Journal Pre-proof

Visible and NIR hyperspectral imaging and chemometrics for prediction of microbial quality of beef *Longissimus dorsi m.* under simulated normal and abuse storage conditions

Achata Em, Oliveira M, Esquerre Ca, Tiwari Bk, O'Donnell Cp

PII: S0023-6438(20)30452-7

DOI: https://doi.org/10.1016/j.lwt.2020.109463

Reference: YFSTL 109463

To appear in: LWT - Food Science and Technology

Received Date: 30 December 2019

Revised Date: 13 April 2020

Accepted Date: 18 April 2020

Please cite this article as: Em, A., M, O., Ca, E., Bk, T., Cp, O'Donnell., Visible and NIR hyperspectral imaging and chemometrics for prediction of microbial quality of beef *Longissimus dorsi m.* under simulated normal and abuse storage conditions, *LWT - Food Science and Technology* (2020), doi: https://doi.org/10.1016/j.lwt.2020.109463.

This is a PDF file of an article that has undergone enhancements after acceptance, such as the addition of a cover page and metadata, and formatting for readability, but it is not yet the definitive version of record. This version will undergo additional copyediting, typesetting and review before it is published in its final form, but we are providing this version to give early visibility of the article. Please note that, during the production process, errors may be discovered which could affect the content, and all legal disclaimers that apply to the journal pertain.

© 2020 Published by Elsevier Ltd.



CRediT authorship contribution statement

Eva Achata: Investigation, Data Curation, Formal analysis, Visualization, Writing - Original draft, **Marcia Oliveira:** Investigation, Data Curation **Carlos Esquerre:** Software, Writing - Review & Editing **Brijesh Tiwari:** Conceptualization, Methodology, Resources **Colm O'Donnell:** Conceptualization, Resources, Writing - Review & Editing, Supervision, Funding acquisition

Journal Pre-proof

1	Visible and NIR hyperspectral imaging and chemometrics for prediction of microbial
2	quality of beef Longissimus dorsi m. under simulated normal and abuse storage
3	conditions
4	Achata EM ¹ , Oliveira M ² , Esquerre CA ¹ , Tiwari BK ² , O'Donnell CP ¹
5	¹ School of Biosystems and Food Engineering, University College Dublin, Ireland
6	² Department of Food Chemistry & Technology, Teagasc Food Research Centre, Ashtown,
7	Dublin 15, Ireland
8	Abstract
9	There is a need to develop a rapid technique to provide real time information on the microbial
10	load of meat along the supply chain. Hyperspectral imaging (HSI) is a rapid, non-destructive
11	technique well suited to food analysis applications. In this study, HSI in both the visible and
12	near infrared spectral ranges, and chemometrics were studied for prediction of the bacterial
13	growth on beef Longissimus dorsi m. (LD) under simulated normal (4 °C) and abuse (10 °C)
14	storage conditions. Total viable count (TVC) prediction models were developed using partial
15	least squares regression (PLS-R), spectral pre-treatments, band selection and data fusion
16	methods. The best TVC prediction models developed for storage at 4 (RMSE_p 0.58 log
17	CFU/g, RPD _p 4.13, R_{p}^{2} 0.96), 10 °C (RMSE _p 0.97 log CFU/g, RPD _p 3.28, R_{p}^{2} 0.94) or at
18	either 4 or 10 °C (RMSE _p 0.89 log CFU/g, RPD _p 2.27, R^2_{p} 0.86) were developed using high-
19	level data fusion of both spectral regions. The use of appropriate spectral pre-treatments and
20	band selection methods was key for robust model development. This study demonstrated the
21	potential of HSI and chemometrics for real time monitoring to predict microbial growth on
22	LD along the meat supply chain.



24 **1. Introduction**

Journal Pre-proot

The Longissimus dorsi m. (LD) of beef is highly valued by consumers and is normally aged 25 to increase tenderness, juiciness and flavour. Meat processors generally age LD for 28 days or 26 longer to improve tenderness and flavour. However colour and microbial load are also 27 affected which potentially impacts product safety and shelf life (Borch, Kant-Muermans, & 28 Blixt, 1996; Vitale, Pérez-Juan, Lloret, Arnau, & Realini, 2014). The design and application 29 of quality and safety assurance systems are based on thorough risk analysis and control of 30 critical parameters through the entire life cycle of meat products including raw material 31 selection and control during processing and distribution. The temperature profiles during 32 transportation and at retail level are not within the direct control of meat processors and may 33 exceed recommended temperatures. Lack of temperature control from retail to the time of 34 preparation and consumption may also be an issue. In southern European countries 30% of 35 refrigerated foods were reported to be stored above 10 °C in retail cabinets and household 36 refrigerators (Nychas, Skandamis, Tassou, & Koutsoumanis, 2008). 37

38 Current microbiological methods are not suitable for real time monitoring of microbial contamination of meat. The traditional plate count technique is the most commonly used 39 method to monitor microbial load. However, it requires time consuming sample preparation 40 and analysis. The total viable count (TVC) method is an important microbiology indicator for 41 quality and safety evaluation of meat (Lytou, Panagou, & Nychas, 2016). The initial 42 microbial load of meat after processing, storage temperature, pH and relative humidity are the 43 main factors influencing microbial load throughout the supply chain. Enzyme-linked immune 44 absorbent assay (ELISA), gene analysis-based methods such as polymerase chain reaction 45 (PCR) and DNA sequencing are also employed for microbial contamination detection (Si et 46 al., 2016) but are not suited to online analysis. 47

Visible (VIS) and near-infrared (NIR) spectroscopy are rapid non-destructive techniques
widely used in environmental, pharmaceutical, fuel and food analysis applications. The VIS

and NIR spectral regions range from 380 - 740 nm and 700 - 2500 nm respectively.
Spectroscopic sensors usually acquire spectra from a limited field of view which limits their
applicability for rapid safety analysis of large volume batches or analysis of heterogeneous
samples such as meat products (Millar, Moss, & Stevenson, 1996).

Hyperspectral imaging (HSI) is a rapid analytical tool for non-destructive measurement of 54 food quality and safety. HSI integrates traditional imaging and spectroscopy to acquire both 55 56 spatial and spectral information from samples. Each pixel in a hyperspectral image contains 57 the spectrum of that specific position, i.e. the light-absorbing and/or scattering properties of the spatial region represented, which can be used to characterise the composition of that 58 59 particular pixel (Gowen, O'Donnell, Cullen, Downey, & Frias, 2007; Kamruzzaman, Makino, & Oshita, 2016). HSI techniques can be employed at different points along meat distribution 60 chains. HSI has been studied to predict microbial growth on fresh beef meat using the VIS 61 range (Peng et al., 2011; Tao, Peng, Gomes, Chao, & Qin, 2015). However, no studies have 62 been reported to date on the use of HSI in the NIR spectral range to predict microbial growth 63 64 on fresh beef.

Chemometric methods are employed to develop prediction models from HSI data. Partial 65 least squares regression (PLS-R) may be used to predict unknown concentrations and 66 67 generate prediction maps to estimate spatial distributions of components in samples (Gowen, Burger, Esquerre, Downey, & O'Donnell, 2014). Spectral pre-treatments are used to correct 68 for the effects of natural variability in the shape and size of samples, light scattering and 69 differences in the effective path length in spectral data, which can present difficulties in the 70 application of HSI for quality assessment (Esquerre, Gowen, Burger, Downey, & O'Donnell, 71 72 2012). Band selection methods have been demonstrated to improve the performance of regression models and to reduce the processing times required to evaluate HSI data by 73 selecting the most informative bands. The variable importance projection method (VIP), the 74 selectivity ratio method (SR) and the ensemble Monte Carlo variable selection method 75

76 (EMCVS) have been demonstrated to be reliable band selection methods for HSI data
77 (Achata, Inguglia, Esquerre, Tiwari, & O'Donnell, 2019; Farrés, Platikanov, Tsakovski, &
78 Tauler, 2015).

79 Data fusion combines information from different sources to produce a more reliable and accurate model or information. Three levels of data fusion may be employed i) low level 80 (data-level) fusion, where data from all sources are properly transformed and concatenated 81 82 for model development, ii) mid-level (feature-level) fusion, where variable selection or feature extraction is applied to each data source before the extracted features are combined; 83 iii) and high-level (decision-level) fusion where a model is constructed for each data source 84 separately and their predictions combined thereafter (Liu & Brown, 2004). Data fusion has 85 been studied to detect volatile basic nitrogen (TVB-N) content in chicken meat using a 86 colorimetric sensor and a VIS system (Khulal, Zhao, Hu, & Chen, 2017). 87

The objective of this study was to investigate the potential of HSI and chemometrics for the prediction of the microbial quality of beef under simulated normal (4 °C) and abuse (10 °C) storage conditions.

91

92 2. Materials and methods

93 2.1. Sample preparation

94 LD samples (n = 104) from 9 cattle (denoted S1 to S9) of ca. 25 mm thickness were obtained from local supermarkets and a meat processing facility. The samples were placed in sealed 95 food containers and randomly assigned for storage at either 4 $^{\circ}C$ (n = 53) for 360 hours or at 96 10 °C (n = 51) for 168 hours. Three randomly selected samples (from 3 cattle) were removed 97 from storage and scanned using a visible short wave near infrared (VIS-SWNIR) and an NIR 98 99 HSI systems. The TVC of samples was measured after scanning using the ISO 48833-1:2013 methodology (ISO, 2013). Briefly 25 g of each sample was suspended in 225 ml of buffered 100 peptone water (BPW, Oxoid, Hampshire, England) and aseptically homogenized in a 101

102	stomacher (Star-Blender LB 400, VWR) for 2 min. Further decimal dilutions were made with
103	maximum recovery diluent (MRD, HyServe, Germany). Three replicates were assessed per
104	sample at each sampling time. Reported populations represent the mean of three values.

105

106 **2.2. Hyperspectral images**

Hyperspectral images of the *LD* samples were obtained using a VIS-SWNIR HSI system (400 – 1000 nm) and an NIR HSI system (880 – 1720 nm) (DV Optics, Padova, Italy). Calibration of both HSI systems was performed as outlined by Achata, Esquerre, O'Donnell, and Gowen (2015). The acquired hypercubes were saved in ENVI formatted files and imported into MATLAB (The MathWorks Inc., Natick, MA, USA) for further spatial and spectral data pre-processing and chemometric analysis, using in-house developed functions and scripts.

114

115 2.2.1. VIS-SWNIR HSI spatial and spectral pre-processing

The noise present at both ends of the spectra was removed by trimming the spectral range to 116 445 - 970 nm. The background was removed using a mask created using the ratio between 117 bands 80 (840 nm) and 20 (540 nm) and removing pixels with a ratio value > 1.5. To improve 118 119 the signal-to-noise ratio (SNR) and reduce processing times and data storage required, 2×2 binning was performed on the obtained hypercubes of 1000 x 580 pixel image with 106 120 spectral bands, resulting in hypercubes of 500 x 290 pixel image with 106 spectral bands. The 121 binned 3-dimensional hypercubes were unfolded into matrices of pixel spectra (14500 pixel x 122 106 spectral bands) to facilitate algorithm development. The mean reflectance (R) spectra of 123 each masked sample was calculated and smoothed using the Savitzky - Golay (SG) 5 points 124 second order polynomial method prior to chemometric analysis (Savitzky & Golay, 1964). 125

127 2.2.2. NIR-HSI spatial and spectral pre-processing

The noise present at both ends of the spectra was removed by trimming the spectral range to 128 957 - 1664 nm. Dead pixels and spikes were removed by replacing the affected values with 129 the mean values of adjacent bands in the same spectrum. The background was removed using 130 a mask which was created with the ratio between bands 90 (1580 nm) and 20 (1090 nm), 131 removing pixels with a ratio value > 0.65. Images were segmented using the pixel ratio 132 between bands 37 (1209 nm) and 43 (1251 nm) to remove fat and connective tissue (ratio 133 value > 0.7). The 3-dimensional hypercubes (500 x 320 pixel image with 102 bands) were 134 unfolded into matrices of pixel spectra (160000 pixel x 102 bands). The mean reflectance 135 spectra of each segmented sample was calculated and smoothed using the Savitzky - Golay 136 (SG) 5 points second order polynomial method prior to chemometric analysis. 137

138

139 **2.3.** Chemometric analysis

140 **2.3.1.** PCA

141 PCA (not reported) was carried out to investigate the relationships between storage 142 temperature over time and spectral data, and to identify potential outliers using the Hoteling 143 T^2 statistic. A sample was considered as an outlier if the T^2 value was > T^2 crit = $A \times F_{(0.05,A,n - A)}$ 144 $A_{A} \times (n-1)/(n-A)$, where A is the number of significant components, n is the number of spectra 145 in the dataset and $F_{(0.05,A,n - A)}$ is the F statistic (with $\alpha = 0.05$, A and n – A degrees of 146 freedom).

147

```
148 2.3.2. PLS-R
```

PLS regression (Wold, Sjöström, & Eriksson, 2001) models were developed to predict TVC
of samples using HSI data, spectral pre-treatments, band selection and data fusion methods.

- 151 Spectral data sets were split into calibration and validation sets to develop and validate the
- 152 prediction models. Smoothed mean spectral data from 4 randomly selected samples (S1, S2,
- 153 S4 and S6 (n=69)) was used for calibration and samples (S3 and S5) were used to validate the
- 154 models (n=35). Predictions models were evaluated using the:
- i) Smoothed mean spectral data of samples stored at 4 $^{\circ}C$ (n = 53)
- 156 ii) Smoothed mean spectral data of samples stored at 10 $^{\circ}$ C (n = 51)
- 157 iii) Smoothed mean spectral data of samples stored at either 4 or 10 $^{\circ}$ C (n = 104).
- 158 The number of latent variables (LV) were selected by analysis of the root mean square error
- of ten-fold cross-validation (RMSE_{CV}) presented in Eq. (1) and roughness of the regression
 vector.

