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1 **Improvement in spectral library-based quantification of soil properties using**
2 **representative spiking and local calibration – the case of soil inorganic carbon**
3 **prediction by mid-infrared spectroscopy**

4

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22

23

24 **Abstract**

25 Mid-infrared reflectance spectroscopy (MIRS) is time- and cost-effective. It was used for
26 quantifying soil inorganic carbon (SIC) concentration in France based on a national library,
27 and performances were evaluated on an independent regional set. Our objective was to
28 improve the accuracy of MIRS predictions based on common multivariate regression, through
29 spiking (enrichment of the national library with some representative target samples) with
30 possible extra-weighting (replication of spiking samples) and local calibration (only using
31 calibration samples that are spectral neighbours of each target samples), which have not been
32 fully explored yet, in combination especially.

33 Global (i.e. common) calibration yielded accurate prediction (standard error of prediction,
34 SEP, was $\approx 5 \text{ g kg}^{-1}$), which could be improved when the library was completed with spiking
35 samples (optimally 10 samples extra-weighted 40 times; $\text{SEP} = 3.3 \text{ g kg}^{-1}$). Using spiking
36 samples only (without the library) yielded slightly less accurate results ($\text{SEP} = 3.6 \text{ g kg}^{-1}$).
37 Prediction was more accurate using local calibration without spiking, but on a validation set
38 that was reduced because some validation samples lacked calibration neighbours ($\text{SEP} = 2.5$ -
39 2.7 g kg^{-1}). Local calibration with spiking (optimally 10 samples without extra-weight)
40 yielded somewhat less accurate prediction but for the full validation set when few calibration
41 neighbours were required ($\text{SEP} = 2.7 \text{ g kg}^{-1}$), or higher accuracy on the reduced validation set
42 when many neighbours were required ($\text{SEP} = 2.3 \text{ g kg}^{-1}$).

43 These accurate predictions demonstrated the usefulness of representative spiking and local
44 calibration for rendering large soil spectral libraries fully operational, while extra-weighting
45 had no additional benefit. Along with more exhaustive spectral libraries, this paves the way
46 for extensive use of MIRS for SIC determination.

47

48 **Highlights**

- 49 • We used a national MIRS library for quantifying soil inorganic C in a French region
- 50 • Combining representative spiking and local calibration improved prediction markedly
- 51 • Requiring more calibration neighbours increased accuracy but on less samples
- 52 • MIRS could increasingly replace calcimetry for SIC analysis

53

54 **Keywords:** Diffuse reflectance spectroscopy, national spectral library, carbonates, France.

55

56 **1. Introduction**

57 In the context of increasing carbon dioxide (CO₂) content in the atmosphere, there is growing
58 interest in the potential of soils to sequester carbon (Lal, 2004; Lal et al., 2015; Dignac et al.,
59 2017). Comprehensive understanding of sinks and sources of soil carbon and their
60 relationships with global change is a priority research topic, in order to produce accurate
61 estimates of future atmospheric CO₂ concentrations. The soil carbon pool comprises two
62 distinct components, soil organic carbon (SOC) and soil inorganic carbon (SIC), which
63 roughly contribute two-thirds and one-third, respectively (Batjes, 2014). Attention is paid
64 firstly to SOC, which is larger and considered more dynamic than SIC and a key component
65 in soil functioning. Nevertheless SIC is a pool of carbon that matters and may evolve with
66 time, land use and climate change (Emmerich, 2003; Sanderman, 2012; Chevallier et al.,
67 2016), thus is worth quantifying. Moreover, to date, quantifying SOC in calcareous soils often
68 requires determining SIC, SOC being consequently calculated by difference between total
69 carbon determined by dry combustion and SIC determined by calcimetry. Direct measurement
70 of SOC in calcareous soils is possible, through wet oxidation (Walkley and Black, 1984) or
71 dry combustion after acid fumigation (Harris et al., 2001), but these methods use hazardous
72 and polluting reactants, and have been regularly criticized for inaccurate results; thus
73 alternative approaches are being looked for (Apesteguia et al., 2018).

74 The usefulness of infrared diffuse reflectance spectroscopy coupled with chemometrics for
75 quantifying soil properties cost- and time-effectively has been demonstrated extensively
76 (Viscarra Rossel et al., 2006; Cécillon et al., 2009; Soriano-Disla et al., 2014; Gredilla et al.,
77 2016). Much attention has been paid to the prediction of SOC concentration by near infrared
78 reflectance spectroscopy (NIRS), but better predictions have often been achieved for SIC
79 concentration and/or using mid-infrared reflectance spectroscopy (MIRS; McCarty et al.,
80 2002; Bellon-Maurel and McBratney, 2011; Barthès et al., 2016). Quantification by NIRS and
81 MIRS requires calibrations, using samples that have been characterized both spectrally and
82 conventionally (e.g. dry combustion for total carbon and calcimetry for SIC). Large soil
83 databases that include conventionally- and spectrally-analysed samples representative at a
84 country or even wider scale are now becoming available (Brown et al., 2006; Genot et al.,
85 2011; Grinand et al., 2012; Viscarra Rossel and Webster, 2012; Stevens et al., 2013; Viscarra
86 Rossel et al., 2016). Such spectral libraries pave the way for the development of many
87 applications, as indicated by several works that used large-scale NIRS databases for making
88 predictions on target sample sets, for instance at local scale (Brown, 2007; Sankey et al.,
89 2008; Guerrero et al., 2010, 2014; Gogé et al., 2014). Some studies tested specific approaches
90 to improve NIRS or MIRS predictions when using large soil spectral libraries, for instance

