceforth termed 286 'bias'. Relative root mean square standard error of prediction (RMSEP as a 287 percentage of the measured mean), ratio of the standard error of prediction to the 288 standard deviation of the measured dataset (RPD) as recommended by Williams 289 (2014), and the R-squared value of the relationship between observed and predicted 290 data (r^2) were used to measure prediction accuracy.

291 2.5.3 Calibration of new NIRS equations. To create new grass-clover prediction 292 equations, different data pre-treatment methods were first assessed by varying use of 293 derivitives, gap, smoothing and scatter correction. All calibrations were performed 294 using the WinISI III v1.50 (Infrasoft International, Port Matilda, PA, USA) software.

295 They were carried out as Modified Partial Least Squares regressions over the range 296 1100-2498 nm using a 2 nm gap. To account for any sub-sampling error the root mean 297 square difference of each sub-sample was calculated using the WinISI III v1.50 298 software. An upper limit of 5000 was used to judge poor replication meaning any 299 sample with a root mean square greater than 5000 would be removed. None of the 300 samples in the calibration set were above this limit. Raw data and two derivatives 301 were tested in the process (Raw (0,0,1,1), 1st Derivative (1,4,4,1) and 2nd Derivative 302 (2,10,5,1)) and three scatter corrections (Standard Normal Variate Detrending 303 (SNVD), Normal Multiplicative Scatter Correction (NMSC) and Weighted Multiplicative 304 Scatter Correction (WMSC)) for each of the derivatives. The maximum number of 305 terms set for each equation was 11. There were three elimination passes carried out 306 and the cross validation value was set at 6 in which the calibration set was divided into 307 six groups with one group removed sequentially and predicted using a calibration 308 formed using the remaining samples. The validation errors were combined to give a 309 standard error of cross validation (SECV). The optimal equations were those with the lowest SECV. The combination of data pre-treatment giving the optimal prediction 310 311 model is shown in supplementary table 1 for each variable. The optimal equation was 312 compared against the industry standard method, based on the study of Park et al. 313 (1997), which was taking the first derivative (1,4,4,1) with SNVD scatter correction and 314 a repeatability file (a file containing multiple spectra from the same sample measured 315 under different conditions, designed to reduce the variability caused by differing 316 environmental conditions and instruments). Differences between the optimal 317 equations and the industry standard equations were small, therefore further validation 318 was performed using the industry standard equations as these were the most likely to 319 be utilised commercially. For the purposes of a validation test, 10 samples were

320 removed from the dataset and tested using the remaining equation. These samples
321 were chosen by including the very first sample to be collected and then every tenth
322 sample in order of their arrival at CEDAR for processing.

323

324 3. Results

325 3.1 Sample chemical composition

The silages contained a wide range of chemical composition with LA, WSC and TVFA being the nutritional characteristics with the greatest variance of those measured. Volatile corrected dry matter of the silages was evenly distributed with a mean of 395 g/kg. Measured CP concentration (calculated from N and VCODM) ranged from 57 to 215 g/kg DM and with a mean of 138 g/kg DM.

331 With the exception of ash, aNDF, and WSC, the concentration of all other 332 measured variables were affected by the CC of the sample when grouped into 333 minimal, low, medium, and high clover groups (Table 2). VCODM and N were 334 significantly increased in the high clover group (>500 g/kg DM CC) relative to the other 335 three groups (both P < 0.001), as was CP with the exception of the medium group 336 which contained an intermediary CP concentration (P < 0.001). Degradability 337 parameters calculated using the Ørskov and McDonald (1979) model and DOMD were 338 lowest in the high clover group (all P < 0.04) and numerically highest in the low clover 339 group (60-240 g/kg DM CC), however, when degradability parameters were calculated 340 using FiM equations, differences between clover groups were non-significant. 341 Fermentation end products (LA, TFVA and TVC) decreased in concentration 342 sequentially as CC increased (all P < 0.003) while pH was similar for minimal, low and 343 medium groups and higher for the high clover group (P < 0.001). NH₃-N was also

highest in the high clover group in comparison to the minimal clover group while the other two groups contained intermediate concentrations of NH₃-N (P < 0.02).