161
$$RMSE = \sqrt{\frac{\sum_{i=1}^{n} (y_i - \hat{y}_i)^2}{n}}$$
 (1)

- 162 y_i and \hat{y}_i are the measured and predicted values of the microbial counts respectively.
- 163

164 **2.3.3. Spectral pre-treatments**

Standard normal variate (SNV), median scaled (MS), Savitzky-Golay 7 points second order 165 polynomial first derivative (FD), Savitzky-Golay 7 points second order polynomial second 166 derivative (SD), Savitzky-Golay 11 points fourth order polynomial third derivative (TD), 167 linear detrending second-order polynomial (LD), asymmetric least squares (AsLs) (Barnes, 168 Dhanoa, & Lister, 1989; Boelens, Eilers, & Hankemeier, 2005; Engel et al., 2013; Savitzky & 169 Golay, 1964) and all combinations of any two spectral pre-treatments were applied. The 170 Savitzky-Golay derivative (FD, SD or TD) window length and polynomial order were 171 selected by preliminary tests on 10 randomly selected spectra. 172

173

174 **2.3.4. Band selection**

The VIP (Eriksson, Hermens, Johansson, Verhaar, & Wold, 1995; Wold et al., 2001), SR (Rajalahti et al., 2009) and the EMCVS (Esquerre, Gowen, O'Gorman, Downey, & O'Donnell, 2017) band selection methods were evaluated and compared with and without spectral pre-treatments.

The performance of the regression models was assessed using the root mean square error 179 (RMSE), the ratio of standard deviation of the reference data of the calibration set and the 180 RMSE (RPD) and the coefficient of determination (R^2) for calibration (C), cross-validation 181 (CV) and prediction (P) sets (Eq. 2-4). The best model was selected based on the number of 182 latent variables, selected wavebands and the geometric mean of the RPD values from 183 184 calibration, cross-validation, and prediction sets. Prediction models developed for complex matrices can be classified as excellent (RPD > 4.1), very good (RPD 3.5 - 4.0), good (RPD 3.0 - 3.4), 185 fair (RPD 2.5 – 2.9) and poor (RPD 2.0 – 2.4) (Williams, 2014). 186

187
$$RPD = \frac{s_{ycal}}{SEP}$$
 (2)

188
$$SEP = \sqrt{\frac{\sum_{i=1}^{n} (y_i - \hat{y}_i - \text{bias})^2}{n-1}}$$
 (3)

189
$$R^{2} = \left(\frac{\sum_{i=1}^{n} (y_{i} \hat{y}_{i}) - n\bar{y} \,\bar{y}}{\sqrt{\sum_{i=1}^{n} y_{i}^{2} - n\bar{y}^{2}} \sqrt{\sum_{i=1}^{n} \hat{y}_{i}^{2} - n\bar{y}^{2}}}\right)^{2} \tag{4}$$

190 Where the *bias* is the average difference between reference value and predicted value (Eq. 5).

191
$$bias = \frac{\sum_{i=1}^{n} (y_i - \hat{y}_i)}{n}$$
 (5)

192

193 **2.3.5. Data fusion**

Prediction models for TVC of samples were developed for low level (LL), medium level (ML) and high-level (HL) data fusion of VIS-SWNIR and NIR HSI data. For the LL data fusion, the spectral data of both systems was concatenated before model development. For the ML data fusion, the selected spectral bands (obtained by the band selection method that achieved the best performance for each system) were combined. When the variance of the pre-treated VIS-SWNIR and NIR HSI data was different, each data block was scaled to unit variance. The calibration, validation and prediction sets were scaled using the inverse of the standard deviation of the calibration set (Forshed, Idborg, & Jacobsson, 2007). For the HL data fusion, prediction models were developed by averaging the predictions of the best performing models identified for each system.

204

205 **3. Results and discussion**

206 3.1. **TVC of samples**

The TVC of LD samples during storage at 4 °C and 10 °C are presented in Table 1. TVC 207 values increased from 3.4 to 14.1 log CFU/g over 360 h storage at 4 °C and increased from 208 3.4 to 13.1 log CFU/g over 168 h storage at 10 °C. Initial TVC values (day 0) varied 209 according to the origin of the LD samples. LD S1 to S6 samples were purchased from local 210 supermarkets and had initial TVC values $> 6 \log \text{ CFU/g}$. However, *LD* samples purchased 211 directly from the meat processing facility (S7 to S9) had lower initial TVC values. LD S9 212 (non aged) samples had the lowest TVC values (3.4 log CFU/g), while samples from the LD 213 S7 and S8 which were both aged for 28 days had TVC values of ca. 5.3 log CFU/g. 214

Previous studies reported that *LD* samples with TVC values < 7 log CFU/g are acceptable,
and samples with values > 7 log CFU/g are spoiled (Tao et al., 2015). Moreover, the presence
of slime and discolouration has been reported for meat samples with TVC values > 7 log
CFU/g (Bell & Garout, 1994).

219

220 **3.2.** Spectral characteristics of *LD* samples

PCA analysis revealed the presence of one outlier in the VIS-SWNIR spectra of the *LD* samples stored at 4 °C, which was removed from the dataset. No outliers were identified in the NIR spectra. Figs. 1 and 2 show the spectral variations between samples during storage at 4 and 10 °C respectively. The SD + AsLs pre-treated log (1/R) NIR spectra were used for TVC prediction at 4 °C, whereas the SD+MS pre-treated reflectance VIS-SWNIR spectra
were selected for TVC prediction at 10 °C. Spectral shifts are more apparent during storage
due to changes in physical characteristics, chemical composition and microbial activity.
Spoiled samples exhibited broader absorption bands compared to unspoiled samples.

Absorbance peaks observed at 1076 and 1342 nm in Fig. 1 may be related to the C-H 229 stretching of the first and second overtone regions respectively, and the peak at 1580 nm may 230 be related to the 1st overtone of O-H stretching (glucose) (Osborne, Fearn, & Hindle, 1993). 231 The selected bands highlighted in Fig.1 provide complementary information on the samples 232 and are related to the 2nd overtone of O-H stretching (978 nm) of water and the 1st overtone of 233 N-H stretching (1496 nm) of protein. Similar spectral bands were observed by Barbin, 234 ElMasry, Sun, Allen, and Morsy (2013) for pork samples. The observed differences between 235 spoiled and fresh meat may relate to the presence of protein, free amino acids, amines or 236 nitrogen bearing substances and their interactions with water. Such observations are 237 consistent with the proteolytic changes which occur during microbial spoilage (Atanassova, 238 Veleva, & Stoyanchev, 2018). 239

Fig. 2 shows the characteristic peaks of the of oxymyoglobin (MbFe^{II}O₂) at 545 and 580 nm 240 (Achata et al., 2019; Alamprese, Casale, Sinelli, Lanteri, & Casiraghi, 2013; Millar et al., 241 1996). These bands are prominent at the start of storage and increase in intensity after 24 h 242 storage at 10 °C (TVC $< 7.3 \log \text{ CFU/g}$). The intensity of these bands decreases after 48 h 243 corresponding to TVC values $> 7.5 \log CFU/g$. These changes may correspond to a decrease 244 in the concentration of red pigments due to microbial growth during storage. The prominent 245 peak at 765 nm may be related to the 3rd overtone C-H stretching. The selected bands 246 highlighted in Fig. 2 provide complementary information on the samples and may be related 247 to the 3rd overtone of C-H stretching (750 - 780 nm) (Osborne et al., 1993). 248

3.3. TVC prediction models

SD, SNV and the combination of SD+LD, and SNV+SD were identified as the best performing spectral pre-treatments after evaluating 50 combinations for each band selection method (Appendix 1). Models developed using the variable selection approach were compared with the best models developed using the full spectral range for both the VIS-SWNIR and NIR HSI spectral regions.

The best performing PLS-R model developed to predict TVC during storage at 4 °C (Table 256 2) was developed using the NIR-HSI data and EMCVS of the SD+AsLs pre-treated log(1/R)257 spectra (17 selected bands, LV 7, RMSE_P 0.81 log CFU/g, RPD_P 3.09, R²_P 0.95). Lower 258 coefficients of determination for the prediction of pork meat TVC were obtained by Barbin 259 et al. (2013). Fig. 3a shows the predicted versus measured TVC values for LD samples 260 stored at 4 °C obtained with the SD+AsLs pre-treated log(1/R) spectra. HL data fusion 261 improved the performance of the prediction models (RMSE_P 0.58 log CFU/g, RPD_P 4.13, 262 R_{P}^{2} 0.96) compared to those obtained with LL data fusion, ML data fusion and the best 263 models selected for the VIS-SWNIR and NIR HSI spectral data (Table 5). 264

The best performing PLS-R model developed to predict TVC during storage at 10 °C (Table 265 3) was developed using the VIS-SWNIR - HSI data and EMCVS of the SD+MS pre-treated 266 reflectance spectra (46 selected bands, LV 6, RMSE_P 0.96 log CFU/g, RPD_P 3.32, R²_P 0.94). 267 Fig.3b. shows the predicted versus measured TVC values for LD samples stored at 10 °C 268 obtained with the SD+MS pre-treated reflectance spectra. The use of derivative pre-269 270 treatments of VIS-SWNIR spectra has been reported to accentuate the differences in myoglobin spectra (Millar et al., 1996) by removing baseline offsets and decreasing 271 scattering effects (Esquerre et al., 2012) as observed in Fig. 2. LL and HL data fusion 272 vielded good prediction models, comparable to those obtained with the VIS-SWNIR data 273 and better than the models developed using ML data fusion (Table 5). 274

The best performing PLS-R models developed to predict TVC for samples stored at either 4 275 or 10 °C (Table 4) were developed using the VIS-SWNIR data and EMCVS of the SNV+SD 276 pre-treated reflectance spectra (8 selected bands, LV 4, RMSEP 0.95 log CFU/g, RPD_P 2.10, 277 R_{P}^{2} 0.85). Improved data fusion PLS-R models were developed using the LL and HL data 278 fusion showed in Table 5 (RMSE_P 0.87 log CFU/g, RPD_P 2.27, R²_P 0.88, and RMSE_P 0.89 279 log CFU/g. RPD_P 2.27, R²_P 0.86 respectively). Fig. 3c shows the predicted versus measured 280 TVC values of *LD* stored at either 4 or 10 °C obtained using HL data fusion of both spectral 281 regions. 282

Selected TVC prediction maps built using the best prediction model developed for storage at
10 °C (SD+MS on the reflectance VIS-SWNIR spectra) are shown in Fig. 4.

Good and excellent TVC prediction models were obtained for beef LD samples stored at 4 285 °C for both the NIR spectral range and HL data fusion of the VIS-SWNIR and NIR spectral 286 regions respectively based on the RPD prediction model performance classifications reported 287 by Williams (2014) for complex matrices. Good TVC prediction models were also obtained 288 for samples stored at 10 °C using the VIS-SWNIR spectral range and HL data fusion of both 289 spectral regions. However poor TVC prediction models were obtained for samples stored at 290 either 4 °C or 10 °C using the VIS-SWNIR spectral range and HL data fusion of both 291 spectral regions. In all cases EMCVS outperformed the other band selection methods 292 evaluated. Combinations of SD, SNV and LD spectral pre-treatments also improved 293 regression model development. Data fusion approaches improved prediction model 294 performance in all cases. This is in agreement with the study reported by (Li, Chen, Zhao, 295 and Wu (2015)) who reported that superior regression models can be obtained using data 296 fusion and appropriate band selection methods. 297

298

300 4. Conclusions

Excellent (RMSE_p 0.58 log CFU/g, RPD_p 4.13, R^2_p 0.96) and good (RMSE_p 0.97 log CFU/g, RPD_p 3.28, R^2_p 0.94) TVC prediction models were developed for beef *LD* samples stored at 4 °C and 10 °C respectively, using the HL data fusion of the two spectral regions. Prediction models were successfully developed using spectral pre-treatments, the full spectral range, selected bands and data fusion of both VIS-SWNIR and NIR spectral regions to predict the TVC of *LD* samples with low prediction errors.

The application of SD and SNV spectral pre-treatments improved the performance of the 307 308 developed models using both spectral ranges and on selected bands. EMCVS improved the performance of the TVC prediction models developed compared with the full spectral range 309 and outperformed VIP and SR methods. Data fusion approaches improved prediction model 310 performance in all cases, HL data fusion yielded the best TVC prediction models (RMSE_p 311 0.89 log CFU/g, RPD_p 2.27, R_p^2 0.86) for beef samples stored at both 4 °C and 10 °C. 312 Appropriate band selection was key for robust model development. This study demonstrated 313 the potential of HSI and chemometrics as a rapid analytical tool for monitoring meat 314 microbial quality along the supply chain. 315

316

317 Acknowledgement

The authors acknowledge funding for this project from FIRM (13/FM/508) as administeredby the Irish Department of Agriculture, Food & the Marine.