91 local calibration: instead of using all library samples for building a prediction model, local
92 calibration selects those most appropriate considering the target set, and in some cases, even
93 builds a prediction model for each target sample separately, using calibration samples that are
94 its spectral neighbours (Shenk et al., 1997; Ramirez-Lopez et al., 2013; Lobsey et al., 2017).
95 Other studies tested spiking, which consists of enriching the calibration set with some samples
96 originating from the target set (Sankey et al., 2008; Guerrero et al., 2010); these spiking
97 samples can even be extra-weighted, to increase their influence in the calibration database
98 (Guerrero et al., 2014). Noticeable improvements in predictions have been reported when
99 applying local calibration (Genot et al., 2011; Rabenarivo et al., 2013; Gogé et al., 2014;
100 Clairotte et al., 2016; Lobsey et al., 2017) or spiking, without extra-weighting (Guerrero et al.,
101 2010; Gogé et al., 2014), and even more, with extra-weighting (Guerrero et al., 2014, 2016).
102 But local calibration and spiking have rarely been used in conjunction (Gogé et al., 2014), and
103 have never been used with spectrally representative spiking samples or extra-weighting.
104 The French soil quality monitoring network (Réseau de mesures de la qualité des sols,
105 RMQS; Arrouays et al., 2002) represents a collection of soil samples originating from more
106 than 2100 sites located regularly over the whole French metropolitan territory. A range of soil
107 properties have been analysed on these samples, and their NIR and MIR spectra have been
108 collected (Gogé et al., 2012; Grinand et al., 2012; Clairotte et al., 2016).
109 The objective of this work was to improve MIRS prediction of SIC concentration in France
110 based on the RMQS spectral library, through spiking with representative target samples
111 (possibly extra-weighted) and local calibration, which have not been fully explored yet, in
112 combination especially (i.e. local calibration after spiking).

113

114 **2. Materials and methods**

115 *2.1. National soil library*

116 The soil samples used as calibration database belong to a large national soil library provided
117 by the French national soil quality monitoring network (RMQS; Arrouays et al., 2002). The
118 RMQS aims at providing a national overview of soil quality, identifying gradients, monitoring
119 the evolution of soil quality over time with a frequency of a decade, and building a bank of
120 soil samples. This soil library was built during a 10-year sampling campaign over the
121 552 000 km² of the French metropolitan territory (Corsica included), which latitude ranges
122 from 41 to 51°N and longitude from 5.0°W to 9.5°E. The sampling design was based upon a
123 square grid with 16-km spacing. At the centre of each square, 25 individual core samples were
124 taken at 0–30 cm depth using an unaligned sampling design within a 20 × 20 m area, and

125 were then bulked to obtain composite samples (Arrouays et al., 2002). Samples were also
126 collected at 30–50 cm, but these were not considered for this study. In total, 2178 samples
127 were considered here, representing numerous soil types: Cambisols, Calcisols, Luvisols,
128 Leptosols, Andosols, Albeluvisols, etc. (IUSS Working Group WRB, 2014).

129

130 2.2. *Target set*

131 The target set was made of 164 composite topsoil samples originating from commercial
132 vineyard plots (one sample per plot) located in nine villages or small towns of the Languedoc-
133 Roussillon region, in southern France. Vine represents the main agricultural production in
134 Languedoc-Roussillon, which is the largest vine-growing region in France, and the studied
135 vineyard plots were considered representative of Languedoc-Roussillon (Coll, 2011). The
136 nine sites are presented in Table 1. Briefly, they have latitude ranging from 42.5 to 44.0°N,
137 longitude from 2.5 to 4.0°E, elevation from 5 to 358 m a.s.l., with either Calcisols, Arenosols,
138 Cambisols or Luvisols (IUSS Working Group WRB, 2014). Samples were taken from plots
139 under different vineyard management systems, which varied according to the types of
140 pesticides and fertilizers used, and possible inter-row weeding and grass cover. Samples were
141 collected in inter-row centre at 0–15 cm depth using a hand-held gouge auger. Ten soil
142 subsamples were taken in each plot and carefully mixed to form a composite sample. More
143 information on the sites and samples has been provided by Coll (2011) and by Salomé et al.
144 (2014, 2016). This set was independent from the national library.

145

146 2.3. *Soil conventional analysis*

147 All determinations of carbonate content were made in the Laboratoire d'analyses des sols
148 from INRA (Institut national de la recherche agronomique) in Arras, France, which is
149 ISO/CEI 17025:2005 accredited. They were carried out on finely ground (< 0.25 mm)
150 aliquots of 2-mm sieved air-dried soils samples using a Bernard calcimeter, according to the
151 standard French procedure NF ISO 10693 (ISO, 1995), which consists of measuring the
152 volume of CO₂ produced after the addition of chlorhydric acid (Pansu and Gautheyrou, 2006).
153 The carbonate content was calculated after calibration with a pure calcium carbonate (CaCO₃)
154 standard and was expressed as equivalent CaCO₃ content. Soil inorganic carbon content was
155 calculated as $0.12 \times \text{soil CaCO}_3 \text{ content}$ (because the molar mass of carbon is 12 g mol⁻¹ and
156 that of CaCO₃ is 100 g mol⁻¹).