346

347 3.2 Validation of current grass-based NIRS equations

348 Using data from the present study grass-clover sample set to verify the accuracy of 349 the current grass-based prediction equations, a wide range of prediction accuracy was 350 observed depending on the chemical component tested (Table 3). Volatile corrected 351 dry matter and N showed good prediction accuracy with RPD values of 4.92 and 2.35 352 respectively, and no significant difference between observed and predicted means. 353 Furthermore, the relationship between the observed and predicted data for both these 354 variables closely followed a line of parity (Figure 1) especially at low concentrations. 355 However, all other variables led to RPD values that were <2 denoting inadequate 356 performance. Digestible organic matter in total dry matter, and aNDF, had low relative 357 RMSEP (both <10% of the observed mean) and no significant difference between the 358 observed and predicted means which could be considered acceptable despite having 359 an RPD value <2. For these variables the slope of the relationship between observed 360 and predicted data followed a line of parity however there was greater variability in the 361 relationship than was seen for VCODM and N (Figure 1). Crude protein prediction 362 showed a relatively high RPD value (1.58) and good correlation between predicted 363 and observed data ($r^2 = 0.75$) however the slope of the relationship did not follow a line 364 of parity (Figure 1) leading to a significant bias (P < 0.005) for under-estimation at 365 higher concentrations with the average under-estimation being 12.4 g/kg DM.

Fermentation characteristics (LA, pH, TVC and TVFA) all showed intermediate
prediction accuracy with RPD values ranging from 1.15 to 1.22. Of these variables, LA
in particular had a very high relative RMSEP at 71% of the observed mean as a result

369 of high variability in prediction accuracy where concentration was low (Figure 1). For 370 both TVC and TVFA there was a significant bias towards over-estimation (both P <371 0.001). Poor prediction accuracy (RPD value <1) was observed for NH₃-N, ADF, EE, 372 EDN_{FIM}, and MDM_{FIM} all of which showed a significant bias between the predicted and 373 observed means (all P < 0.001). Of special note, EDNFIM and MDMFIM showed the least 374 prediction accuracy of all the variables tested with a significant over-estimation for 375 EDN_{FIM} of 139 g/kg N and an under-estimation for MDM_{FIM} of 17 g/kg DM. Moreover, 376 predicted and observed data showed little correlation (Figure 1) indicated by r^2 values 377 of 0.01.

378 The degree of variation and the magnitude of bias in relation to sample CC is 379 illustrated in Figure 2 using CP and EDNFIM as examples which are crucial to diet 380 formulation. In the case of CP, prediction bias in samples containing 800-1000 g/kg 381 DM CC is greater than 30 g/kg DM (Figure 2a), and similarly for EDNFIM, a prediction 382 bias greater than 200 g/kg N was observed in this very high CC range. Meanwhile, 383 bias was comparatively lower in the minimal clover group (< 70 g/kg DM CC) at 6 g/kg 384 DM for CP and 103 g/kg N for EDN_{FIM} reflecting the degree of bias that might be 385 expected for a pure grass sample.

386

387 3.3 Validation of new grass-clover equations

Following production of new equations using the NIRS spectra from the grass-clover silages in the sample set, a cross validation test indicated 12 out of 21 new equations had a relative SECV of 10% of the observed mean, suggesting a good calibration was achieved for these variables (Table 4). VCODM, pH, aNDF, ADF, and EDDM_{0.08} were amongst the strongest calibrations according to cross validation while TVC, WSC, TVFA, Alcohol and LA were the least robust. For variables where both a fresh and a

394 DM basis equation were produced, the equation that predicted on a fresh basis gave 395 the more accurate result for ash, EE and WSC, whereas the opposite was true for 396 ADF and aNDF, where the equation that predicted concentration on a DM basis was 397 more accurate.

398 A validation test was also applied to the new grass-clover prediction equations 399 through removal of 10 samples from the calibration data-set (Table 5). Seven variables 400 gave an RPD value > 2 denoting good accuracy including VCODM, ADF, aNDF, EDN 401 and N. Additionally the RPD score of all values were improved relative to prediction 402 accuracy using the grass-based equations, which was reflected in greatly reduced 403 bias, for example, new equations reduced crude protein mean bias from -12.4 to -0.82 404 g/kg DM and EDN mean bias improved from 139 to 12 g/kg N on average. The new 405 alcohol and EE (DM basis) equations gave a low RPD value (>1) suggesting these 406 equations are unlikely to be suitable for use without further improvement.

407

408 **4. Discussion**

409 4.1 Chemical composition and clover concentration

410 The wide range of samples collected in this study provided a robust test for the current 411 grass-based prediction equations. The sample set was dominated by samples 412 containing predominantly grass with only a guarter of the samples obtained containing 413 > 500 g/kg DM CC. Roughly half the total number of samples obtained were below the 414 minimum optimum clover inclusion rate of 300 g/kg DM suggested by Nyfeler et al. 415 (2011). This may be due to the sample set comprising a greater number of first cut 416 silage samples than second, third or fourth cuts in which CC would have been greater 417 due to warmer and drier conditions in the latter half of the year (Chmelikova et al., 418 2015).