320

Journal Pre-proof

322 **References**

- Achata, E. M., Esquerre, C. A., O'Donnell, C. P., & Gowen, A. A. (2015). A study on the
 application of near infrared hyperspectral chemical imaging for monitoring moisture
 content and water activity in low moisture systems. *Molecules*, 20(2), 2611.
- Achata, E. M., Inguglia, E. S., Esquerre, C. A., Tiwari, B. K., & O'Donnell, C. P. (2019).
 Evaluation of Vis-NIR hyperspectral imaging as a process analytical tool to classify
 brined pork samples and predict brining salt concentration. *Journal of Food Engineering*, 246, 134-140. doi: 10.1016/j.jfoodeng.2018.10.022
- Alamprese, C., Casale, M., Sinelli, N., Lanteri, S., & Casiraghi, E. (2013). Detection of
 minced beef adulteration with turkey meat by UV–vis, NIR and MIR spectroscopy.
 LWT Food Science and Technology, 53(1), 225-232. doi:
 https://doi.org/10.1016/j.lwt.2013.01.027
- Atanassova, S., Veleva, P., & Stoyanchev, T. (2018). Chapter 10 Near-Infrared Spectral
 Informative Indicators for Meat and Dairy Products, Bacterial Contamination, and
 Freshness Evaluation. In A. M. Holban & A. M. Grumezescu (Eds.), *Microbial Contamination and Food Degradation* (pp. 315-340): Academic Press.
- Barbin, D. F., ElMasry, G., Sun, D.-W., Allen, P., & Morsy, N. (2013). Non-destructive
 assessment of microbial contamination in porcine meat using NIR hyperspectral
 imaging. *Innovative Food Science & Emerging Technologies*, 17, 180-191. doi:
 https://doi.org/10.1016/j.ifset.2012.11.001
- Barnes, R. J., Dhanoa, M. S., & Lister, S. J. (1989). Standard Normal Variate Transformation
 and De-Trending of Near-Infrared Diffuse Reflectance Spectra. *Applied Spectroscopy*,
 43(5), 772-777. doi: 10.1366/0003702894202201
- Bell, R. G., & Garout, A. M. (1994). The effective product life of vacuum-packaged beef
 imported into Saudi Arabia by sea, as assessed by chemical, microbiological and
 organoleptic criteria. *Meat Science*, *36*(3), 381-396. doi: https://doi.org/10.1016/03091740(94)90134-1
- Boelens, H. F. M., Eilers, P. H. C., & Hankemeier, T. (2005). Sign Constraints Improve the
 Detection of Differences between Complex Spectral Data Sets: LC–IR As an
 Example. Analytical Chemistry, 77(24), 7998-8007. doi: 10.1021/ac051370e
- Borch, E., Kant-Muermans, M.-L., & Blixt, Y. (1996). Bacterial spoilage of meat and cured
 meat products. *International Journal of Food Microbiology*, *33*(1), 103-120. doi:
 https://doi.org/10.1016/0168-1605(96)01135-X
- Engel, J., Gerretzen, J., Szymańska, E., Jansen, J. J., Downey, G., Blanchet, L., & Buydens,
 L. M. C. (2013). Breaking with trends in pre-processing? *TrAC Trends in Analytical Chemistry*, 50, 96-106. doi: https://doi.org/10.1016/j.trac.2013.04.015
- Eriksson, L., Hermens, J. L. M., Johansson, E., Verhaar, H. J. M., & Wold, S. (1995).
 Multivariate analysis of aquatic toxicity data with PLS. *Aquatic Sciences*, 57(3), 217-241. doi: 10.1007/BF00877428
- Esquerre, C., Gowen, A. A., Burger, J., Downey, G., & O'Donnell, C. P. (2012). Suppressing
 sample morphology effects in near infrared spectral imaging using chemometric data
 pre-treatments. [Article]. *Chemometrics and Intelligent Laboratory Systems*, 117,
 129-137. doi: 10.1016/j.chemolab.2012.02.006
- Esquerre, C. A., Gowen, A. A., O'Gorman, A., Downey, G., & O'Donnell, C. P. (2017). 365 Evaluation of ensemble Monte Carlo variable selection for identification of metabolite 366 367 markers on NMR data. Analytica Chimica Acta, 964, 45-54. doi: https://doi.org/10.1016/j.aca.2017.01.027 368
- Farrés, M., Platikanov, S., Tsakovski, S., & Tauler, R. (2015). Comparison of the variable
 importance in projection (VIP) and of the selectivity ratio (SR) methods for variable

- 371 selection and interpretation. Journal of Chemometrics, 29(10), 528-536. doi: 10.1002/cem.2736 372 Forshed, J., Idborg, H., & Jacobsson, S. P. (2007). Evaluation of different techniques for data 373 374 fusion of LC/MS and 1H-NMR. Chemometrics and Intelligent Laboratory Systems, 85(1), 102-109. doi: https://doi.org/10.1016/j.chemolab.2006.05.002 375 Gowen, A., Burger, J., Esquerre, C., Downey, G., & O'Donnell, C. (2014). Near infrared 376 377 hyperspectral image regression: On the use of prediction maps as a tool for detecting model overfitting. [Article]. Journal of Near Infrared Spectroscopy, 22(4), 261-270. 378 379 doi: 10.1255/jnirs.1114 380 Gowen, A. A., O'Donnell, C. P., Cullen, P. J., Downey, G., & Frias, J. M. (2007). Hyperspectral imaging - an emerging process analytical tool for food quality and 381
- Hyperspectral imaging an emerging process analytical tool for food quality and safety control. *Trends in Food Science and Technology*, 18(12), 590-598. doi: 10.1016/j.tifs.2007.06.001
 10.0101/j.tifs.2007.06.001
- ISO. (2013). International standard ISO 4833-2:2013: Microbiology of the food chain Horizontal method for the enumeration of microorganisms Part 1: Colony count at
 30 °C by the pour plate technique.
- Kamruzzaman, M., Makino, Y., & Oshita, S. (2016). Hyperspectral imaging for real-time
 monitoring of water holding capacity in red meat. *LWT Food Science and Technology*, 66, 685-691. doi: https://doi.org/10.1016/j.lwt.2015.11.021
- Khulal, U., Zhao, J., Hu, W., & Chen, Q. (2017). Intelligent evaluation of total volatile basic
 nitrogen (TVB-N) content in chicken meat by an improved multiple level data fusion
 model. *Sensors and Actuators B: Chemical*, 238, 337-345. doi:
 https://doi.org/10.1016/j.snb.2016.07.074
- Li, H., Chen, Q., Zhao, J., & Wu, M. (2015). Nondestructive detection of total volatile basic
 nitrogen (TVB-N) content in pork meat by integrating hyperspectral imaging and
 colorimetric sensor combined with a nonlinear data fusion. *LWT Food Science and Technology*, 63(1), 268-274. doi: https://doi.org/10.1016/j.lwt.2015.03.052
- Liu, Y., & Brown, S. D. (2004). Wavelet multiscale regression from the perspective of data
 fusion: new conceptual approaches. [journal article]. *Analytical and Bioanalytical Chemistry*, 380(3), 445-452. doi: 10.1007/s00216-004-2776-x
- 401 Lytou, A., Panagou, E. Z., & Nychas, G.-J. E. (2016). Development of a predictive model for
 402 the growth kinetics of aerobic microbial population on pomegranate marinated
 403 chicken breast fillets under isothermal and dynamic temperature conditions. *Food*404 *Microbiology*, 55, 25-31. doi: https://doi.org/10.1016/j.fm.2015.11.009
- Millar, S. J., Moss, B. W., & Stevenson, M. H. (1996). Some observations on the absorption
 spectra of various myoglobin derivatives found in meat. *Meat Science*, 42(3), 277288. doi: https://doi.org/10.1016/0309-1740(94)00045-X
- 408 Nychas, G.-J. E., Skandamis, P. N., Tassou, C. C., & Koutsoumanis, K. P. (2008). Meat
 409 spoilage during distribution. *Meat Science*, 78(1), 77-89. doi:
 410 https://doi.org/10.1016/j.meatsci.2007.06.020
- 411 Osborne, B. G., Fearn, T., & Hindle, P. T. (1993). *Practical NIR spectroscopy with*412 *applications in food and beverage analysis*. Essex, England: Longman Scientific &
 413 Technical; Wiley.
- Peng, Y., Zhang, J., Wang, W., Li, Y., Wu, J., Huang, H., . . . Jiang, W. (2011). Potential
 prediction of the microbial spoilage of beef using spatially resolved hyperspectral
 scattering profiles. *Journal of Food Engineering*, 102(2), 163-169. doi:
 https://doi.org/10.1016/j.jfoodeng.2010.08.014
- Rajalahti, T., Arneberg, R., Kroksveen, A. C., Berle, M., Myhr, K.-M., & Kvalheim, O. M.
 (2009). Discriminating Variable Test and Selectivity Ratio Plot: Quantitative Tools
 for Interpretation and Variable (Biomarker) Selection in Complex Spectral or

Journal Pre-proof

- 421
 Chromatographic
 Profiles.
 Analytical
 Chemistry,
 81(7),
 2581-2590.
 doi:

 422
 10.1021/ac802514y
 10.1021/ac802514y
 10.1021/ac802514y
 10.1021/ac802514y
- 423 Savitzky, A., & Golay, M. J. E. (1964). Smoothing and Differentiation of Data by Simplified
 424 Least Squares Procedures. *Analytical Chemistry*, *36*(8), 1627-1639. doi:
 425 10.1021/ac60214a047
- Si, Y., Grazon, C., Clavier, G., Rieger, J., Audibert, J.-F., Sclavi, B., & Méallet-Renault, R.
 (2016). Rapid and accurate detection of Escherichia coli growth by fluorescent pHsensitive organic nanoparticles for high-throughput screening applications. *Biosensors and Bioelectronics*, 75, 320-327. doi: https://doi.org/10.1016/j.bios.2015.08.028
- Tao, F., Peng, Y., Gomes, C. L., Chao, K., & Qin, J. (2015). A comparative study for
 improving prediction of total viable count in beef based on hyperspectral scattering
 characteristics. *Journal of Food Engineering*, *162*, 38-47. doi:
 https://doi.org/10.1016/j.jfoodeng.2015.04.008
- Vitale, M., Pérez-Juan, M., Lloret, E., Arnau, J., & Realini, C. E. (2014). Effect of aging time
 in vacuum on tenderness, and color and lipid stability of beef from mature cows
 during display in high oxygen atmosphere package. *Meat Science*, 96(1), 270-277.
 doi: https://doi.org/10.1016/j.meatsci.2013.07.027
- Williams, P. (2014). The RPD Statistic: A Tutorial Note. *NIR news*, 25(1), 22-26. doi: 10.1255/nirn.1419
- Wold, S., Sjöström, M., & Eriksson, L. (2001). PLS-regression: a basic tool of chemometrics. *Chemometrics and Intelligent Laboratory Systems*, 58(2), 109-130. doi: https://doi.org/10.1016/S0169-7439(01)00155-1

Journe

Appendix 1. Performance of the TVC PLS-R models developed using VIS-SWNIR (445 - 970 nm) data from beef *LD* samples stored at 4 °C. PLS full spectral range on reflectance (R) and logarithmic transformed (log(1/R)) spectral data is compared with spectral pre-treatments (SNV, SD, SD+LD, SNV+SD) and band selection methods (VIP, SR and EMCVS).

	Spectral	#	#	С	alibration		Cros	ss validati	on	Prediction			
Chem mo	nometric ethod	Pre- treatment	Bands	LV	RMSEc	RPDc	R ² c	RMSEcv	RPDcv	R ² cv	RMSEp	RPDp	R ² p
		None	106	7	0.91	2.73	0.87	1.48	1.69	0.68	1.72	<mark>1.20</mark>	0.72
		SNV	106	7	0.84	2.97	0.89	1.20	2.08	0.78	1.62	<mark>1.37</mark>	0.78
	R	SD	100	8	0.61	4.09	0.94	1.08	2.30	0.82	2.16	<mark>1.21</mark>	0.71
		SD+LD	100	8	0.61	4.10	0.94	1.09	2.30	0.82	2.05	<mark>1.23</mark>	0.71
		SNV+SD	100	6	0.79	3.15	0.90	1.34	1.86	0.74	1.35	<mark>1.61</mark>	0.81
FL3	Log(1/R)	None	106	3	1.27	1.96	0.74	1.59	1.56	0.60	1.79	<mark>1.20</mark>	0.58
		SNV	106	5	0.89	2.79	0.87	1.16	2.16	0.78	1.45	<mark>1.45</mark>	0.75
		SD	100	6	0.63	3.94	0.94	1.06	2.37	0.82	1.93	<mark>1.22</mark>	0.67
		SD+LD	100	6	0.64	3.87	0.93	1.05	2.39	0.82	1.83	<mark>1.25</mark>	0.67
		SNV+SD	100	4	0.82	3.06	0.89	1.24	2.04	0.76	1.45	<mark>1.44</mark>	0.75
		None	7	2	1.27	1.97	0.74	1.41	1.77	0.68	1.79	<mark>1.33</mark>	0.66
		SNV	27	6	0.97	2.58	0.85	1.20	2.07	0.77	1.80	<mark>1.25</mark>	0.75
	R	SD	28	7	0.94	2.65	0.86	1.38	1.81	0.70	1.78	<mark>1.25</mark>	0.70
		SD+LD	33	10	0.51	4.89	0.96	0.90	2.79	0.87	2.50	<mark>1.40</mark>	0.74
		SNV+SD	18	11	0.46	5.47	0.97	0.79	3.18	0.90	1.31	<mark>1.92</mark>	0.81
VIP		None	4	3	1.21	2.06	0.76	1.39	1.79	0.70	1.77	<mark>1.25</mark>	0.60
		SNV	17	11	0.84	2.98	0.89	1.21	2.07	0.77	1.65	<mark>1.33</mark>	0.71
	Log(1/R)	SD	18	6	0.79	3.16	0.90	1.13	2.21	0.80	1.72	<mark>1.30</mark>	0.68
		SD+LD	15	3	1.10	2.27	0.81	1.28	1.95	0.74	1.34	<mark>1.66</mark>	0.76
		SNV+SD	13	8	0.55	4.51	0.95	0.70	3.56	0.92	1.38	<mark>1.86</mark>	0.80