157 According to the NF ISO 10693 procedure (ISO, 1995), acceptable repeatability for this
158 analysis is 0.4 g kg⁻¹ when SIC ≤ 6 g kg⁻¹, 6% between 6 and 18 g kg⁻¹, 1.1 g kg⁻¹ between 18

159 and 21.6 g kg⁻¹, and 5% when SIC > 21.6 g kg⁻¹. Replicated measurements allowed
160 calculating the expanded uncertainty associated with SIC analysis in the above-mentioned
161 laboratory. Expanded uncertainty increases the standard measurement uncertainty, calculated
162 as the quadratic sum of measurement repeatability and intermediate precision, by multiplying
163 it by a coverage factor (> 1) related to the confidence level requested (JCGM, 2012).
164 Measurement repeatability was calculated based on triplicate SIC analysis of 20 samples from
165 an inter-laboratory network; intermediate precision, based on the analysis of control samples
166 over time; and the coverage factor was fixed at 2 for achieving a confidence level of 0.95. In
167 such conditions, and using a modelling approach, the expanded uncertainty was empirically
168 estimated according to Equation 1:

$$\text{Expanded uncertainty} = 0.025 \text{ SIC} + 0.13 \text{ (SIC in g kg}^{-1}\text{)} \quad \text{Equation 1}$$

169 Expanded uncertainty was considered as standard error of laboratory (SEL). On the target site,
170 where SIC averaged 19 g kg⁻¹ (Table 1), SEL could thus be estimated at 0.6 g kg⁻¹ [i.e.
171 (0.025×19)+0.13].
172

173

174 *2.4. Measurement of mid-infrared reflectance (MIR)*

175 Air-dried, 2-mm sieved then finely ground samples were oven-dried overnight at 40°C before
176 spectral analysis. Reflectance spectra in the mid-infrared region were acquired at 934
177 wavenumbers between 4000 and 400 cm⁻¹ (i.e. 2500 and 25 000nm, respectively) at
178 3.86 cm⁻¹ interval using a Fourier transform Nicolet 6700 (Thermo Fischer Scientific,
179 Madison, WI, USA). This spectrophotometer is equipped with a silicon carbide source, a
180 Michelson interferometer as dispersive element, and a DTGS (deuterated triglycine sulfate)
181 detector. Soil samples were placed in a 17-well plate, where their surface was flattened with
182 the flat section of a glass cylinder, and they were then scanned using an auto-sampler (soil
183 surface area scanned: ca. 10 mm²). Each MIR spectrum resulted from 32 co-added scans, and
184 the body of the plate (beside wells) was used as reference standard and scanned once per plate
185 (i.e. every 17 samples). Reflectance was converted into apparent absorbance, which is the
186 decimal logarithm of the inverse of reflectance. Twenty wavenumbers were removed due to
187 often noisy spectrum end, and MIR spectra were used in the range from 4000 to 478 cm⁻¹
188 (2500 and 20 909 nm, respectively).

189

190 *2.5. Data analysis*

191 *2.5.1. Principles*

192 Mid-infrared absorbance spectra and SIC measurements of the national library samples were
 193 used to build a calibration model that expressed SIC content as a function of sample spectrum.
 194 The model was then applied on validation samples in order to predict their SIC content from
 195 their MIR spectrum. Modified Partial least squares regression (PLSR) was the linear
 196 multivariate regression procedure used to infer SIC content from spectra. To date, PLSR is the
 197 most common procedure for analysing infrared spectral data, in soil science in particular
 198 (Stenberg et al., 2010). It reduces a complex spectral matrix into a few orthogonal
 199 components (or terms, or latent variables LV), which are built in order to maximise their
 200 covariance with the variable of interest (here SIC; Bjørsvik and Martens, 2001). The
 201 modification proposed by Shenk and Westerhaus (1991) consisted of scaling the conventional
 202 data and the absorbance data at each wavelength to have a standard deviation of 1.0 before
 203 each PLS term. All calculations were done using the WinISI 4 software (Foss
 204 NIRSystems/Tecator Infracore International, State College, PA, USA). The accuracy of the
 205 prediction models was estimated on the validation set by computing the standard error of
 206 prediction (SEP, calculated according to Equation 2), bias (mean residual), coefficient of
 207 determination (R^2_{val}) and RPD_{val} ratio (ratio of SD_{val} to SEP, where SD_{val} is the standard
 208 deviation of the validation set).

$$209 \quad \text{SEP} = \sqrt{\frac{\sum_1^n (y_i - \hat{y}_i)^2}{n}} \quad \text{Equation 2}$$

210 where y_i and \hat{y}_i are the observed and MIRS-predicted values for sample i , and n the total
 211 number of samples in the validation set.

212

213 2.5.2. The validation set

214 Out of the 164 samples of the target set, 30 were not used for validation but were kept for
 215 spiking (cf. 2.5.5). The validation set then included 134 samples. The 30 samples kept for
 216 spiking were the most representative spectrally, selected according to a procedure proposed
 217 by Shenk and Westerhaus (1991):

- 218 - a principal component analysis (PCA) was performed on the spectra of the 164 target
 219 samples;
- 220 - distances between samples in the PCA space were calculated using the Mahalanobis distance
 221 H (Mark and Tunnell, 1985);
- 222 - the sample that had the most neighbours closer than a given distance d was selected while its
 223 neighbours were discarded;
- 224 - the process was continued until no samples remained with neighbours closer than d ;

225 - the distance d was set so that 30 samples could be selected.

226 A PCA was also performed on the spectra of the national library. The spectra of the target
227 samples were then projected onto this PCA space to calculate their Mahalanobis distance H
228 and evaluate how they were represented spectrally by the library (Mark and Tunnell, 1985).