Crude protein concentration (ranging from 57 to 215 g/kg DM with a mean concentration of 138g/kg DM) indicated that, although some of the samples contained very high levels of crude protein, mean concentration was similar to that expected for well fertilised modern grass silages which have been observed ranging from 120-270 g/kg DM (Burns *et al.*, 2015). This mean is also significantly lower than those reported in published feed composition tables for crude protein concentration of grass-clover silages e.g. 173 g/kg DM; AFRC (1993).

426 The supply of effective degradable N (EDN) is another important factor in diet 427 formulation. High concentrations (>700 g/kg N) of rapidly degraded protein in the 428 rumen can be wasteful as there is insufficient time for bacterial N capture, which is 429 often a characteristic of legume silages (Coblentz and Grabber, 2013; Dewhurst, 430 2013). In this study, average EDN_{FIM} was 623 g/kg N so within the optimal range and 431 lower than values cited in other studies, for example, in another study, measured 432 average EDN of grass-clover silages was 880 g/kg N at an assumed passage rate of 433 0.05/hr (Hvelplund and Weisbjerg, 2000). The discrepancy may be due to clover 434 varieties in this sample set being predominantly comprised of red clovers containing 435 the enzyme poly-phenol oxidase which is thought to reduce proteolysis in the rumen 436 (Lee *et al.*, 2009). Digestibility, EDDM_{0.08}, and EDN_{0.08} all showed a similar pattern 437 where the low group (70 – 250 g/kg DM CC) gave the highest value and the high group 438 (> 500 g/kg DM CC) the lowest suggesting inclusion of clover between 70 - 250 g/kg 439 DM is an optimal range for digestibility and degradability. Poor digestibility in the high 440 group may relate to an increased maturity of clover and grass with higher lignification 441 in samples in that range (Nousiainen et al., 2009). Increasing ratios of ADF:aNDF in 442 samples with a higher CC indicates the differing fractions of fibre present in legumes 443 in comparison to grasses, especially red clover which is largely comprised of stem

444 where ADF concentration is higher than in leaves (Alstrup et al., 2016). There was a 445 notable decrease in volatiles content (LA, TVC and TVFA) and an increase in pH in 446 the high group relative to the other quartiles where values were generally similar. This 447 suggests samples with a very high CC were more difficult to ensile, perhaps due to 448 reduced availability of sugar to fuel bacterial activity that may also indicate an 449 increased maturity. Using a factor of 0.0157 of DOMD, mean ME within the sample 450 set was predicted as 9.9 MJ/kg DM which is considerably lower than recently 451 measured values for modern monoculture grass silages which ranged from 12-13 452 MJ/kg DM (Burns *et al.*, 2015).

453

454 4.2 Using grass-based NIRS equations for clover containing samples

455 The key objective of this project was to determine whether the current grass NIRS 456 equations could be applied to grass-clover samples and predict the concentrations of 457 chemical components with good accuracy. Variates that were considered most 458 important for correct diet formation included CP, EDN, MDM and DOMD as these are 459 the variables involved in balancing rumen degradable protein and energy supply. 460 Whilst the prediction accuracy of DOMD and some other variables (including VCODM, 461 N and aNDF) could be considered suitably accurate with relative RMSEPs of <10 % 462 of the measured mean, CP, MDM, and EDN were amongst the variables with high 463 relative RMSEPs and RPD values of less than 2.0 combined with a significant bias. 464 Similar results were seen in a smaller preliminary study of 58 grass-clover silages in 465 which the same equations were tested and crude protein was significantly under-466 estimated by 22 g/kg DM on average (Davies et al., 2012). The consequence of this 467 bias would be an imbalance in microbial N supply in the ration which is likely to lead 468 to reduced N use efficiency in cattle resulting in higher levels of N excretion in urine

469 and faeces contributing to environmental loading (Kebreab et al., 2002). Under-470 estimation of CP in silage samples could result in farmers under-valuing grass-clover 471 silages as protein sources, and compensating through an oversupply of expensive 472 bought-in protein within the concentrate portion of the diet. For CP and EDN, 473 increasing bias correlated with increasing CC. This might be explained by samples 474 containing a high concentration of grass being more similar in composition to the 475 calibration samples used to create the current grass-based equations. Also, bias may 476 be created due to different N fractionation within clover, with some fractions present 477 that are absent (or present in different concentrations) in grass, such as the 478 concentration of non-protein N (Chrenkova et al., 2014).