	1.0.1			
	10.272	100 M		
			C	

		None	14	8	0.96	2.59	0.85	1.39	1.79	0.71	1.66	<mark>1.35</mark>	0.76
		SNV	1	1	1.78	1.40	0.49	1.86	1.34	0.44	2.21	<mark>1.31</mark>	0.60
	R	SD	7	4	1.01	2.47	0.84	1.27	1.96	0.74	1.64	<mark>1.35</mark>	0.67
		SD+LD	3	3	1.08	2.30	0.81	1.27	1.97	0.74	1.55	<mark>1.42</mark>	0.73
CD.		SNV+SD	1	1	1.34	1.87	0.71	1.43	1.74	0.67	1.84	<mark>1.28</mark>	0.66
SK		None	11	8	0.95	2.64	0.86	1.29	1.94	0.74	1.44	<mark>1.59</mark>	0.75
		SNV	15	6	1.15	2.16	0.79	1.50	1.67	0.64	1.36	<mark>1.63</mark>	0.74
	Log(1/R)	SD	1	1	1.26	1.97	0.74	1.34	1.87	0.71	2.00	<mark>1.10</mark>	0.48
		SD+LD	5	3	1.09	2.28	0.81	1.26	1.98	0.75	1.96	<mark>1.23</mark>	0.56
		SNV+SD	7	5	1.00	2.48	0.84	1.30	1.92	0.73	1.95	<mark>1.64</mark>	0.76
		None	6	3	1.05	1.94	0.73	1.18	1.72	0.67	2.07	<mark>1.10</mark>	0.57
		SNV	8	6	0.93	2.69	0.86	1.16	2.16	0.79	1.72	<mark>1.30</mark>	0.79
	R	SD	5	3	0.82	3.06	0.89	0.94	2.66	0.86	1.84	<mark>1.33</mark>	0.70
		SD+LD	10	5	0.74	3.38	0.91	0.89	2.81	0.87	2.11	<mark>1.25</mark>	0.68
		SNV+SD	7	3	0.90	2.77	0.87	1.02	2.44	0.83	1.52	<mark>1.47</mark>	0.72
EIVICVS		None	43	8	0.85	2.93	0.88	1.18	2.12	0.79	1.54	<mark>1.43</mark>	0.71
		SNV	9	3	1.02	2.45	0.83	1.13	2.20	0.79	1.64	<mark>1.44</mark>	0.71
	Log(1/R)	SD	10	7	0.69	3.63	0.92	0.89	2.82	0.87	1.42	<mark>1.58</mark>	0.73
		SD+LD	8	5	0.70	3.54	0.92	0.83	3.01	0.89	1.56	<mark>1.48</mark>	0.72
		SNV+SD	6	5	0.88	2.83	0.88	1.03	2.42	0.83	1.17	<mark>1.88</mark>	0.83

SR, selectivity ratio: VIP, variable importance projection: EMCVS, ensemble Monte Carlo variable selection; SD, second derivative; SNV, standard normal variate; LD, linear detrend; #Bands, wavelengths used for model development; #LVs, latent variables. The overall best model for 4 °C is highlighted in bold.

Chemometric	Spectral	#	#	C	alibration		Cros	s validati	on	Prediction			
Chen m	nometric ethod	Pre- treatment	Bands	LV	RMSEc	RPDc	R ² c	RMSEcv	RPDcv	R ² cv	RMSEp	RPDp	R ² p
		None	106	8	0.37	5.84	0.97	0.64	3.38	0.92	1.38	<mark>1.66</mark>	0.75
		SNV	106	7	0.39	5.59	0.97	0.63	3.45	0.92	1.39	<mark>1.56</mark>	0.74
	R	SD	100	6	0.66	3.27	0.91	1.00	2.17	0.79	1.18	<mark>1.67</mark>	0.75
		SD+LD	100	5	0.69	3.13	0.90	1.02	2.12	0.78	1.15	<mark>1.72</mark>	0.76
חוכ		SNV+SD	100	4	0.60	3.59	0.92	0.81	2.68	0.86	0.80	<mark>2.63</mark>	0.89
PLS		None	106	8	0.42	5.14	0.96	0.71	3.05	0.89	1.14	<mark>2.48</mark>	0.88
		SNV	106	7	0.42	5.13	0.96	0.69	3.15	0.90	1.02	<mark>2.32</mark>	0.86
	Log(1/R)	SD	100	4	0.64	3.41	0.91	0.84	2.58	0.85	1.21	<mark>1.84</mark>	0.80
		SD+LD	100	5	0.53	4.11	0.94	0.74	2.92	0.88	1.16	<mark>2.02</mark>	0.82
		SNV+SD	100	5	0.54	4.05	0.94	0.73	2.98	0.89	1.01	<mark>2.31</mark>	0.87
		None	20	8	0.48	4.49	0.95	0.73	2.97	0.89	1.06	<mark>1.99</mark>	0.84
		SNV	37	8	0.41	5.29	0.96	0.63	3.42	0.91	1.58	<mark>1.27</mark>	0.72
	R	SD	33	5	0.76	2.86	0.88	1.03	2.10	0.78	1.09	<mark>1.77</mark>	0.81
		SD+LD	27	6	0.72	3.03	0.89	1.05	2.06	0.76	0.91	<mark>1.97</mark>	0.82
		SNV+SD	9	4	0.65	3.34	0.91	0.80	2.72	0.87	0.92	<mark>2.27</mark>	0.87
VIP		None	15	6	0.52	4.13	0.94	0.72	3.03	0.89	1.10	<mark>2.42</mark>	0.88
		SNV	33	7	0.53	4.11	0.94	0.79	2.75	0.87	0.83	<mark>3.10</mark>	0.93
	Log(1/R)	SD	10	5	0.63	3.44	0.92	0.80	2.70	0.86	1.12	<mark>1.94</mark>	0.83
		SD+LD	13	4	0.66	3.29	0.91	0.80	2.71	0.86	1.34	<mark>1.59</mark>	0.78
		SNV+SD	10	5	0.60	3.59	0.92	0.78	2.78	0.87	1.04	<mark>2.16</mark>	0.87
SR	R	None	8	5	1.07	2.02	0.75	1.30	1.66	0.65	0.82	2.67	0.93

Appendix 2. Performance of the TVC PLS-R models developed using VIS-SWNIR (445 - 970 nm) data from beef *LD* samples stored at 10 °C. PLS full spectral range on reflectance (R) and logarithmic transformed ($\log(1/R)$) spectral data is compared with spectral pre-treatments (SNV, SD, SD+LD, SNV+SD) and band selection methods (VIP, SR and EMCVS).

lournal Pre-proof

		SNV	21	9	0.55	3.97	0.94	0.87	2.50	0.84	0.95	<mark>2.20</mark>	0.86
		SD	13	6	0.67	3.25	0.91	0.92	2.35	0.82	1.00	<mark>2.07</mark>	0.87
		SD+LD	17	6	0.74	2.94	0.88	1.04	2.08	0.77	0.88	<mark>2.37</mark>	0.89
		SNV+SD	20	7	0.60	3.59	0.92	0.88	2.46	0.84	0.91	<mark>2.09</mark>	0.86
		None	8	5	0.90	2.39	0.83	1.12	1.93	0.73	1.07	<mark>2.12</mark>	0.86
		SNV	16	7	0.52	4.16	0.94	0.82	2.64	0.86	0.81	<mark>3.16</mark>	0.93
	Log(1/R)	SD	9	7	0.54	4.05	0.94	0.73	2.96	0.89	1.38	<mark>1.81</mark>	0.79
		SD+LD	17	4	0.67	3.25	0.91	0.85	2.55	0.85	1.34	<mark>1.64</mark>	0.79
		SNV+SD	17	4	0.65	3.32	0.91	0.86	2.51	0.84	1.15	<mark>1.98</mark>	0.85
		None	24	5	0.44	4.94	0.96	0.55	3.91	0.93	1.51	<mark>1.47</mark>	0.77
		SNV	18	4	0.45	4.85	0.96	0.56	3.85	0.93	1.73	<mark>1.19</mark>	0.73
	R	SD	13	3	0.75	2.90	0.88	0.87	2.49	0.84	1.00	<mark>1.78</mark>	0.80
		SD+LD	7	5	0.63	3.43	0.91	0.77	2.80	0.87	1.15	<mark>1.60</mark>	0.75
		SNV+SD	6	3	0.63	3.44	0.92	0.72	3.01	0.89	1.02	<mark>1.84</mark>	0.79
EIVICVS		None	9	5	0.44	4.93	0.96	0.55	3.93	0.94	1.03	<mark>2.30</mark>	0.87
		SNV	7	6	0.48	4.51	0.95	0.65	3.34	0.91	1.00	<mark>2.40</mark>	0.88
	Log(1/R)	SD	10	3	0.66	3.29	0.91	0.75	2.89	0.88	1.04	<mark>1.96</mark>	0.83
		SD+LD	11	3	0.67	3.23	0.90	0.77	2.83	0.88	1.12	<mark>1.87</mark>	0.82
		SNV+SD	9	4	0.53	4.08	0.94	0.65	3.36	0.91	0.84	<mark>2.50</mark>	0.89

SR, selectivity ratio: VIP, variable importance projection: EMCVS, ensemble Monte Carlo variable selection; SD, second derivative; SNV, standard normal variate; LD, linear detrend; #Bands, wavelengths used for model development; #LVs, latent variables. The overall best model for 10 °C is highlighted in bold.

Char	Chemometric	Spectral	#	#	Calib	ration		Cross va	lidation		Prediction			
Chen m	ethod	Pre- treatment	Bands	LV	RMSEc	RPDc	R ² c	RMSEcv	RPDcv	R ² cv	RMSEp	RPDp	R ² p	
		None	106	6	0.95	2.46	0.83	1.11	2.11	0.78	1.40	<mark>1.52</mark>	0.75	
		SNV	106	6	0.85	2.76	0.87	1.01	2.32	0.81	1.33	<mark>1.47</mark>	0.78	
	R	SD	100	5	0.96	2.43	0.83	1.19	1.96	0.74	1.11	<mark>1.76</mark>	0.79	
		SNV+SD	100	7	0.86	2.72	0.87	1.09	2.15	0.78	0.98	<mark>1.99</mark>	0.84	
ыс		SD+LD	100	6	0.92	2.53	0.84	1.15	2.04	0.76	1.22	<mark>1.61</mark>	0.76	
PL3		None	106	6	0.92	2.54	0.85	1.17	2.00	0.76	1.23	<mark>1.71</mark>	0.77	
		SNV	106	6	0.83	2.83	0.87	0.98	2.39	0.83	1.36	<mark>1.44</mark>	0.73	
	Log(1/R)	SD	100	5	0.90	2.59	0.85	1.11	2.11	0.78	1.11	<mark>1.80</mark>	0.79	
		SNV+SD	100	4	0.94	2.50	0.84	1.09	2.14	0.78	1.11	<mark>1.76</mark>	0.79	
		SD+LD	100	4	0.94	2.49	0.84	1.13	2.07	0.77	1.16	<mark>1.71</mark>	0.77	
		None	8	5	1.03	2.28	0.81	1.13	2.06	0.77	1.51	<mark>1.42</mark>	0.69	
		SNV	22	7	0.86	2.73	0.87	0.96	2.43	0.83	1.55	<mark>1.28</mark>	0.74	
	R	SD	7	4	1.19	1.96	0.74	1.29	1.82	0.70	1.08	<mark>1.85</mark>	0.80	
		SNV+SD	8	3	1.21	1.93	0.73	1.32	1.77	0.68	1.31	<mark>1.63</mark>	0.74	
		SD+LD	19	6	1.12	2.09	0.77	1.26	1.85	0.71	1.32	<mark>1.51</mark>	0.72	
VIP		None	8	3	1.04	2.26	0.80	1.09	2.14	0.78	1.51	<mark>1.37</mark>	0.65	
		SNV	10	6	0.92	2.55	0.85	1.03	2.28	0.81	1.24	<mark>1.68</mark>	0.77	
	Log(1/R)	SD	30	4	0.99	2.37	0.82	1.12	2.08	0.77	1.15	<mark>1.76</mark>	0.78	
		SNV+SD	24	5	0.95	2.46	0.83	1.10	2.12	0.78	1.06	<mark>1.88</mark>	0.80	
		SD+LD	15	5	1.03	2.27	0.81	1.15	2.03	0.76	1.28	<mark>1.57</mark>	0.72	
CD	P	None	7	5	1.41	1.66	0.64	1.59	1.47	0.54	1.44	<mark>1.60</mark>	0.74	
лс	n	SNV	14	8	1.11	2.11	0.78	1.24	1.89	0.72	1.27	<mark>1.74</mark>	0.81	

Appendix 3. Performance of the TVC PLS-R models developed using VIS-SWNIR (445 - 970 nm) data from beef *LD* samples stored at 4 °C or 10 °C. PLS full spectral range on reflectance (R) and logarithmic transformed (log(1/R)) spectral data is compared with spectral pre-treatments (SNV, SD, SD+LD, SNV+SD) and band selection methods (VIP, SR and EMCVS).