229

230 2.5.3. Global calibration

231 Global PLSR, which is the common PLSR procedure, uses all calibration samples for
232 building a unique model that is then applied uniformly to all validation samples. The number
233 of PLSR latent variables that minimized the standard error of cross-validation (SECV) was
234 retained for the prediction model, with an upper limit set to 16 (SECV was calculated using
235 Equation 2, with n being the number of calibration samples). The cross-validation was carried
236 out by dividing the calibration set into four groups composed cyclically (i.e. the 1st, 5th, 9th
237 samples in the first group, the 2nd, 6th, 10th samples in the second group, etc.), after the
238 samples had been ranked by increasing SIC values. Three groups were used to develop the
239 model and one to test it, and the procedure was performed four times to use all samples for
240 both model development and prediction. The residuals of the four predictions were pooled to
241 calculate SECV.

242

243 2.5.4. Local calibration

244 In contrast, local PLSR makes prediction for each validation sample individually, only using
245 calibration samples that are its spectral neighbours (Shenk et al., 1997). These neighbours
246 were selected according to the correlation coefficient R between the spectra of calibration
247 samples and each validation sample, with an R cut-off value below which samples were not
248 considered neighbours. According to preliminary tests, this R cut-off value was set at 0.95.

249 The minimum number of calibration neighbours requested was varied from 4 to 50. It is
250 important to note that prediction was not made for validation samples that did not have
251 enough calibration neighbours. There was firstly no requirement on the maximum number of
252 calibration neighbours; then this maximum was set to 300 and 100, but this had limited
253 interest and the corresponding results will not be presented.

254 Actually local calibration differs from global calibration on two points: (i) prediction is made
255 on each validation sample individually, only using calibration neighbours, not all calibration
256 samples; and (ii) prediction is not made for validation samples that do not have enough
257 calibration neighbours (i.e. that are poorly represented by the calibration set). To separate
258 these both aspects, prediction was also made using all calibration samples (i.e. global

259 calibration), but only on the validation samples that had enough calibration neighbours. In
260 that case, predictions by local and global calibration were made on the same validation
261 samples, and the difference between local and global calibrations was only due to calibration
262 by neighbours.

263 In local calibration the number of latent variables was not determined through cross-
264 validation. Instead, each prediction was calculated as the weighted average of the predicted
265 values generated with 3 to 16 latent variables, each weight being calculated as the inverse of
266 the product of the root mean square (RMS; cf. Equation 3) of spectral residuals (i.e. the
267 difference between the actual spectrum and the spectrum approximated using the considered
268 number of latent variables) and RMS of the regression coefficients using the considered
269 number of latent variables (Shenk et al., 1997).

$$270 \quad \text{RMS} = \sqrt{\frac{\sum_1^n x_i^2}{n}} \quad \text{Equation 3}$$

271 where x_i are the values considered (i.e. spectral residuals or regression coefficients) and n
272 their number.

273

274 2.5.5. Spiking

275 Spiking consists of adding a few target samples to the calibration set, so that the prediction
276 model better suits the validation set (Guerrero et al., 2010). This was achieved using the 30
277 representative target samples removed from the validation set (cf. 2.5.2), which will thereafter
278 be called "spiking samples". Spiking was performed for both global and local calibrations and
279 different sizes were tested for the spiking subset: 7, 10, 15, 20 and 30 samples. When 7 to 20
280 samples were used for spiking, they were selected as the most spectrally representative in the
281 set of 30 samples removed from the target set (in which they were the most spectrally
282 representative, cf. 2.5.2).

283 Moreover, the spiking subset could be extra-weighted, which consisted of adding several
284 copies of this subset to the calibration set, in order to increase its influence in the calibration
285 (Guerrero et al., 2014). Different weights were tested, from 10 to 200, which means that each
286 spiking sample was represented by 10 to 200 replicates in the calibration set, respectively. Of
287 course spiking uses (some) target samples for calibration, but the improvement in model
288 performance really seems worthwhile (Guerrero et al., 2010, 2014, 2016). Spiking without or
289 with extra-weighting was performed for both global and local calibration.

290

291 2.5.6. Pre-processing

292 In addition, different common spectrum pre-processing methods, or pretreatments, were
293 tested: none (no scatter correction); standard normal variate transformation (SNV), which
294 consists of mean-centring and variance-scaling the spectrum; detrending (D), which consists
295 of removing a linear trend from the spectrum; SNVD (i.e. both SNV and D); and
296 multiplicative scatter correction (MSC), which consists of centering and scaling the spectrum
297 with, respectively, the intercept and slope calculated when regressing linearly this spectrum
298 against the average calibration spectrum (Geladi et al., 1985; Barnes et al., 1989). These
299 transformations were possibly followed by first derivation. Derivation aims at reducing
300 baseline variation and enhancing spectral features, and was calculated over a 5-, 15- or 20-
301 point gap, with 5-point smoothing, in order to reduce signal random noise (Bertrand, 2000).
302 The conditions of derivation were denoted 155, 1155 and 1205 for first derivation with 5-, 15-
303 and 20-point gap, respectively, and 5-point smoothing. No derivation and no smoothing was
304 denoted 001.

305

306 **3. Results**

307 *3.1. Distributions of SIC content*

308 In the national library and target set, minimum SIC content was ≈ 0 g kg⁻¹ (under the
309 detection limit), maximum was 104 and 86 g kg⁻¹, mean 6.4 and 18.8 g kg⁻¹, median ≈ 0 and
310 13.3 g kg⁻¹, and standard deviation 16.0 and 21.4 g kg⁻¹, respectively. The national library was
311 dominated by samples with no SIC (53%) or very low SIC content (22% with
312 $0 < \text{SIC} < 1$ g kg⁻¹), while soils with SIC > 20 and > 50 g kg⁻¹ accounted for 12% and 4% of
313 the library, respectively (Fig. 1). Indeed, the French territory has large mountain regions with
314 acid rocks and soils (Arrouays et al., 2011). In the target set, samples with no SIC accounted
315 for a noticeable proportion of the set (31%), but SIC-rich soils were more frequent: samples
316 with SIC > 20 and > 50 g kg⁻¹ accounted for 43% and 11% of the set, respectively (Fig. 1).
317 Carbonated rocks and soils cover an important proportion of the region considered, in
318 southern France.