479 When considerring the impact of the observed inaccuracies on diet formulation 480 it is estimated metabolisable protein, and not crude protein that is often the protein 481 fraction used for diet formulation. Crude protein multiplied by EDN (as a proportion of 482 total N) is used to calculate effective rumen degradable protein (**eRDP**) which is one 483 of the factors that determines metabolisable protein (alongside digestable undegraded 484 true protein, **DUP**) in diet formulation software. The effect of CC on calculated eRDP 485 bias is shown in Figure 2c. The opposing bias in EDN and CP cancel out to some 486 extent at low CC however the overall effect is an over-estimation of eRDP that 487 increases at higher CC. This may lead to an oversupply of fermentable energy in 488 relation to available protein for microbial N capture, creating an imbalance that could 489 reduce the efficiency of dietary nutrient utilisation. This would only be further compounded by the inaccuracy seen in MDM prediction which is used to determine 490 491 the requirement for fermentable energy.

492

493 4.3 Performance of new grass-clover-based equations

494 Comparing the performance of new grass-clover equations with the grass-based 495 equations for use on clover-containing samples, and using relative SECV as a 496 measure of potential performance of the calibration, some of the new grass-clover 497 equations produced in this study are likely to perform well (including important variates 498 such as VCODM, N, EDN, and DOMD) whereas others have very high errors 499 (particularly the volatile compounds) and would require further development. The 500 accuracy of prediction for volatile compounds (LA, TVC, and TVFA) is notable in all 501 equations tested (both grass and grass-clover based) for producing poor reliability, 502 and volatile concentrations showed some of the greatest variation within the measured 503 sample set. Some of the lack of reliability in these equations could be due to the 504 variability of scanning undried and unmilled material rather than presenting the sample 505 in an homogenous, dried form (Sorensen, 2004).

506 For variates where the measured value was calculated on a dry sample (ash, 507 EE, aNDF, ADF and WSC) equations have been calibrated to give both a fresh and a 508 DM basis value. In most instances, the calibration for the fresh value was more robust, however, because presenting information on a DM basis is widely practiced, fresh 509 510 values would be transformed based on VCODM values which would introduce further 511 error. Overall however, in a small validation test, new equations were better able to 512 predict all variables when compared to the accuracy of grass-based prediction 513 equations. The prediction of EDN_{0.08} and EDDM_{0.08} showed marked improvement over 514 the previous prediction accuracy for EDN_{FIM} and MDM_{FIM} using the grass-based equations perhaps due to the reduced complexitiy of calculating these variables from 515 516 measured data. For example, calculating reference values for MDM_{FIM} from measured 517 degradability at different timepoints in vivo is a multi-step process involving many 518 different variables (such as corrections for solubility, fatty acid content and crude

519 protein concentration) and therefore it may not be feasible to predict such a value 520 based on NIRS spectra alone. These improvements would have a significant impact 521 on the accuracy of rumen degradable protein and fermentable energy prediction.

522

523 4.4 Implementation of new equations

524 The implementation of new grass-clover equations requires that nutritionists, feed 525 company representatives, and farmers are widely aware of the new option, and that 526 samples are correctly identified as containing a viable quantity of clover. Additionally, 527 grass-clover mixtures are just one example of alternative forages that are currently 528 gaining popularity, and it is unlikely that new equations can be created and 529 implemented for all of them due to the time needed to collect a sufficiently large group 530 of calibration samples. Therefore, it would be more convenient if one equation (such 531 as the current grass-based equation or alternatively a seperate general 'legume' 532 equation) could be adapted to analyse many different grass and legume based 533 forages. Another solution would be to use a two step process in which the CC of the 534 sample is predicted using NIRS, and then used to apply a correction to nutritional 535 predictions. A number of previous studies have used NIRS to determine the botanical 536 composition of a mixed sample (containing two species) with success for both grass-537 clover (Cougnon et al., 2014) and lucerne-grass silages (Karayilanli et al., 2016) 538 however in all instances the calibration was performed on dry samples and therefore 539 further work is required to create an analysis for fresh samples that would be 540 appropriate for use in the UK.

541

542 **5. Conclusions**

543 For some variables, notably VCODM, N, DOMD and aNDF, current UK grass-based 544 equations were able to be applied to clover-containing samples with adequate 545 accuracy. However, in general, it was concluded from the evidence observed in this 546 study that the NIRS calibration equations developed for use on grass silages, could 547 not predict a number of key chemical components (including CP and EDN) with 548 sufficient accuracy, when used for grass-clover mixture silages. This was consistent 549 with the findings of a previous study (Davies *et al.*, 2012). Therefore, we suggest two 550 possible solutions that would be appropriate for uptake by UK laboratories: (i) the 551 introduction of new grass-clover prediction equations calibrated using the sample set 552 obtained for this study or (ii) the use of a correction factor that could be applied based 553 on the CC of the sample. Furthermore, in a wider sense, this study provides some 554 evidence that caution should be used whenever NIRS equations are applied to forage 555 mixtures where only one component of the mixture was represented within the equation calibration set. Where possible, using an equation based on a specific 556 557 calibration set that is very similar to the material requiring analysis is likely to produce 558 the most accurate predictions.