ourn	$\mathbf{D}_{\mathbf{f}}$	\mathbf{nr}		
		$\mathbf{p}_{\mathbf{L}}$	U	

		SD	16	6	1.08	2.16	0.79	1.24	1.89	0.72	1.17	<mark>1.70</mark>	0.78
		SNV+SD	11	6	1.10	2.13	0.78	1.25	1.87	0.72	1.25	<mark>1.58</mark>	0.74
		SD+LD	17	6	1.02	2.29	0.81	1.20	1.96	0.74	1.24	<mark>1.60</mark>	0.75
		None	14	7	1.02	2.30	0.81	1.23	1.90	0.73	1.27	<mark>1.63</mark>	0.74
		SNV	17	5	1.02	2.29	0.81	1.22	1.92	0.73	1.24	<mark>1.61</mark>	0.75
	Log(1/R)	SD	13	7	0.97	2.40	0.83	1.15	2.03	0.76	1.23	<mark>1.70</mark>	0.76
		SNV+SD	17	5	1.07	2.18	0.79	1.24	1.89	0.72	1.41	<mark>1.43</mark>	0.68
		SD+LD	14	6	0.99	2.37	0.82	1.16	2.02	0.76	1.11	<mark>1.81</mark>	0.79
		None	3	2	1.02	1.65	0.63	1.07	1.57	0.59	1.66	<mark>1.29</mark>	0.60
		SNV	9	6	0.81	2.90	0.88	0.90	2.61	0.85	1.56	<mark>1.27</mark>	0.74
	R	SD	3	3	1.10	2.13	0.78	1.17	2.01	0.75	1.01	<mark>1.97</mark>	0.83
		SNV+SD	8	4	0.94	2.48	0.84	1.05	2.22	0.80	0.95	<mark>2.10</mark>	0.85
ENACVS		SD+LD	4	3	0.92	2.55	0.85	0.96	2.43	0.83	1.24	<mark>1.61</mark>	0.76
EIVICVS		None	23	13	0.61	3.82	0.93	0.77	3.04	0.89	0.88	<mark>2.25</mark>	0.86
		SNV	15	6	0.91	2.57	0.85	1.01	2.32	0.81	1.20	<mark>1.71</mark>	0.78
	Log(1/R)	SD	8	6	0.93	2.50	0.84	1.05	2.22	0.80	1.21	<mark>1.69</mark>	0.76
		SNV+SD	8	3	0.95	2.45	0.83	1.01	2.31	0.81	0.96	<mark>2.10</mark>	0.84
		SD+LD	9	3	0.95	2.45	0.83	1.03	2.27	0.81	1.04	<mark>1.92</mark>	0.81

SR, selectivity ratio: VIP, variable importance projection: EMCVS, ensemble Monte Carlo variable selection; SD, second derivative; SNV, standard normal variate; LD, linear detrend; #Bands, wavelengths used for model development; #LVs, latent variables. The overall best model for 10 °C is highlighted in bold.

Appendix 4. Performance of the TVC PLS-R models developed using NIR (957 - 1664 nm) data from beef LD samples stored at 4 °C, PLS full
spectral range on reflectance (R) and logarithmic transformed (log(1/R)) spectral data is compared with spectral pre-treatments (SNV, SD,
SD+LD, SNV+SD) and band selection methods (VIP, SR and EMCVS).

Chor	Chemometric	Spectral	#	#	С	alibration		Cros	s validatio	n	Prediction			
m	ethod	Pre- treatment	Bands	LV	RMSEc	RPDc	R ² c	RMSEcv	RPDcv	R ² cv	RMSEp	RPDp	R ² p	
		None	102	8	0.66	3.58	0.92	1.14	2.07	0.78	2.75	<mark>0.94</mark>	0.45	
	Р	SD	96	5	0.84	2.79	0.87	1.17	2.00	0.75	1.97	<mark>1.68</mark>	0.84	
	ĸ	SD+LD	96	5	0.84	2.79	0.87	1.15	2.04	0.76	1.91	<mark>1.72</mark>	0.84	
		SNV+SD	96	5	0.87	2.70	0.86	1.19	1.97	0.74	1.66	<mark>1.69</mark>	0.83	
PL5		None	102	6	0.82	2.87	0.88	1.21	1.94	0.75	3.50	<mark>0.69</mark>	0.08	
	l = -(1/D)	SD	96	5	0.92	2.55	0.85	1.45	1.62	0.63	0.87	<mark>2.23</mark>	0.91	
	LOG(1/K)	SD+LD	96	5	0.93	2.52	0.84	1.43	1.64	0.64	0.83	<mark>2.33</mark>	0.91	
		SNV+SD	96	4	0.72	3.28	0.91	0.97	2.42	0.83	1.19	<mark>1.63</mark>	0.82	
	R	None	12	7	0.66	3.55	0.92	1.03	2.28	0.82	2.89	<mark>1.33</mark>	0.61	
		SD	17	5	0.85	2.76	0.87	1.14	2.06	0.77	2.74	<mark>1.23</mark>	0.57	
		SD+LD	17	5	0.93	2.52	0.84	1.20	1.95	0.74	1.91	<mark>1.59</mark>	0.72	
		SNV+SD	12	7	0.83	2.82	0.87	1.08	2.17	0.79	1.94	<mark>1.45</mark>	0.68	
VIP		None	14	7	0.60	3.93	0.94	0.86	2.72	0.87	2.88	<mark>0.96</mark>	0.27	
		SD	11	6	0.86	2.72	0.86	1.11	2.11	0.78	1.58	<mark>2.25</mark>	0.86	
	Log(1/R)	SD+LD	8	6	0.88	2.68	0.86	1.08	2.17	0.79	1.22	<mark>2.07</mark>	0.86	
		SNV+SD	7	4	0.76	3.07	0.89	0.94	2.50	0.84	1.24	<mark>1.87</mark>	0.80	
		None	42	6	1.02	2.29	0.81	1.32	1.78	0.69	1.57	<mark>1.78</mark>	0.86	
	P	SD	4	3	1.21	1.94	0.73	1.38	1.70	0.66	1.22	<mark>1.91</mark>	0.82	
CP	ĸ	SD+LD	2	1	1.57	1.49	0.55	1.64	1.43	0.51	1.73	<mark>1.37</mark>	0.64	
SR		SNV+SD	1	1	1.51	1.55	0.58	1.59	1.48	0.54	2.28	<mark>1.01</mark>	0.32	
		None	29	6	1.04	2.26	0.80	1.40	1.68	0.66	1.12	<mark>2.26</mark>	0.87	
	LO2(1/ N)	SD	6	2	1.44	1.63	0.62	1.58	1.49	0.55	1.23	<mark>2.00</mark>	0.90	

		SD+LD	4	4	1.35	1.74	0.67	1.60	1.47	0.54	1.13	<mark>2.12</mark>	0.91
		SNV+SD	1	1	1.67	1.40	0.49	1.75	1.34	0.44	2.56	<mark>0.90</mark>	0.15
		None	8	5	0.72	3.24	0.90	1.00	2.34	0.82	3.91	<mark>0.82</mark>	0.07
	D	SD	16	4	0.94	2.50	0.84	1.14	2.06	0.76	2.83	<mark>1.30</mark>	0.63
	n	SD+LD	6	4	0.93	2.52	0.84	1.10	2.14	0.78	1.98	<mark>1.59</mark>	0.73
	Log(1/R)	SNV+SD	8	6	0.65	3.63	0.92	0.79	2.95	0.89	1.15	<mark>2.18</mark>	0.85
EIVICVS		None	19	5	0.86	2.69	0.86	1.06	2.20	0.79	3.85	<mark>0.75</mark>	0.01
		SD	10	4	0.80	2.93	0.88	1.01	2.32	0.82	0.93	<mark>2.48</mark>	0.89
		SD+LD	13	4	0.78	3.03	0.89	1.01	2.31	0.81	0.92	<mark>3.00</mark>	0.92
		SNV+SD	5	4	0.73	3.22	0.90	0.88	2.68	0.86	1.51	<mark>1.54</mark>	0.70

SR, selectivity ratio: VIP, variable importance projection: EMCVS, ensemble Monte Carlo variable selection; SD, second derivative; SNV, standard normal variate; LD, linear detrend; #Bands, wavelengths used for model development; #LVs, latent variables. The overall best model for 4 °C is highlighted in bold.

Journal

Appendix 5. Performance of the TVC PLS-R models developed using NIR (957 - 1664 nm) data from beef *LD* samples stored at 10 °C, PLS full spectral range on reflectance (R) and logarithmic transformed ($\log(1/R)$) spectral data is compared with spectral pre-treatments (SNV, SD, SD+LD, SNV+SD) and band selection methods (VIP, SR and EMCVS).

Cham		Due	#	#	Cal	ibration		Cross	s validatio	n	Pre	diction	
me	ethod	treatment	Bands	LV	RMSEc	RPDc	R ² c	RMSEcv	RPDcv	R ² cv	RMSEp	RPDp	R ² p
		None	102	11	0.59	3.68	0.93	1.06	2.04	0.77	1.90	<mark>1.16</mark>	0.62
		SNV	102	5	1.03	2.10	0.77	1.39	1.55	0.60	2.19	<mark>1.23</mark>	0.54
	R	SD	96	7	0.71	3.07	0.89	1.21	1.79	0.70	2.05	<mark>0.98</mark>	0.47
		SD+LD	96	4	1.27	1.71	0.66	1.52	1.43	0.51	1.67	<mark>1.28</mark>	0.55
PLS		SNV+SD	96	6	0.73	2.97	0.89	1.07	2.02	0.76	1.59	<mark>1.31</mark>	0.63
PLS		None	102	11	0.55	3.91	0.93	1.07	2.02	0.76	2.38	<mark>0.80</mark>	0.35
		SNV	102	12	0.41	5.29	0.96	0.90	2.41	0.83	1.67	<mark>1.53</mark>	0.73
	Log(1/R)	SD	96	7	0.68	3.18	0.90	1.11	1.95	0.74	1.38	<mark>1.33</mark>	0.64
		SD+LD	96	6	0.79	2.75	0.87	1.29	1.68	0.65	1.37	<mark>1.31</mark>	0.63
		SNV+SD	96	7	0.61	3.56	0.92	1.03	2.12	0.78	1.19	<mark>1.57</mark>	0.72
		None	19	9	0.68	3.19	0.90	1.00	2.17	0.79	1.79	<mark>1.02</mark>	0.51
		SNV	4	4	1.00	2.10	0.77	1.17	1.79	0.69	1.93	<mark>1.05</mark>	0.41
	R	SD	8	4	0.90	2.40	0.83	1.04	2.08	0.77	1.59	<mark>1.34</mark>	0.64
		SD+LD	7	4	0.81	2.67	0.86	0.96	2.26	0.81	1.37	<mark>1.31</mark>	0.65
EMCVS		SNV+SD	14	3	0.80	2.72	0.86	0.90	2.39	0.83	1.64	<mark>1.43</mark>	0.69
LIVICVJ		None	12	8	0.63	3.46	0.92	0.88	2.45	0.84	2.57	<mark>0.70</mark>	0.31
		SNV	19	9	0.44	4.93	0.96	0.67	3.22	0.90	1.15	<mark>2.23</mark>	0.87
	Log(1/R)	SD	12	5	0.78	2.78	0.87	0.96	2.25	0.80	1.71	<mark>1.35</mark>	0.67
		SD+LD	9	4	1.03	2.11	0.78	1.14	1.91	0.73	1.49	<mark>1.15</mark>	0.51
		SNV+SD	15	5	0.71	3.05	0.89	0.92	2.36	0.82	1.45	<mark>1.30</mark>	0.60
SR		None	1	1	2.14	1.01	0.03	2.32	0.93	0.12	1.93	<mark>0.90</mark>	0.29
	ĸ	SNV	7	5	1.28	1.69	0.65	1.58	1.37	0.49	2.22	<mark>0.84</mark>	0.26

	11 12 24 24	105	Sec. 15 (a)	6	
			1 Mar 10		

		SD	2	1	1.45	1.49	0.55	1.55	1.40	0.49	1.80	<mark>0.96</mark>	0.30
		SD+LD	1	1	1.72	1.26	0.37	1.84	1.18	0.29	2.26	<mark>0.77</mark>	0.12
		SNV+SD	1	1	1.91	1.14	0.23	2.03	1.07	0.14	1.90	<mark>1.04</mark>	0.36
		None	1	1	2.14	1.01	0.02	2.32	0.93	0.15	1.95	<mark>0.89</mark>	0.26
		SNV	1	1	1.92	1.13	0.21	2.04	1.06	0.13	1.50	<mark>1.23</mark>	0.72
	Log(1/R)	SD	1	1	1.91	1.13	0.22	2.00	1.08	0.15	1.74	<mark>0.99</mark>	0.29
		SD+LD	1	1	1.91	1.14	0.23	2.00	1.09	0.16	1.73	<mark>1.00</mark>	0.30
		SNV+SD	4	3	1.49	1.46	0.53	1.78	1.22	0.36	1.82	<mark>1.07</mark>	0.39
		None	6	4	1.32	1.64	0.63	1.55	1.40	0.49	2.17	<mark>0.89</mark>	0.28
	R	SNV	21	8	0.74	2.92	0.88	1.14	1.90	0.74	1.95	<mark>1.30</mark>	0.60
		SD	6	4	1.07	2.02	0.76	1.22	1.78	0.69	1.47	<mark>1.64</mark>	0.76
		SD+LD	5	3	1.40	1.55	0.58	1.59	1.36	0.47	1.91	<mark>1.14</mark>	0.54
		SNV+SD	9	4	0.87	2.48	0.84	1.03	2.10	0.77	1.82	<mark>1.71</mark>	0.77
VIP		None	13	9	0.82	2.65	0.86	1.18	1.84	0.71	1.78	<mark>1.79</mark>	0.86
		SNV	20	10	0.54	4.00	0.94	1.04	2.07	0.78	1.82	<mark>1.12</mark>	0.53
	Log(1/R)	SD	24	8	0.67	3.23	0.90	0.98	2.22	0.80	1.40	<mark>1.35</mark>	0.66
		SD+LD	22	6	0.90	2.42	0.83	1.18	1.84	0.70	1.61	<mark>1.15</mark>	0.54
		SNV+SD	31	8	0.67	3.23	0.90	0.98	2.21	0.80	1.14	<mark>1.65</mark>	0.75

SR, selectivity ratio: VIP, variable importance projection: EMCVS, ensemble Monte Carlo variable selection; SD, second derivative; SNV, standard normal variate; LD, linear detrend; #Bands, wavelengths used for model development; #LVs, latent variables. The overall best model for 10 °C is highlighted in bold.