319

320 *3.2. Prediction of SIC content using global calibration without spiking*

321 Global calibration without spiking represents usual calibration, built with the whole national
322 library. In these conditions, the best MIRS predictions were achieved using the pretreatment
323 None1155 (i.e. first derivation with 15-point gap and 5-point smoothing) and yielded
324 SEP = 5.2 g kg⁻¹ and RPD_{val} = 3.7 (Table 2 and Fig. 2a).

325 Predictions were particularly poor for the samples that originated from Aigues-Mortes. When
326 projected onto the PCA space of the national library, most Aigues-Mortes samples were not
327 close to national library samples (Fig. 3), though only two were spectral outliers formally (i.e.
328 their Mahalanobis distance H with the national library, i.e. its centre, was > 3). Salomé et al.
329 (2014), who studied the same sample set, also observed that the soils from Aigues-Mortes
330 were particular, being quite rich in SIC though very sandy.

331

332 *3.3. Prediction of SIC content using global calibration with spiking*

333 3.3.1. Global calibration using spiking samples only (and not the national library)

334 Firstly calibrations were built using only spiking samples, to address their importance, and the
335 national library was not used. This led to surprisingly accurate predictions on the 134
336 validation samples (SEP ranged from 4.5 to 3.6 g kg⁻¹ and RPD_{val} from 4.3 to 5.3; Table 2 for
337 10 and 30 spiking-calibration samples, and Fig. 4). Actually global calibration with 10
338 representative samples from the target site led to more accurate predictions than global
339 calibration using the national library (RPD_{val} = 5.3 vs. 3.7, respectively).

340

341 3.3.2. Global calibration using the national library; effect of the number of spiking samples

342 Then calibrations were built using the national library enriched with representative spiking
343 samples. Adding one copy of the spiking samples had little effect on prediction, due to the
344 large size of the library (SEP = 4.9 g kg⁻¹ and RPD_{val} = 3.9 with 10 to 30 spiking samples;
345 Table 2 and Fig. 4), as also observed by Guerrero et al. (2014). So the spiking samples were
346 extra-weighted, meaning that each was replicated 10 to 200 times (Guerrero et al., 2014).

347 As could be expected, prediction accuracy tended to increase with the number of spiking
348 samples; but rather unexpectedly, the additional benefit was limited beyond 10 spiking
349 samples (Fig. 4). Predictions using 7 spiking samples were rather poor (SEP \approx 5 g kg⁻¹ and
350 RPD_{val} $<$ 4 with extra-weight \geq 10), probably because 7 spiking samples were not sufficient to
351 represent the validation set correctly.

352

353 3.3.3. Global calibration using the national library; effect of the weight of spiking samples

354 As could also be expected, prediction accuracy increased with the weight of spiking samples,
355 but the additional benefit was limited beyond 40 replicates (Fig. 4): with 10 spiking samples
356 replicated 40 times vs. 30 spiking samples replicated 200 times, SEP was 3.3 vs. 3.0 g kg⁻¹
357 and RPD_{val} 5.9 vs. 6.4, respectively (Table 2). Thus prediction accuracy was little improved
358 when multiplying the number of spiking samples by three and their weight by five, which

359 required three times more conventional analyses, noticeably more computational time, thus
360 did not seem relevant. Computational time was not measured. However 10 spiking samples
361 replicated 40 times vs. 30 spiking samples replicated 200 times represented additions of 400
362 vs. 6000 samples to a library already including 2178 samples, resulting in calibration datasets
363 of 2578 vs. 8178 samples, respectively.

364 It seemed that SEP continued to decrease slightly when the weight of spiking samples
365 increased beyond 200, but this was not explored because it required much computational time.
366 It would be expected that, with ever increasing weight of spiking samples, prediction
367 performance would approach that achieved when using spiking samples only (SEP = 3.6-
368 4.0 g kg⁻¹ for ≥ 10 spiking samples; Fig. 4).

369

370 *3.4. Prediction of SIC content using local calibration without spiking*

371 3.4.1. Number of calibration neighbours

372 Local calibration makes prediction on each validation sample separately, only using
373 calibration samples that are its spectral neighbours according to correlation between spectra,
374 with an R cut-off value that was set to 0.95; and there is no prediction for validation samples
375 that do not have enough calibration neighbours. Fig. 5a presents the distribution of validation
376 samples according to their number of spectral neighbours in the national library; in particular,
377 1 validation sample had no calibration neighbour, 5 samples had 4 to 9 calibration neighbours,
378 and all 6 samples with < 10 calibration neighbours originated from Aigues-Mortes (cf. 3.2).
379 Most validation samples had more than 100 calibration neighbours, except those from
380 Aigues-Mortes, which had 22 in average (data not shown).

381

382 3.4.2. Local calibration without spiking; effect of the number of calibration neighbours 383 required

384 When the minimum number of calibration neighbours was set to 4, local calibration had
385 limited benefit when compared with global calibration (SEP = 4.4-4.8 g kg⁻¹ and
386 RPD_{val} = 4.1-4.4 on 133 validation samples; Table 3). When the minimum number of required
387 neighbours was increased from 4 to 10, 20, 30 then 50, progressively the number of validation
388 samples that had enough neighbours decreased to 115 samples, while prediction accuracy
389 increased (SEP decreased to 2.7 g kg⁻¹ and RPD_{val} increased to 7.6; Table 3 for ≥ 50
390 neighbours, other data not shown).