559

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569

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- 714 J. Agric. Sci. 92, 499-503. http://dx.doi.org/10.1017/S0021859600063048
- 715
- 716

- 717 **Table 1** The means, ranges and variation coefficients (CV) of chemical components
- 718 measured in a set of 94 diverse grass-clover silages from UK farms (in g/kg DM
- 719 unless otherwise stated).

Item	Min	Max	Mean	CV, %
ADF	229	513	335	10.2
Ash	58	158	97	20.6
aNDF	299	585	447	10.0
CC	4	1000	310	91.3
СР	57	215	138	24.7
Degradability				
EDDM _{0.08} †	217	626	472	16.4
EDN _{0.08} , g/kg N†	55	821	625	18.0
MDM _{FIM} ‡	60	274	146	34.8
EDN _{FIM} , g/kg N ‡	297	811	623	14.3
DOMD	400	766	632	10.6
EE	14.6	42.9	26.6	26.0
LA, g/kg	0.0	64.4	13.4	91.5
рН	3.6	6.7	4.6	13.5
N, g/kg	3.6	17.7	8.8	42.2
NH ₃ -N, g/kg DM*100	17.5	203	62.5	42.2
TVC, g/kg§	2.3	76.1	23.6	57.8
TVFA, g/kg¶	1.1	74.3	19.7	66.0
VCODM, g/kg	182	793	395	33.4
WSC	3.9	164	41.4	86.3

720 CC = clover concentration; EDDM = effective degradability of dry matter; EDN = effective degradable

721 nitrogen; DOMD = digestible organic matter in total dry matter; EE = ether extract; LA = lactic acid; MDM = microbial dry matter yield; NH₃-N = ammonia nitrogen; TVC = total volatile content, TVFA =

722 723 total volatile fatty acids; WSC = water soluble carbohydrates.

724 † Degradability parameters determined by in situ incubation in the rumen, using the model of Ørskov 725 and McDonald (1979) ED = a+b[c/(c+k)] where a = rapidly soluble material; b = non-soluble but 726 degradable material; c = rate f degradation of b; and k = an assumed outflow rate of 0.08/hr. 727 **‡** Degradability parameters determined by *in situ* incubation in the rumen, using the Feed Into Milk

728 (FIM) Consortium (2004) model $ED_{FIM} = (0.9s/(0.9+k_{liq}))+(b_D c/(c+k_{liq}))+(bc/(c+k_f))$ where s = soluble

729 proportion, k_{lig} = fractional outflow rate of the liquid pool (0.075/hr), b_{D} = degradable small particle 730 proportion, b = degradable large particle proportion, c = fractional degradation rate of b, and k_f is the731 fractional outflow rate of the large particle pool (0.045/hr).

732 § TVC is the sum of acetic, butyric, lactic, propionic and valeric acids plus ethanol and propanol.

733 ¶ TVFA is calculated as for TVC but excluding ethanol and propanol.

735 **Table 2** Differences in chemical components in 94 grass-clover silages grouped into

four quartiles (Minimal (Mi), Low (L), Medium (M) and High (H)) according to their

737 clover concentration (mean of each quartile, in g/kg DM unless otherwise stated).

	Clover concentration quartiles†						
Item	Mi	L	М	Н	SED	P value ‡	
CC	34 ^a	145 ^b	335°	743 ^d	28.9	0.001	
n	23	24	24	23			
Chemical components							
ADF	311 ^a	329 ^{ab}	345 ^b	356 ^b	12.5	0.003	
Ash	91.2	94.9	103.4	96.8	5.84	0.201	
aNDF	465	452	443	432	16.5	0.229	
СР	122 ^a	130 ^a	143 ^{ab}	158 ^b	9.3	0.001	
Degradability							
EDDM _{0.08} §	470 ^{ab}	501 ^b	478 ^{ab}	436 ^a	21.7	0.032	
EDN _{0.08} , g/kg N§	643 ^b	682 ^b	640 ^b	531 ^a	28.6	0.001	
MDM _{FIM} ¶	130	135	127	122	6.3	0.218	
EDN _{FIM} , g/kg N¶	627	645	629	589	25.9	0.182	
DOMD	647 ^b	668 ^b	631 ^b	581 ^a	18.5	0.001	
EE	26.7 ^{ab}	28.7 ^b	27.4 ^{ab}	23.2 ^a	1.99	0.044	
LA, g/kg	17.6 ^b	16.2 ^b	14.0 ^{ab}	5.6 ^a	3.42	0.003	
рН	4.45 ^a	4.41 ^a	4.44 ^a	5.23 ^b	0.155	0.001	
N, g/kg	7.8 ^a	7.3 ^a	7.9 ^a	12.3 ^b	0.93	0.001	
NH ₃ -N, g/kg DM*100	48.3ª	56.3 ^{ab}	68.0 ^{ab}	77.5 ^b	9.71	0.018	
TVC, g/kg I	28.4 ^b	26.8 ^b	24.0 ^{ab}	14.3ª	3.74	0.001	
TVFA, g/kg ¥	23.3 ^b	22.4 ^b	20.8 ^{ab}	11.6ª	3.65	0.001	
VCODM, g/kg	397 ^a	350ª	347 ^a	498 ^b	35.1	0.001	
WSC	56.6	39.6	32.9	38.6	10.37	0.125	