Appendix 6. Performance of the TVC PLS-R models developed using NIR (957 - 1664 nm) data from beef LD samples stored either 4 °C or 10
°C. PLS full spectral range on reflectance (R) and logarithmic transformed (log(1/R)) spectral data is compared with spectral pre-treatments
(SNV, SD, SD+LD, SNV+SD) and band selection methods (VIP, SR and EMCVS).

Cha	Chemometric method	Spectral	#	#	Ca	libration		Cross va	lidatio	n	Pro	ediction	
n	nethod	Pre- treatment	Bands	LV	RMSEc	RPDc	R ² c	RMSEcv	PDcv	R ² cv	RMSEp	RPDp	R ² p
		None	102	6	1.36	1.72	0.66	1.57	1.49	0.55	2.19	<mark>0.89</mark>	0.29
		SNV	102	13	0.86	2.72	0.86	1.27	1.85	0.71	2.58	<mark>1.00</mark>	0.44
	R	SD	96	6	1.19	1.97	0.74	1.40	1.67	0.64	1.27	<mark>1.53</mark>	0.71
		SNV+SD	96	7	1.06	2.20	0.79	1.36	1.72	0.67	1.58	<mark>1.25</mark>	0.61
PLS		SD+LD	96	6	1.20	1.94	0.73	1.42	1.65	0.64	1.29	<mark>1.52</mark>	0.71
		None	102	5	1.51	1.55	0.58	1.74	1.34	0.45	2.78	<mark>0.72</mark>	0.14
		SNV	102	13	0.84	2.79	0.87	1.38	1.70	0.67	1.44	<mark>1.35</mark>	0.64
	Log(1/R)	SD	96	6	1.18	1.98	0.74	1.47	1.59	0.61	1.21	<mark>1.62</mark>	0.74
		SNV+SD	96	6	1.15	2.03	0.76	1.48	1.58	0.61	1.29	<mark>1.51</mark>	0.70
		SD+LD	96	6	1.17	2.01	0.75	1.47	1.59	0.61	1.18	<mark>1.66</mark>	0.76
VIP	R	None	9	5	1.41	1.65	0.63	1.56	1.50	0.56	2.52	<mark>0.79</mark>	0.22
		SNV	20	13	0.92	2.53	0.84	1.19	1.97	0.75	1.45	<mark>1.42</mark>	0.68
		SD	27	6	1.18	1.99	0.75	1.37	1.70	0.66	1.41	<mark>1.40</mark>	0.65
		SNV+SD	17	6	1.17	2.01	0.75	1.35	1.73	0.67	1.41	<mark>1.41</mark>	0.65
		SD+LD	6	4	1.36	1.72	0.66	1.47	1.59	0.60	1.34	<mark>1.46</mark>	0.67
	Log(1/R)	None	32	10	1.24	1.89	0.72	1.52	1.54	0.59	2.33	<mark>0.84</mark>	0.27
		SNV	10	5	1.35	1.73	0.67	1.52	1.53	0.58	1.93	<mark>1.09</mark>	0.44
		SD	17	6	1.23	1.90	0.72	1.41	1.66	0.64	1.29	<mark>1.58</mark>	0.72
		SNV+SD	12	6	1.21	1.93	0.73	1.40	1.67	0.65	1.17	<mark>1.66</mark>	0.75
		SD+LD	17	6	1.27	1.83	0.70	1.43	1.63	0.63	1.30	<mark>1.51</mark>	0.70
SR	R	None	28	7	1.40	1.67	0.64	1.93	1.21	0.41	1.31	<mark>1.49</mark>	0.70
SR	N	SNV	1	1	1.92	1.22	0.33	1.98	1.18	0.28	2.19	<mark>0.89</mark>	0.13

Journal Pre-proof

		SD	1	1	1.77	1.32	0.43	1.81	1.29	0.40	1.88	<mark>1.05</mark>	0.39
		SNV+SD	5	4	1.49	1.57	0.59	1.62	1.44	0.52	1.58	<mark>1.24</mark>	0.55
		SD+LD	1	1	1.72	1.36	0.46	1.77	1.32	0.43	1.93	<mark>1.02</mark>	0.37
		None	27	7	1.44	1.63	0.62	1.84	1.27	0.42	1.48	<mark>1.33</mark>	0.63
		SNV	11	3	1.91	1.23	0.34	2.07	1.13	0.23	2.32	<mark>0.86</mark>	0.11
	Log(1/R)	SD	1	1	1.94	1.20	0.31	1.98	1.18	0.28	1.84	<mark>1.13</mark>	0.45
		SNV+SD	1	1	2.10	1.12	0.20	2.16	1.08	0.15	2.14	<mark>0.92</mark>	0.18
		SD+LD	1	1	1.93	1.21	0.32	1.97	1.19	0.29	1.83	<mark>1.14</mark>	0.46
		None	6	4	1.42	1.65	0.63	1.55	1.52	0.57	2.12	<mark>1.00</mark>	0.37
		SNV	19	11	0.85	2.76	0.87	1.09	2.15	0.79	1.77	<mark>1.23</mark>	0.56
	R	SD	10	4	1.31	1.78	0.69	1.42	1.65	0.63	1.48	<mark>1.45</mark>	0.67
		SNV+SD	12	5	1.04	2.24	0.80	1.18	1.98	0.74	1.15	<mark>1.75</mark>	0.77
		SD+LD	11	4	1.32	1.77	0.68	1.43	1.63	0.63	1.55	<mark>1.28</mark>	0.59
EIVICVS		None	12	6	1.36	1.70	0.65	1.55	1.49	0.55	2.08	<mark>0.95</mark>	0.32
		SNV	19	11	0.90	2.60	0.85	1.14	2.05	0.77	1.03	<mark>1.92</mark>	0.81
	Log(1/R)	SD	5	4	1.29	1.81	0.70	1.41	1.66	0.64	1.46	<mark>1.35</mark>	0.63
		SNV+SD	5	3	1.17	1.96	0.74	1.28	1.80	0.69	1.63	<mark>1.20</mark>	0.53
		SD+LD	6	4	1.25	1.87	0.72	1.37	1.71	0.66	1.48	<mark>1.35</mark>	0.64

Appendix 7. Performance of the best TVC PLS-R models developed using the LL data fusion of VIS-SWNIR (445 - 970 nm) and NIR (957 -
1664 nm) HSI data from beef LD samples stored at 4 °C. PLS full spectral range on reflectance (R) and logarithmic transformed (log(1/R))
spectral data is compared with spectral pre-treatments (SNV, SD, SD+LD, SNV+SD) and band selection methods (VIP, SR and EMCVS).

Ch		Spectral	#	#	Ca	libration		Cros	s validati	on	Pre	ediction	
	method	Pre- treatment	Bands	LV	RMSEcv	RPDc	R ² c	RMSEcv	RPDcv	R ² cv	RMSEp	RPDp	R ² p
		None	208	7	0.87	2.85	0.88	1.53	1.63	0.66	1.73	<mark>1.23</mark>	0.68
		SNV	208	7	0.81	3.06	0.89	1.34	1.86	0.73	1.69	<mark>1.30</mark>	0.67
	R	SD	202	6	0.84	2.97	0.89	1.35	1.86	0.71	1.18	<mark>1.75</mark>	0.86
		SD+LD	202	6	0.85	2.94	0.88	1.38	1.81	0.70	1.17	<mark>1.76</mark>	0.86
DIC		SNV+SD	202	6	0.82	3.05	0.89	1.15	2.17	0.79	1.35	<mark>1.52</mark>	0.77
PLS		None	198	6	0.75	3.30	0.91	1.45	1.74	0.67	1.42	<mark>1.51</mark>	0.78
		SNV	208	6	0.75	3.32	0.91	1.06	2.35	0.82	1.36	<mark>1.57</mark>	0.75
	Log(1/R)	SD	208	7	0.71	3.49	0.92	1.11	2.25	0.80	1.32	<mark>1.66</mark>	0.78
		SD+LD	208	9	0.71	3.51	0.92	1.28	1.95	0.75	1.38	<mark>1.86</mark>	0.85
		SNV+SD	202	6	0.72	3.44	0.92	1.17	2.13	0.78	1.25	<mark>1.66</mark>	0.83
VIP		None	4	3	1.11	2.25	0.80	1.23	2.03	0.76	1.62	<mark>1.45</mark>	0.71
		SNV	24	5	0.97	2.56	0.85	1.26	1.97	0.74	1.74	<mark>1.27</mark>	0.63
	R	SD	22	11	0.56	4.42	0.95	1.11	2.27	0.81	1.74	<mark>1.65</mark>	0.83
		SD+LD	22	8	0.76	3.28	0.91	1.09	2.30	0.81	1.16	<mark>1.90</mark>	0.88
		SNV+SD	32	6	0.90	2.78	0.87	1.23	2.03	0.76	1.22	<mark>1.87</mark>	0.82
		None	10	7	0.88	2.82	0.87	1.31	1.91	0.75	1.48	<mark>1.52</mark>	0.72
		SNV	18	12	0.52	4.82	0.96	0.97	2.56	0.87	1.51	<mark>1.45</mark>	0.75
	Log(1/R)	SD	35	9	0.57	4.35	0.95	1.00	2.50	0.84	1.45	<mark>1.56</mark>	0.75
		SD+LD	29	12	0.50	4.99	0.96	0.95	2.64	0.86	1.58	<mark>1.61</mark>	0.74
		SNV+SD	10	7	0.86	2.90	0.88	1.32	1.90	0.72	1.34	<mark>1.71</mark>	0.77
SR	R	None	15	7	1.07	2.34	0.82	1.56	1.60	0.66	1.53	<mark>1.62</mark>	0.79

lournal Pre-proof

		SNV	17	9	0.92	2.72	0.87	1.36	1.83	0.71	1.00	<mark>2.22</mark>	0.87
		SD	8	4	1.00	2.49	0.84	1.27	1.96	0.74	1.60	<mark>1.37</mark>	0.68
		SD+LD	8	3	1.11	2.25	0.80	1.29	1.94	0.73	1.72	<mark>1.28</mark>	0.66
		SNV+SD	2	1	1.35	1.85	0.71	1.45	1.72	0.66	1.77	<mark>1.29</mark>	0.66
		None	15	8	0.91	2.73	0.87	1.39	1.80	0.73	1.52	<mark>1.48</mark>	0.74
		SNV	21	6	1.14	2.19	0.79	1.58	1.59	0.62	1.41	<mark>1.58</mark>	0.77
	Log(1/R)	SD	7	6	0.85	2.94	0.88	1.09	2.29	0.81	2.64	<mark>1.35</mark>	0.64
		SD+LD	1	1	1.24	2.01	0.75	1.31	1.90	0.72	2.08	<mark>1.05</mark>	0.44
		SNV+SD	8	5	0.91	2.72	0.87	1.13	2.21	0.80	1.97	<mark>1.78</mark>	0.83
		None	9	7	0.97	2.58	0.85	1.28	1.95	0.75	1.33	<mark>1.66</mark>	0.76
		SNV	5	4	1.02	2.32	0.81	1.15	2.06	0.76	1.74	<mark>1.29</mark>	0.58
	R	SD	9	6	0.71	3.51	0.92	0.91	2.74	0.87	1.71	<mark>1.29</mark>	0.81
		SD+LD	10	6	0.74	3.36	0.91	0.94	2.64	0.86	1.69	<mark>1.36</mark>	0.76
		SNV+SD	28	4	0.92	2.72	0.86	1.08	2.30	0.81	1.28	<mark>1.81</mark>	0.82
EMCVS		None	25	11	0.53	4.73	0.96	0.93	2.68	0.87	3.12	<mark>1.30</mark>	0.60
		SNV	11	4	0.70	3.54	0.92	0.94	2.66	0.86	1.94	<mark>1.34</mark>	0.61
	Log(1/R)	SD	6	4	0.77	3.22	0.90	0.92	2.72	0.87	1.54	<mark>1.43</mark>	0.68
		SD+LD	5	4	0.85	2.93	0.88	0.98	2.54	0.85	1.33	<mark>1.66</mark>	0.80
		SNV+SD	8	5	0.75	3.32	0.91	0.92	2.72	0.87	1.30	<mark>1.71</mark>	0.77

-

SR, selectivity ratio: VIP, variable importance projection: EMCVS, ensemble Monte Carlo variable selection; SD, second derivative; SNV, standard normal variate; LD, linear detrend; #Bands, wavelengths used for model development; #LVs, latent variables. The overall best model for the LL data fusion of 4 °C data is highlighted in bold.

Appendix 8. Performance of the best TVC PLS-R models developed using the LL data fusion of VIS-SWNIR (445 - 970 nm) and NIR (957 - 1664 nm) HSI data from beef LD samples stored at 10 °C. PLS full spectral range on reflectance (R) and logarithmic transformed (log(1/R)) spectral data is compared with spectral pre-treatments (SNV, SD, SD+LD, SNV+SD) and band selection methods (VIP, SR and EMCVS).