391

392 3.4.3. Reasons for differences between local and global calibration without spiking

393 Actually local calibration did not yield much better prediction than global calibration on the
394 same validation samples ($RPD_{val} = 4.1$ vs. 4.0 on 133 samples with ≥ 4 neighbours required
395 and 7.6 vs. 6.4 on 115 samples with ≥ 50 neighbours, respectively, i.e. an increase ≤ 1.2 ;
396 Table 3). By contrast, global calibration on these validation samples that had enough
397 calibration neighbours yielded noticeably more accurate prediction in general than global
398 calibration on the whole validation set ($RPD_{val} = 6.4$ vs. 3.7 on 115 vs. 134 samples,
399 respectively, i.e. an increase of 2.6 when ≥ 50 neighbours were required), except when few
400 calibration neighbours were required ($RPD_{val} = 4.0$ vs. 3.7 on 133 vs. 134 samples,
401 respectively, with ≥ 4 neighbours; Tables 3 and 2). These comparisons show that without
402 spiking, most improvement in prediction accuracy from global to local calibration resulted in
403 general from removing validation samples with too few calibration neighbours; while the
404 benefit of calibration by neighbours was limited. This was however not the case when few
405 calibration neighbours were required for local calibration: in that case, improvement from
406 global to local calibration resulted mainly from calibration by neighbours, because almost all
407 validation samples had enough neighbours thus were predicted. But this improvement from
408 global to local calibration, due to calibration by neighbours, was limited.

409

410 *3.5. Prediction of SIC content using local calibration with spiking*

411 Local calibration was carried out using the national library enriched with 7 to 30 spiking
412 samples originating from the target sites, and extra-weighted 10 to 50 times, or not extra-
413 weighted. When extra-weighted, a spiking sample neighbouring a validation sample was
414 represented by several copies, among the calibration samples used for making a prediction on
415 this validation sample.

416

417 *3.5.1. Local calibration with spiking; effect of the number of spiking samples*

418 Local calibration with 7 spiking samples had little effect on prediction results when compared
419 with local calibration without spiking (e.g. with ≥ 50 calibration neighbours required,
420 $RPD_{val} = 7.6$ without spiking or with 7 spiking samples extra-weighted 10 times, on
421 133 samples; Table 3 without spiking, data not shown for 7 spiking samples).

422 Using 10 spiking samples was more useful, but the effects varied, depending firstly on the
423 number of calibration neighbours required (Table 3):

- 424 • with few neighbours required (i.e. 4), prediction was much better with than without
425 spiking, on all validation samples ($RPD_{val} = 7.2-7.3$ on 134 samples vs. 4.1 on

426 133 samples, respectively; Fig. 2b for extra-weight 1, i.e. no extra-weighting), because
427 spiking increased the number of calibration neighbours thus improved prediction accuracy;
428 • with many neighbours required (i.e. 50), the effect was less clear and depended on the
429 extra-weight of spiking samples:
430 - with low extra-weight ($\times 1$ or $\times 10$), prediction was moderately better with than without
431 spiking, on the same (reduced) validation set ($RPD_{val} = 8.9-9.1$ vs. 7.6 on 115 samples,
432 respectively; Fig. 2c for extra-weight 10), because the small number of replicates from
433 spiking samples did not provide enough calibration neighbours;
434 - with high extra-weight ($\times 50$), prediction was moderately less accurate with than without
435 spiking, but on all validation samples ($RPD_{val} = 6.8$ on 134 samples vs. 7.6 on
436 115 samples, respectively; Table 3), because the large number of replicates from
437 spiking samples provided enough calibration neighbours.
438 Using more than 10 spiking samples did not improve prediction thus had no interest (with
439 either 10 or 30 spiking samples $\times 1$ or $\times 10$, $RPD_{val} = 7.2-7.3$ on 134 samples with ≥ 4
440 neighbours required; and $RPD_{val} = 8.8-9.2$ on 115-116 samples with ≥ 50 neighbours;
441 Table 3). Surprisingly, the number of spiking samples had little effect on the number of
442 validation samples that had enough calibration neighbours (using either 10 or 30 spiking
443 samples, this number was 134 with ≥ 4 neighbours requested, and 115 or 116 with ≥ 50
444 neighbours; Table 3). Actually, the distribution of validation samples according to their
445 number of calibration neighbours was not much affected by the number of spiking samples
446 (Fig. 5b vs. 5e, 5c vs. 5f, and 5d vs. 5g).

447

448 3.5.2. Local calibration with spiking; effect of the weight of spiking samples

449 Extra-weighting the spiking samples had little effect on prediction results, with either few
450 calibration neighbours required (i.e. 4; with 10 spiking samples $\times 1$ to $\times 50$, $RPD_{val} = 7.2-7.3$
451 on 134 samples) or many neighbours required (i.e. 50; with 10 spiking samples $\times 1$ or $\times 10$,
452 $RPD_{val} = 8.8-9.2$ on 115-116 samples; Table 3). Thus extra-weighting had limited interest in
453 local calibration, which contrasted with global calibration. High extra-weight of spiking
454 samples allowed making prediction on all 134 validation samples when many neighbours
455 were required (i.e. 50; for 10 spiking samples $\times 50$, $RPD_{val} = 6.8-7.2$); but similar result was
456 achieved when few neighbours were required (for 10 spiking samples $\times 1$, $RPD_{val} = 7.3$;
457 Table 3).