738 CC = clover concentration; EDDM = effective degradability of dry matter; EDN = effective degradable
 739 nitrogen; DOMD = digestible organic matter in total dry matter; EE = ether extract; LA = lactic acid;

MDM = microbial dry matter yield; NH₃-N = ammonia nitrogen; SED = standard error of the difference
 between means; TVC = total volatile content, TVFA = total volatile fatty acids; WSC = water soluble
 carbohydrates.

743 † The 94 samples were sorted by ascending clover concentration and divided into four evenly sized

744 guartiles: 0-6%DM clover (Mi); 6-24% clover (L); 25-49% clover (M); and 50-100% clover (H).

The probability of there being no significant difference between treatment means determined using
 Analysis of Variance (ANOVA).

For and McDonald (1979) ED = a+b[c/(c+k)] where a = rapidly soluble material; b = non-soluble but

- degradable material; c = rate f degradation of b; and k = an assumed outflow rate of 0.08/hr.
- 750 ¶ Degradability parameters determined by *in situ* incubation in the rumen, using the Feed Into Milk
- (FIM) Consortium (2004) model $ED_{FIM} = (0.9s/(0.9+k_{Iiq}))+(b_D c/(c+k_{Iiq}))+(bc/(c+k_f))$ where s = soluble proportion, k_{Iiq} = fractional outflow rate of the liquid pool (0.075/hr), b_D = degradable small particle proportion, b = degradable large particle proportion, c = fractional degradation rate of b, and k_f is the fractional outflow rate of the large particle pool (0.045/hr).
- 755 I TVC is the sum of acetic, butyric, lactic, propionic and valeric acids plus ethanol and propanol.
- 756 ¥TVFA is calculated as for TVC but excluding ethanol and propanol.
- 757 ^{a,b} Values within a row with different superscripts differ significantly at *P*<0.05.
- 758

759 **Table 3** The results of a validation in which 94 grass-clover silages were used to test

760 the prediction accuracy of grass-based NIRS equations for chemical composition

761 when used on clover-containing samples (in g/kg DM unless otherwise stated).

ltem	Measured mean	Predicted mean	Bias†	P value ‡	r²§	Relative RMSEP, %¶	RPD
ADF	336	292	43.0	0.001	0.61	17.6	0.87
aNDF	448	438	9.65	0.209	0.56	8.9	1.45
Ash	96.6	91.6	5.0	0.033	0.52	16.5	1.32
CP	138	126	12.4	0.005	0.75	17.1	1.58
DOMD	632	645	-13.0	0.195	0.64	6.7	1.56
EDN _{FIM} , g/kg N II	623	762	139	0.001	0.01	24.5	0.48
EE	26.5	30.1	-3.6	0.001	0.25	25.9	0.89
LA, g/kg	13.4	14.3	-0.9	0.622	0.48	70.6	1.22
MDM _{FIM} II	129	146	-17	0.003	0.01	38.1	0.39
N, g/kg	8.8	8.1	0.7	0.187	0.86	19.4	2.35
NH₃-N, g/kgDM*100	62.5	85.2	-22.6	0.001	0.34	45.0	0.89
рН	4.6	4.8	-0.1	0.122	0.48	10.8	1.21
TVC, g/kg ¥	23.4	30.2	-6.8	0.001	0.52	39.3	1.15
TVFA, g/kg #	19.6	25.6	-5.9	0.001	0.51	43.5	1.17
VCODM, g/kg	397	409	-12.0	0.558	0.98	6.6	4.92
WSC	41.8	48.8	-7.0	0.113	0.40	58.4	1.25

762 DOMD = digestible organic matter in total dry matter; EDN = effective degradable nitrogen; EE = ether 763 extract; LA = lactic acid; MDM = microbial dry matter yield; NH₃-N = ammonia nitrogen; RMSEP = root 764 mean standard error of prediction; RPD = ratio of standard deviation of the measured population to

765 the standard error of prediction; TVC = total volatile content, TVFA = total volatile fatty acids; WSC = 766 water soluble carbohydrates.