Chara	· · · · · · ·	Spectral	#	#	Cal	ibration		Cros	s validatio	on	P	rediction	
Cnem me	ethod	Pre- treatment	Bands	LV	RMSEcv	RPDc	R ² c	RMSEcv	RPDcv	R ² cv	RMSEp	RPDp	R ² p
		None	208	8	0.43	5.04	0.96	0.76	2.87	0.88	1.39	<mark>1.61</mark>	0.73
		SNV	208	7	0.42	5.20	0.96	0.67	3.25	0.91	1.41	<mark>1.87</mark>	0.79
	R	SD	202	7	0.58	3.72	0.93	0.95	2.28	0.81	1.11	<mark>1.86</mark>	0.79
		SD+LD	202	7	0.58	3.73	0.93	0.95	2.29	0.81	1.10	<mark>1.87</mark>	0.79
DIS		SNV+SD	202	4	0.70	3.08	0.89	0.93	2.32	0.82	0.88	<mark>2.46</mark>	0.89
FLJ		None	208	8	0.44	4.90	0.96	0.77	2.82	0.88	1.64	<mark>1.59</mark>	0.76
		SNV	208	6	0.55	3.95	0.94	0.83	2.61	0.85	1.33	<mark>1.97</mark>	0.81
	Log(1/R)	SD	202	6	0.54	4.02	0.94	0.84	2.59	0.85	1.41	<mark>1.75</mark>	0.79
		SD+LD	202	6	0.54	4.05	0.94	0.83	2.61	0.85	1.39	<mark>1.78</mark>	0.80
		SNV+SD	202	6	0.48	4.49	0.95	0.76	2.85	0.88	1.23	<mark>1.97</mark>	0.83
		None	20	6	0.43	5.09	0.96	0.59	3.68	0.93	1.26	<mark>1.89</mark>	0.81
		SNV	128	6	0.42	5.11	0.96	0.61	3.54	0.92	1.30	<mark>1.92</mark>	0.82
	R	SD	9	6	0.52	4.16	0.94	0.65	3.36	0.91	0.86	<mark>2.09</mark>	0.84
		SD+LD	10	7	0.49	4.45	0.95	0.64	3.40	0.91	0.97	<mark>1.98</mark>	0.84
		SNV+SD	12	5	0.49	4.39	0.95	0.68	3.17	0.90	1.20	<mark>1.76</mark>	0.77
ENICVS		None	11	6	0.46	4.66	0.95	0.58	3.71	0.93	0.94	<mark>3.03</mark>	0.94
		SNV	7	5	0.59	3.69	0.93	0.75	2.90	0.88	1.13	<mark>2.22</mark>	0.86
	Log(1/R)	SD	7	3	0.62	3.49	0.92	0.71	3.04	0.89	0.98	<mark>2.05</mark>	0.84
		SD+LD	20	3	0.63	3.41	0.91	0.73	2.96	0.89	1.11	1.87	0.82
		SNV+SD	9	4	0.55	3.94	0.94	0.66	3.30	0.91	1.03	2.12	0.85
	_	None	8	5	1.07	2.02	0.75	1.30	1.66	0.65	0.82	2.67	0.93
SR	R	SNV	20	8	0.65	3.32	0.91	0.98	2.21	0.80	0.89	<mark>2.56</mark>	0.91

ournal Pre-proof

		SD	13	6	0.67	3.25	0.91	0.92	2.35	0.82	1.00	<mark>2.07</mark>	0.87
		SD+LD	23	7	0.74	2.93	0.88	1.08	2.01	0.75	0.74	<mark>2.51</mark>	0.90
		SNV+SD	11	6	0.73	2.96	0.89	0.97	2.24	0.80	1.20	<mark>1.81</mark>	0.84
		None	8	5	0.90	2.39	0.83	1.12	1.93	0.73	1.07	<mark>2.12</mark>	0.86
		SNV	25	8	0.70	3.08	0.89	1.14	1.91	0.73	1.50	<mark>1.40</mark>	0.84
	Log(1/R)	SD	25	5	0.54	4.04	0.94	0.80	2.70	0.86	1.13	<mark>2.02</mark>	0.83
		SD+LD	10	8	0.53	4.11	0.94	0.80	2.71	0.86	1.20	<mark>1.86</mark>	0.80
		SNV+SD	24	5	0.55	3.95	0.94	0.77	2.82	0.87	1.02	<mark>2.20</mark>	0.86
	R	None	21	10	0.40	5.48	0.97	0.59	3.68	0.93	1.21	<mark>1.77</mark>	0.78
		SNV	27	5	0.48	4.48	0.95	0.68	3.20	0.90	1.04	<mark>2.15</mark>	0.86
		SD	20	8	0.60	3.62	0.92	0.88	2.47	0.84	1.21	<mark>1.72</mark>	0.77
		SD+LD	16	8	0.61	3.56	0.92	0.89	2.43	0.83	1.24	<mark>1.66</mark>	0.75
		SNV+SD	14	5	0.73	2.97	0.89	0.94	2.31	0.81	0.84	<mark>2.50</mark>	0.89
VIP	Log(1/R)	None	14	7	0.48	4.52	0.95	0.68	3.17	0.90	1.11	<mark>2.35</mark>	0.87
		SNV	7	5	0.61	3.57	0.92	0.75	2.88	0.88	1.10	<mark>2.42</mark>	0.88
		SD	10	5	0.63	3.44	0.92	0.80	2.70	0.86	1.12	<mark>1.94</mark>	0.83
		SD+LD	14	4	0.63	3.43	0.92	0.80	2.70	0.86	1.16	<mark>1.85</mark>	0.82
		SNV+SD	10	5	0.58	3.72	0.93	0.74	2.92	0.88	1.01	<mark>2.12</mark>	0.85

SR, selectivity ratio: VIP, variable importance projection: EMCVS, ensemble Monte Carlo variable selection; SD, second derivative; SNV, standard normal variate; LD, linear detrend; #Bands, wavelengths used for model development; #LVs, latent variables. The overall best model for the LLDF of 10 °C data is highlighted in bold.

Appendix 9. Performance of the best TVC PLS-R models developed using the LL data fusion of VIS-SWNIR (445 - 970 nm) and NIR (957 - 1664 nm) HSI data from beef *LD* samples stored at either 4 °C or 10 °C. PLS full spectral range on reflectance (R) and logarithmic transformed (log(1/R)) spectral data is compared with spectral pre-treatments (SNV, SD, SD+LD, SNV+SD) and band selection methods (VIP, SR and EMCVS).

Regression		Pre-	#	#	Cali	ibration		Cross val	lidation		Prediction			
M	odel	treatment	Bands	LV	RMSEc	RPDc	$\mathbf{R}^2 \mathbf{c}$	RMSEcv	RPDcv	R ² cv	RMSEp	RPDp	R ² p	
		None	208	8	0.86	2.72	0.86	1.06	2.21	0.8	1.38	<mark>1.43</mark>	0.72	
		SNV	208	7	0.79	2.95	0.89	1.01	2.32	0.82	1.42	<mark>1.38</mark>	0.70	
	R	SD	202	6	0.96	2.45	0.83	1.20	1.95	0.74	0.95	<mark>2.08</mark>	0.85	
		SD+LD	202	7	0.89	2.64	0.86	1.16	2.02	0.75	1.00	<mark>2.04</mark>	0.85	
		SNV+SD	202	7	0.83	2.82	0.87	1.07	2.19	0.79	1.12	<mark>1.86</mark>	0.81	
PL5		None	208	8	0.83	2.81	0.87	1.12	2.09	0.78	1.44	<mark>1.36</mark>	0.66	
		SNV	208	5	0.94	2.50	0.84	1.09	2.14	0.78	1.33	<mark>1.49</mark>	0.71	
	Log(1/R)	SD	202	6	0.88	2.66	0.86	1.10	2.13	0.78	1.17	<mark>1.71</mark>	0.77	
		SD+LD	202	6	0.89	2.63	0.86	1.11	2.11	0.78	1.15	<mark>1.76</mark>	0.78	
		SNV+SD	202	5	0.93	2.52	0.84	1.14	2.06	0.76	1.18	<mark>1.75</mark>	0.78	
		None	30	8	0.96	2.44	0.83	1.09	2.15	0.78	1.48	<mark>1.43</mark>	0.69	
		SNV	13	7	0.96	2.45	0.83	1.09	2.15	0.78	1.40	<mark>1.44</mark>	0.73	
	R	SD	15	7	1.03	2.27	0.81	1.22	1.91	0.73	1.07	<mark>1.86</mark>	0.80	
		SD+LD	11	6	1.02	2.30	0.81	1.18	1.98	0.75	1.15	<mark>1.73</mark>	0.79	
		SNV+SD	29	10	0.82	2.84	0.88	1.10	2.13	0.78	1.09	<mark>1.83</mark>	0.81	
VIP		None	10	4	1.06	2.21	0.8	1.14	2.06	0.76	1.49	<mark>1.35</mark>	0.66	
		SNV	22	6	0.92	2.55	0.85	1.04	2.24	0.8	1.31	<mark>1.53</mark>	0.73	
	Log(1/R)	SD	6	4	1.10	2.12	0.78	1.19	1.97	0.74	1.18	<mark>1.74</mark>	0.77	
		SD+LD	6	3	1.08	2.17	0.79	1.12	2.10	0.77	1.21	<mark>1.67</mark>	0.75	
		SNV+SD	10	4	1.04	2.25	0.8	1.17	2.00	0.75	1.27	<mark>1.66</mark>	0.75	
SR	R	None	7	5	1.41	1.66	0.64	1.59	1.47	0.54	1.44	<mark>1.60</mark>	0.74	

10.10

		SNV	20	9	0.97	2.42	0.83	1.33	1.75	0.68	1.24	<mark>1.75</mark>	0.78
		SD	12	8	1.05	2.23	0.80	1.23	1.91	0.73	1.06	<mark>1.87</mark>	0.82
		SD+LD	24	6	1.07	2.19	0.79	1.23	1.90	0.72	1.24	<mark>1.61</mark>	0.75
		SNV+SD	11	6	1.13	2.06	0.76	1.32	1.78	0.69	1.33	<mark>1.50</mark>	0.71
		None	14	7	1.02	2.3	0.81	1.23	1.90	0.73	1.27	<mark>1.63</mark>	0.74
		SNV	18	8	1.09	2.15	0.78	1.31	1.79	0.69	1.19	<mark>1.96</mark>	0.82
	Log(1/R)	SD	13	7	0.97	2.4	0.83	1.15	2.03	0.76	1.23	<mark>1.70</mark>	0.76
		SD+LD	19	5	0.98	2.38	0.82	1.1	2.12	0.78	1.25	<mark>1.61</mark>	0.74
		SNV+SD	15	7	0.95	2.46	0.84	1.15	2.04	0.76	1.36	<mark>1.59</mark>	0.72
		None	15	5	0.93	2.51	0.84	1.07	2.18	0.79	1.32	<mark>1.63</mark>	0.77
		SNV	52	6	0.86	2.73	0.87	0.98	2.4	0.83	1.37	<mark>1.48</mark>	0.73
	R	SD	6	3	1.06	2.21	0.80	1.13	2.06	0.76	1.15	<mark>1.73</mark>	0.79
		SD+LD	35	4	0.96	2.45	0.83	1.12	2.08	0.77	0.87	<mark>2.27</mark>	0.88
		SNV+SD	15	4	0.90	2.61	0.85	0.99	2.36	0.82	1.03	<mark>2.10</mark>	0.85
EIVICVS		None	7	5	0.97	2.40	0.83	1.06	2.21	0.8	1.58	<mark>1.37</mark>	0.65
		SNV	17	8	0.87	2.68	0.86	1.03	2.26	0.81	1.32	<mark>1.70</mark>	0.76
	Log(1/R)	SD	6	3	0.99	2.36	0.82	1.07	2.18	0.79	1.04	<mark>1.91</mark>	0.81
		SD+LD	5	5	0.93	2.51	0.84	1.01	2.31	0.81	0.98	<mark>2.09</mark>	0.84
		SNV+SD	3	2	1.04	2.18	0.79	1.08	2.10	0.77	1.24	<mark>1.60</mark>	0.73

_

SR, selectivity ratio: VIP, variable importance projection: EMCVS, ensemble Monte Carlo variable selection; SD, second derivative; SNV, standard normal variate; LD, linear detrend; #Bands, wavelengths used for model development; #LVs, latent variables. The overall best model for the LLDF of 4 and 10 °C combine data is highlighted in bold.