458

459 3.5.3. Local calibration with spiking; effect of the number of calibration neighbours required
460 As observed for local calibration without spiking (cf. 3.4.2), increasing the minimum number
461 of calibration neighbours improved prediction accuracy on a validation set that was reduced
462 (with 10 spiking samples $\times 1$ and ≥ 4 vs. ≥ 50 neighbours required, $SEP = 2.7$ vs. 2.3 g kg^{-1} and
463 $RPD_{\text{val}} = 7.3$ vs. 8.9 on 134 vs. 115 samples, respectively); but this effect vanished at high
464 extra-weight (with 10 spiking samples $\times 50$ and ≥ 4 vs. ≥ 50 neighbours required, $SEP = 2.7$
465 vs. 2.7 g kg^{-1} and $RPD_{\text{val}} = 7.2$ vs. 6.8 on 134 samples, respectively; Table 3).

466

467 3.5.4. Reasons for differences between local and global calibration with spiking

468 To compare local and global calibration with spiking, it was useful to consider whether
469 prediction was possible for all validation samples or not, as was also the case without spiking.
470 Prediction was possible for all validation samples when all had enough calibration
471 neighbours, which was achieved (i) when few calibration neighbours were required, or
472 (ii) when many were required but spiking samples were highly extra-weighted. In that case
473 where prediction was possible for all validation samples, the difference with global calibration
474 was not due to the removal of validation samples with too few calibration neighbours. The
475 difference was only due to calibration by neighbours, and the benefit of local calibration was
476 noticeable, but it decreased when extra-weighting increased (with 10 spiking samples $\times 1$, $\times 10$
477 and $\times 50$, local calibration with ≥ 4 neighbours vs. global calibration yielded $RPD_{\text{val}} = 7.3$ vs.
478 3.9 , 7.2 vs. 5.0 and 7.2 vs. 5.9 on 134 samples, respectively; Table 3). Indeed, extra-weighting
479 the spiking samples improved prediction using global calibration but not using local
480 calibration. By contrast, when prediction was not possible for some validation samples with
481 too few calibration neighbours, which in general were poorly predicted otherwise, the
482 difference between local and global calibration was larger and had two causes: the removal of
483 poorly predicted samples from the validation set, which was the dominant cause in general,
484 and calibration by neighbours (with 10 spiking samples $\times 1$ and $\times 10$, local calibration with
485 ≥ 50 neighbours vs. global calibration on the same validation samples vs. global calibration
486 on the complete validation set yielded $RPD_{\text{val}} = 8.9$ - 9.1 vs. 7.5 - 7.8 vs. 3.9 - 5.0 on 115 vs. 115
487 vs. 134 samples, respectively; Tables 3 and 2, and Fig. 2c and 2d for local and global
488 calibration with same calibration and validation samples).

489

490 3.6. *Complementary considerations*

491 The absolute value of bias was large ($< -1 \text{ g kg}^{-1}$) with global calibration and < 10 spiking
492 samples, or ≥ 10 spiking samples extra-weighted < 20 times. With ≥ 10 spiking samples or, in
493 local calibration, with ≥ 50 neighbours required, prediction improvement was mainly an effect
494 of bias reduction. Guerrero et al. (2014) also attributed to large bias the poor prediction of
495 SOC content of target samples when using a large unspiked NIRS library.
496 Mean SIC was 17.8 g kg^{-1} on the validation set, thus SEL could be estimated at 0.6 g kg^{-1}
497 according to Equation 1 (with SEL = expanded uncertainty), and its contribution to SEP was
498 limited.

499

500 **4. Discussion**

501 *4.1. Global calibration without spiking*

502 Studying almost the same national library than in the present work, also with MIRS, Grinand
503 et al. (2012) obtained $\text{SEP} = 2.9 \text{ g kg}^{-1}$ and $\text{RPD}_{\text{val}} = 5.6$ in average when calibrating
504 randomly (with five replicates) on 20% of the library and validating on 80%, and
505 $\text{SEP} = 2.4 \text{ g kg}^{-1}$ and $\text{RPD}_{\text{val}} = 6.8$ when calibrating on 80% and validating on 20%. This
506 seems better than in the present study, but actually, mean SIC was smaller in the national
507 library than in our target set, and SEP represented 38% to 45% of the mean vs. 29% here.
508 McCarty et al. (2002) made global MIRS calibrations on profile samples from 14 locations in
509 nine US states, with RPD_{val} that could be estimated at 8.0 in external random validation (i.e.
510 possibly non-independent) and at 2.4 in independent validation (one-site-out). This pointed
511 out that, even for soil attributes linked to molecular functional groups that absorb in the MIR
512 range, such as SIC content, global calibration without spiking may lead to poor independent
513 validation when validation samples are poorly represented by calibration samples, as noticed
514 by Guerrero et al. (2014) from their study and several other works. Similar performances
515 ($\text{RPD}_{\text{val}} = 2.4$) were achieved in independent validation after global visible-and-NIRS
516 (VNIRS) calibration by Brown et al. (2006) on a very large and diverse set of
517 > 4100 independent samples from four continents, and by Morgan et al. (2009) on a set of
518 core samples from six central Texas fields. Brown et al. (2006) used boosted regression trees,
519 which outperformed PLS regression, and achieved $\text{RPD}_{\text{val}} = 2.7$ when using sand content as
520 auxiliary predictor. Comstock et al. (2019) used a national US library made of 1268 MIR
521 spectra and PLS global calibration for predicting SIC on 209 independent samples in two US
522 states, and achieved $\text{SEP} = 3.1 \text{ g kg}^{-1}$ and $\text{RPD}_{\text{val}} = 6.4$. This suggests that the target samples
523 were better represented by the national library than in the present study, though this aspect
524 was not specified, from spectral viewpoint especially. These references and our work

525 underline that prediction accuracy is greatly affected by sample set diversity, especially by the
526 distribution of the variable of interest and by the ability of the calibration set to represent the
527 validation samples, which vary across studies thus render comparisons difficult.