767 † Bias is the measured mean minus the predicted mean, therefore minus values indicate over-

768 estimation and positive values indicate under-estimation of the equation.

769 **‡** The probability of there being no significant difference between the measured mean and the

770 predicted mean analysed using student's t-test.

771 § Simple linear regression coefficient

772 ¶ Root mean square error of prediction presented as a percentage of the measured mean for 773 standardisation

774 || Degradability parameters determined by *in situ* incubation in the rumen, using the Feed Into Milk

(FIM) Consortium (2004) model $ED_{FIM} = (0.9s/(0.9+k_{Iiq}))+(b_D c/(c+k_{Iiq}))+(bc/(c+k_f))$ where s = soluble

775 776 proportion, k_{lig} = fractional outflow rate of the liquid pool (0.075/hr), b_D = degradable small particle

proportion, b = degradable large particle proportion, $c = fractional degradation rate of b, and k_f is the$ 777 fractional outflow rate of the large particle pool (0.045/hr). 778

779 ¥TVFA is calculated as for TVC but excluding ethanol and propanol.

- 780 # TVC is the sum of acetic, butyric, lactic, propionic and valeric acids plus ethanol and propanol.
- 781 782

783 **Table 4** Indicators of calibration strength and prediction accuracy using cross-

validation for a range of optimised new NIRS equations calibrated on spectra from

785 95 diverse grass-clover silages.

Item†	n ‡	SEC	r²§	Relative SECV, % ¶
ADF (DM)	183	13.4	0.90	4.49
ADF (Fresh)	181	6.22	0.98	5.71
Alcohol II	178	1.08	0.83	37.1
aNDF (DM)	183	18.5	0.89	4.80
aNDF (Fresh)	182	7.79	0.98	5.26
Ash (DM)	185	10.4	0.70	12.5
Ash (Fresh)	179	3.30	0.91	11.1
DOMD	172	3.10	0.83	5.47
EDDM _{0.08} ¥	174	2.15	0.88	5.28
EDN _{0.08} ¥	174	3.93	0.79	7.03
EE (DM)	180	2.67	0.83	11.2
EE (Fresh)	179	0.94	0.90	10.8
LA	173	4.76	0.81	41.5
Ν	180	0.65	0.97	8.33
NH ₃ -N	176	0.01	0.88	18.8
рН	180	0.16	0.93	4.18
TVC#	185	5.39	0.82	27.9
TVFA ††	183	5.17	0.81	31.8
VCODM	181	7.17	1.00	2.10
WSC (DM)	180	10.1	0.92	31.4
WSC (Fresh)	181	4.62	0.93	29.6

786 EDDM = effective degradability of dry matter; EDN = effective degradable nitrogen; DOMD =

digestible organic matter in total dry matter; EE = ether extract; LA = lactic acid; $NH_3-N =$ ammonia nitrogen; SEC = standard error of calibration; SECV = standard error of cross-validation; TVC = total

volatile content, TVFA = total volatile fatty acids; WSC = water soluble carbohydrates.

790 † For variables that are measured on a dry sample (Ash, ADF, aNDF and WSC) two equations were
 791 produced, one predicting on a fresh basis and one on a DM basis.

- 792 **‡** The number of spectra that were included in the prediction equation.
- 793 § Simple linear regression coefficient
- 794 ¶ Standard error of cross validation presented as a percentage of the measured mean for
- 795 standardisation
- 796 || Alcohol is the sum of ethanol and propanol
- YDegradability parameters determined by *in situ* incubation in the rumen, using the model of Ørskov
- and McDonald (1979) where a = rapidly soluble material; b = non-soluble but degradable material; c =
- rate f degradation of b; effective degradability = a+b[c/(c+k)] where k = an assumed outflow rate of 0.08/hr.
- 801 # TVC is the sum of acetic, butyric, lactic, propionic and valeric acids plus ethanol and propanol.
- 802 *††* TVFA is calculated as for TVC but excluding ethanol and propanol.
- 803

Table 5 The results of a validation in which 10 grass-clover silages were used to test
the prediction accuracy of new clover/grass-based NIRS equations generated from
the spectra of 85 other grass-clover silages (in g/kg DM unless otherwise stated).
Industry standardised data pre-treatment methods were used (1st derivative and
SNVD scatter correction) in the calibration of these equations.