Journal Pre-proof

	Time			Sample i	d. /TVC		
Temperature	(hours)	S1	S2	S 3	S7	S8	S9
	0	7.1±0.4 ^{a,c}	6.9±0.2 ^{ª,c}	7.0±0.5 ^{a,b}	5.3±0.1 ^ª	5.3±0.4 ^a	3.4±0.5 <mark>ª</mark>
	24	7.1±0.3 ^{a,c}	6.9±0.4 ^{a,c}	7.2±0.6 ^{ª,b}	5.3±0.2 <mark>ª</mark>	5.1±0.4 ^a	3.3±0.0 <mark>ª</mark>
	96	8.9±0.1 ^{b,c,d,e}	$8.6\pm0.2^{b,c,d,f,g}$	8.3±0.3 ^{a,b,c}	6.6±0.4 ^b	7.1±0.7 <mark>^b</mark>	4.1±0.3 ^a
	120	8.1±0.0 ^{a,b,c,e}	7.7±0.1 ^{a,b,c,d,g}	8.1±0.6 ^{a,b,c}	8.7±0.6 <mark>°</mark>	8.3±0.2 [°]	5.6±0.0 <mark>^b</mark>
4.00	192	8.8±0.4 ^{b,c,d,e}	9.1±0.2 ^{b,c,d,e,f,g}	8.4±0.3 ^{a,b,c}	9.4±0.1 <mark>°</mark>	9.4±0.2 <mark>°</mark>	7.7±0.5 ^{c,d}
4 %	264	10.1±0.7 ^{b,d,e,f}	10.5±0.7 ^{d,e,f}	9.0±0.1 ^{b,c,d}	10.9±0.8 ^d	12.2±0.1 ^{d,e}	8.9±0.1 ^{c,d,e}
	288	9.5±0.3 ^{b,c,d,e,f}	$9.7 \pm 0.6^{b,d,e,f,g,h}$	10.2±0.3 ^{c,d,e}	12.2±0.3 ^{e,f}	12.1±0.3 ^{d,e,f}	9.2±0.2 ^{c,d,e}
	312	9.7±1.2 ^{b,c,d,e,f}	$9.0\pm1.0^{b,c,d,f,g,h}$	10.6±0.7 ^{d,e}	-	-	-
	336	-	-		13.2±0.2 ^{e,f,g}	13.5±0.7 ^{e,f}	10.5±1.3 ^{d,e}
	360	10.6±0.5 ^{d,e,f}	11.1±0.3 ^{f,h}	11.4±1.0 ^{d,e}	14.±0.3 ^{f,g}	14.1±0.1 ^{f,g}	-
	Time			Sample i	d. /TVC		
Temperature	(hours)	S4	S5	S 6	S7	S8	S9
	0	6.1±0.1 <mark>ª</mark>	6.7±0.1 <mark>ª</mark>	6.7±0.3 ^{a,b}	5.3±0.1 ^a	5.3±0.4 <mark>ª</mark>	3.4±0.5 <mark>ª</mark>
	12	6.7±0.3 ^{b,c}	7.7±0.2 ^b	$6.9 \pm 0.1 \frac{a,b}{c}$	-	-	-
	24	6.8±0.2 ^{b,c}	7.3±0.1 ^b	7.1±0.1 ^{a,b,c}	6.0±0.3 ^a	6.1±0.8 <mark>ª</mark>	3.4±0.5 ^a
	36	$6.9 \pm 4.0^{b,c,d}$	7.7±0.1 ^b	7.1±0.2 ^{a,b,c}	-	-	-
	48	7.4±0.1 ^{c,d}	7.6±0.1 <mark>^b</mark>	7.5±0.1 ^{b,c}	-	-	-
10.00	84	$9.2 \pm 0.1^{e,f,h}$	9.4±0.1 ^{c,d,e,g}	9.3±0.2 ^{d,e}	-	-	-
10 °C	96	$9.4\pm0.2^{\text{e,f,h,i}}$	9.1±0.2 ^{c,d}	9.3±0.1 ^{d,e}	8.8±0.2 ^{b,c}	9.0±0.1 ^b	7.6±0.1 ^b
	108	9.9±0.1 ^{g,h,i}	$9.8 \pm 0.2^{\frac{c,e,f,g}{c,e,f,g}}$	9.6±0.1 ^{d,e,f}	-	-	-
	120	9.6±0.2 ^{e,f,g,h,i}	9.8 ± 0.1 ^{c,e,f,g}	9.8±0.1 ^{e,f}	10.5±0.4 ^{b,c,d}	10.2±0.1 <mark>°</mark>	9.3±0.5 <mark>°</mark>
	132	10.1±0.2 ^{g,h,i}	10.0±0.1 ^{e,f,g}	9.9±0.1 ^{e,f}	-	-	-
	144	$9.8\pm0.1^{f,g,h,i}$	9.9 ± 0.1	$10.0 \pm 0.2^{\frac{e,f}{c}}$	11.6±1.4 ^{c,d}	12.2±0.2 ^d	11.1±0.2 ^d
	168	_	-	_	12.3±0.0 ^{c,d}	13.1±0.1 ^d	12.3±0.3 ^e

1 Table 1. TVC of beef *LD* samples (log CFU/g) stored at 4 and 10 °C. Average value \pm standard deviation (n = 3). Superscripts show the statistical significance between samples obtained with Tukey-Kramer test at $\alpha = 0.05$.

3 Values followed by different letters in the same column are significantly different using ANOVA and Tukey test (p < 0.05).

Table 2. Performance of the best TVC PLS-R models developed for beef *LD* samples stored at 4 °C using the full spectral range and optimum
band selection method evaluated for VIS-SWNIR (445 - 970 nm) and NIR (957 - 1664 nm) HSI data.

Regression model		Pre- treatment		#	#	Calibration			Cross v	alidation	Prediction			
		1^{st} 2^{nd}		Bands	LV	RMSEc	RMSEc RPDc R²c		RMSEcv	RMSEcv RPDcv		RMSEp	RPDp	R ² p
445 - 970	nm								Č.					
PLS	R	SNV	SD	100	6	0.79	3.15	0.9	1.34	1.86	0.74	1.35	<mark>1.61</mark>	0.81
EMCVS	$\log(1/R)$	SNV	SD	6	5	0.88	2.83	0.88	1.03	2.42	0.83	1.17	<mark>1.88</mark>	0.83
957 - 166	4 nm													
PLS	log(1/R)	SD	AsLs	96	9	0.42	5.63	0.97	1.03	2.29	0.81	0.99	<mark>2.26</mark>	0.94
EMCVS	log(1/R)	SD	AsLs	17	7	0.5	4.71	0.95	0.7	3.37	0.91	0.81	<mark>3.09</mark>	0.95

6 EMCVS, ensemble Monte Carlo variable selection; SD, second derivative; SNV, standard normal variate; AsLs, asymmetric least squares;

#Bands, wavelengths used for model development; #LVs, latent variables. The best overall model for 4 °C is highlighted in bold.

Journe

9 Table 3. Performance of the best TVC PLS-R models developed for beef *LD* samples stored at 10 °C using the full spectral range and optimum
10 band selection method evaluated for VIS-SWNIR (445 - 970 nm) and NIR (957 - 1664 nm) HSI data.

Regression model		Pre- treatment		#	#	# Calibration			Cross	Prediction				
		1^{st}	2^{nd}	Bands	LV	RMSEc	RPDc	R ² c	RMSEcv	RPDcv	R ² cv	RMSEp	RPDp	R ² p
445 - 970	nm								¢.,					
PLS	R	SD	MS	100	6	0.48	4.52	0.95	0.7	3.08	0.89	1.09	<mark>2.84</mark>	0.92
EMCVS	R	SD	MS	46	6	0.47	4.60	0.95	0.69	3.16	0.90	0.96	<mark>3.32</mark>	0.94
957 - 1664	4 nm													
PLS	$\log(1/R)$	SD	SNV	96	7	0.62	3.49	0.92	1.03	2.11	0.78	1.2	<mark>1.59</mark>	0.72
EMCVS	$\log(1/R)$	SNV		19	9	0.44	4.93	0.96	0.67	3.22	0.90	1.15	<mark>2.23</mark>	0.87

11 EMCVS, ensemble Monte Carlo variable selection; MS, median scaled; SD, second derivative; SNV, standard normal variate; #Bands,

12 wavelengths used for model development; #LVs, latent variables. The best overall model for 10 °C is highlighted in bold.

OUTRIC

Table 4. Performance of the best TVC PLS-R models developed for beef LD samples stored at either 4 °C or 10 °C using the full spectral range 14 and optimum band selection method evaluated for VIS-SWNIR (445 - 970 nm) and NIR (957 - 1664 nm) HSI data. 15

Regression model		Pre- treatment		#	#	Calibration			Cross	validatior	l	Prec	Prediction		
		1^{st}	2^{nd}	Bands	LV	RMSEc	RPDc	R ² c	RMSEcv	RPDcv	R ² cv	RMSEp	RPDp	R ² p	
445 - 970 m	m								6.						
PLS	R	SNV	SD	100	7	0.86	2.72	0.87	1.09	2.15	0.78	0.98	<mark>1.99</mark>	0.84	
EMCVS	R	<mark>SNV</mark>	<mark>SD</mark>	<mark>8</mark>	<mark>4</mark>	<mark>0.94</mark>	<mark>2.48</mark>	<mark>0.84</mark>	<mark>1.05</mark>	<mark>2.22</mark>	<mark>0.80</mark>	<mark>0.95</mark>	<mark>2.10</mark>	<mark>0.85</mark>	
957 - 1664 i	nm														
PLS	log(1/R)	SD	LD	202	4	1.17	2.01	0.75	1.47	1.59	0.61	1.18	<mark>1.66</mark>	0.76	
EMCVS	R	SNV	SD	96	6	1.04	2.24	0.80	1.18	1.98	0.74	1.15	<mark>1.75</mark>	0.77	

EMCVS, ensemble Monte Carlo variable selection; SNV, standard normal variate; LD, linear detrend; SD, second derivative; #Bands, 16

wavelengths used for model development; #LVs, latent variables. The overall best model for the combined data of both temperatures is 17 18 highlighted in bold. OUTRI

Table 5. Performance of the best TVC PLS-R models developed using VIS-SWNIR (445 - 970 nm) and NIR (957 - 1664 nm) HSI data from

21 beef LD samples stored at (i) 4 °C, (ii) 10 °C and (iii) either 4 °C or 10 °C using the optimum band selection method and LL, ML and HL data

22 fusion

Reg	ression mo	del	Pre-tr	eatment	#	#	Ca	libration	ı	Cross	s validatio	on	Prediction		
			1 st	2 nd	Bands	LV	RMSEc	RPDc	R ² c	RMSEcv	RPDcv	R ² cv	RMSEp	RPDp	R ² p
4	ŀ°C								<u>k</u>						
VIS-SWNIR	EMCVS	log(1/R)	SNV	SD	6	5	0.88	2.83	0.88	1.03	2.42	0.83	1.17	<mark>1.88</mark>	0.83
NIR	EMCVS	log(1/R)	SD	AsLs	17	7	0.50	4.71	0.95	0.70	3.37	0.91	0.81	<mark>3.09</mark>	0.95
LL	VIP	R	SD	LD	22	8	0.76	3.28	0.91	1.09	2.30	0.81	1.16	<mark>1.90</mark>	0.88
ML						11	0.50	4.66	0.95	0.92	2.55	0.85	0.80	<mark>2.41</mark>	0.93
HL							0.83	2.82	0.87	0.91	2.58	0.85	0.58	<mark>4.13</mark>	0.96
1	LO °C														
VIS-SWNIR	EMCVS	R	SD	MS	46	6	0.47	4.60	0.95	0.69	3.16	0.90	0.96	<mark>3.32</mark>	0.94
NIR	EMCVS	log(1/R)	SNV		19	9	0.44	4.93	0.96	0.67	3.22	0.90	1.15	<mark>2.23</mark>	0.87
LL	EMCVS	log(1/R)	None		11	6	0.46	4.66	0.95	0.58	3.71	0.93	0.94	<mark>3.03</mark>	0.94
ML						6	0.47	4.64	0.95	0.75	2.89	0.88	1.17	<mark>2.17</mark>	0.84
HL							0.36	6.07	0.97	0.51	4.30	0.95	0.97	<mark>3.28</mark>	0.94
4	•C and 10	D°C													
VIS-SWNIR	EMCVS	R	SNV	SD	8	4	0.94	2.48	0.84	1.05	2.22	0.80	0.95	<mark>2.10</mark>	0.85
NIR	EMCVS	R	SNV	SD	96	6	1.04	2.24	0.80	1.18	1.98	0.74	1.15	<mark>1.75</mark>	0.77
LL	EMCVS	R	SD	LD	35	4	0.96	2.45	0.83	1.12	2.08	0.77	0.87	<mark>2.27</mark>	0.88
ML						5	0.79	2.96	0.89	0.93	2.51	0.84	1.32	<mark>1.49</mark>	0.74
HL							0.84	2.79	0.88	0.94	2.47	0.84	0.89	<mark>2.27</mark>	0.86

23 EMCVS, ensemble Monte Carlo variable selection; VIP, variable importance projection ; SD, second derivative; SNV, standard normal variate;

AsLs, asymmetric least squares; LD, linear detrend; MS, medium scaled: #Bands, wavelengths used for model development; #LVs, latent variables. The best model for each storage temperature is highlighted in bold.



Fig. 1. Log (1/R) NIR pre-treated (SD+AsLs) spectra of beef *LD* samples stored at 4 °C for selected storage times. Bands selected by the EMCVS method to predict TVC of samples are highlighted in blue.



Fig. 2. Reflectance VIS-SWNIR pre-treated (SD+MS) spectra of beef *LD* samples stored at 10 °C for selected storage times. Bands selected by the EMCVS method to predict TVC of samples are highlighted in blue.



Fig. 3. Measured vs predicted TVC for the best performing PLS-R models developed for beef *LD* samples stored at (a) 4 °C, (b) 10 °C and (c) either 4 °C or 10 °C.

Journal Pre-proof



Fig. 4. Prediction maps for TVC of beef *LD* samples (log CFU/g) stored at 10 °C for selected times using the reflectance VIS-SWNIR pre-treated (SD+MS) spectra which selected 46 bands).

- Microbial quality of beef stored under normal or abuse conditions can be predicted
- Spectral pre-treatments, band selection and data fusion methods are key for robust model development
- Hyperspectral imaging and chemometrics have potential for real-time monitoring of microbial quality

Journal Pre-proof

Conflict of Interest and Authorship Conformation Form

Please check the following as appropriate:

- ✓ All authors have participated in (a) conception and design, or analysis and interpretation of the data; (b) drafting the article or revising it critically for important intellectual content; and (c) approval of the final version.
- ✓ This manuscript has not been submitted to, nor is under review at, another journal or other publishing venue.
- ✓ The authors have no affiliation with any organization with a direct or indirect financial interest in the subject matter discussed in the manuscript
- The following authors have affiliations with organizations with direct or indirect financial interest in the subject matter discussed in the manuscript:

Author's name	Affiliation
	0
30-	