	Moasurod	Prodicted			Relative	
	moon	Fredicied	Bias†	<i>r</i> ² ‡	RMSEP,	RPD
Item	mean	mean			% §	
ADF	343	352	-9.28	0.93	7.31	2.94
ADF, g/kg	162	159	1.93	0.99	6.17	8.66
Alcohol, g/kg ¶	3.5	4.9	-1.41	0.19	75.7	0.93
aNDF	459	479	-20.6	0.85	8.02	2.15
aNDF, g/kg	212	215	-2.54	0.98	5.81	7.87
Ash	88.5	87.0	1.49	0.26	16.5	1.19
Ash, g/kg	39.2	40.6	-1.38	0.73	18.8	1.82
СР	120	120	-0.82	0.74	12.3	1.92
DOMD	637	637	-0.2	0.71	9.18	1.76
EDDM _{0.08}	452	438	14.7	0.68	16.3	1.58
EDN₀.₀ଃ, g/kg N ∥	600	588	11.9	0.92	6.74	3.43
EE	22.8	21.6	1.22	0.46	23.2	0.95
EE, g/kg	9.9	9.8	0.15	0.67	16.9	1.82
LA, g/kg	15.1	12.7	2.41	0.72	49.0	1.74
N, g/kg	8.9	9.0	-0.02	0.92	13.4	3.76
NH ₃ -N, g/kg DM*100	104	112	-8.01	0.64	25.0	1.64
рН	4.7	4.6	0.06	0.70	10.8	1.86
TVC, g/kg ¥	22.9	21.7	1.20	0.76	33.5	1.85
TVFA, g/kg #	19.5	18.4	1.03	0.73	37.0	1.86
VCODM, g/kg	451	448	2.87	0.99	2.46	14.2
WSC	51.7	58.7	-7.01	0.69	35.3	1.76
WSC, g/kg	22.4	27.2	-4.84	0.69	44.4	1.57

809 DOMD = digestible organic matter in total dry matter; EDN = effective degradable nitrogen; EE = ether

810 extract; LA = lactic acid; NH₃-N = ammonia nitrogen; RMSEP = root mean standard error of

811 prediction; RPD = ratio of standard deviation of the measured population to the standard error of

prediction; SNVD, standard normal variate de-trending; TVC = total volatile content, TVFA = total
 volatile fatty acids; WSC = water soluble carbohydrates.

814 † Bias is the measured mean minus the predicted mean, therefore minus values indicate over-

815 estimation and positive values indicate under-estimation of the equation.

816 **‡** Simple linear regression coefficient

817 § Root mean square error of prediction presented as a percentage of the measured mean for

818 standardisation

819 ¶ Alcohol is the sum of ethanol and propanol

820 || Degradability parameters determined by *in situ* incubation in the rumen, using the model of Ørskov

and McDonald (1979) where a = rapidly soluble material; b = non-soluble but degradable material; c =

rate f degradation of b; effective degradability = a+b[c/(c+k)] where k = an assumed outflow rate of

823 0.08/hr.

824 ¥TVC is the sum of acetic, butyric, lactic, propionic and valeric acids plus ethanol and propanol.

TVFA is calculated as for TVC but excluding ethanol and propanol.

827 Figure captions

828 Figure 1 The relationship between predicted and measured values where 94 grass-829 clover silages were utilised to assess prediction accuracy of grass-based near infra-830 red reflectance spectrometry (NIRS) equations for 15 chemical components when 831 used on clover-containing samples. Graphs for each chemical component show a line 832 of parity. VCODM = volatile corrected oven dry matter; TVFA = total volatile fatty acids; 833 TVC = total volatile content; DOMD = digestible organic matter in total dry matter; ADF 834 = acid detergent fibre; aNDF neutral detergent fibre; MDM = microbial dry matter yield; 835 NH₃-N = ammonia nitrogen; EDN = effective degradable nitrogen; WSC = water

- 836 soluble carbohydrate.
- **Figure 2** The relationship between bias and sample clover concentration in a
- 838 validation test where 94 grass-clover silages were utilised to assess prediction
- 839 accuracy of grass-based near infra-red reflectance spectroscopy (NIRS) equations
- for a) crude protein b) effective degradable nitrogen (EDN_{FIM}) and c) calculated
- 841 effective rumen degradable protein (eRPD) concentration (eRDP = CP * (0.8 *
- 842 EDN_{FIM})). Linear lines of best fit are shown for measured (——) and NIRS predicted
- 844
- 843 (— —) data.