528

529 *4.2. Global calibration with spiking*

530 The results achieved when using spiking samples only (and not the national library) for
531 calibration were not very different from those reported by Grinand et al. (2012). These
532 authors used MIRS for predicting SIC within the same library than in the present study:
533 random calibration on 10% of the library yielded $RPD_{val} = 5.0$ on the remaining samples,
534 close to $RPD_{val} = 5.3$ achieved here when using 10 representative target samples to make
535 prediction on 134 (i.e. 7%).

536 Moreover, several literature papers have reported NIRS or VNIRS predictions at local scale
537 through global calibration on a large soil library completed with spiking samples. Sankey et
538 al. (2008) performed global VNIRS calibration at local scale in north-western USA and
539 reported better SIC predictions when enriching a national soil library with spiking samples,
540 but predictions were not necessarily better than when using these spiking samples only (i.e.
541 without the national library). These authors used more spiking samples than in the present
542 study (50% of the target set), which could explained the limited benefit of the national library;
543 but we also achieved comparably accurate predictions when using local samples only than
544 when using them for spiking the national library (Table 2). Overall, Sankey et al. (2008)
545 achieved poor predictions ($RPD < 1.6$, $SEP > 80\%$ of the mean), probably due to less
546 informative spectral range (VNIR vs. MIR) and to random selection of spiking samples,
547 which did not optimize their representativeness. Guerrero et al. (2010) also observed that
548 global NIRS-PLSR calibration of soil nitrogen (N) in south-eastern Spain using only a few
549 samples from the target site could yield similarly and even more accurate validations than
550 large regional libraries spiked with these samples; thus additional information provided by
551 regional libraries was not particularly useful. The spiking samples, which were not replicated,
552 had more influence in general when the library size decreased (different sizes being
553 considered). This led to the idea of extra-weighting the spiking samples, the interest of which
554 was demonstrated next by Guerrero et al. (2014). These authors used a national NIRS library
555 from Spain to build global PLS calibrations of SOC content for target sites in Spain, UK and
556 Sweden. They got better predictions when the spiking subset was selected according to
557 spectral representativeness, which was our choice. Moreover, they observed for large libraries
558 that increasing the weight of spiking samples was more useful than increasing their number.

559 They did not fully explore the effect of varying extra-weighting but tested a weight
560 determined as the ratio of library size to target set size, which would yield 13 in the present
561 study. We also observed that extra-weighting of spiking samples was more fruitful than
562 spiking our national library with more than 10 spiking samples, because these represented the
563 target set correctly. Guerrero et al. (2014) reported accurate predictions when using spiking
564 samples only, but more accurate predictions when using libraries enriched with extra-
565 weighted spiking samples, which was also our observation. In contrast, predicting SIC in a
566 24-km² area using the same national library than in the present study but in the VNIR range,
567 Gogé et al. (2014) found that spiking the national library had limited benefit. They attributed
568 this result to the strong spectral features of carbonates, which would render the national
569 library sufficient, while spiking improved prediction accuracy for soil properties supposedly
570 less strongly featured (SOC, clay, etc.). In this work, the spiking samples were selected at
571 random, thus not particularly representative of the target site, moreover they were not extra-
572 weighted, and we can assume this also limited the benefit of spiking.

573

574 *4.3. Local calibration without spiking*

575 We observed that some validation samples had few calibration neighbours, thus were poorly
576 represented by the national library, though most of them were not spectral outliers (according
577 to the Mahalanobis distance H calculated between each validation sample and the library
578 centre, cf. 3.2). Brown et al. (2005) developed and applied global VNIRS calibrations on six
579 sites of north-western USA having similar features, and similarly observed, in one-site-out
580 cross-validations, that SIC predictions might be poor in some individual sites though they
581 were well represented spectrally by the other sites. This underlines that target samples may be
582 well represented globally by a calibration set according to usual tests (e.g. H distance between
583 target samples and the calibration set, i.e. its centre), but nevertheless have few spectral
584 neighbours in this set, which often leads to poor predictions.

585 In their above-cited paper that dealt with VNIRS prediction of SIC in a 24-km² area, based on
586 the same national library than in the present study, Gogé et al. (2014) compared global
587 calibration with locally weighted calibration involving 300 neighbours (in this local
588 calibration approach, neighbours' contribution to prediction increases with their spectral
589 similarity to the target sample). They found little difference in prediction accuracy between
590 both approaches. This again was attributed to the strong spectral features of carbonates, which
591 would be correctly represented by the whole library (i.e. would render specific representation
592 by spectral neighbours useless), while local calibration was more useful for predicting soil

593 properties supposedly less strongly featured (e.g. SOC or clay). Moreover, the authors
594 underlined that the optimal number of calibration neighbours, which was fixed according to
595 preliminary work, depended on the soil property considered and on the library.

596 Local calibration involves two aspects: not predicting validation samples with too few
597 calibration neighbours, on the one hand, and calibration by neighbours, on the other hand. In
598 the present study, calibration by neighbours had limited benefit without spiking: when local
599 calibration outperformed global calibration clearly, this was mainly due to not predicting
600 validation samples with too few calibration neighbours; and this occurred when many
601 calibration neighbours were required. This suggests that strongly improving prediction
602 accuracy at the level of the validation set (i.e. SEP, RPD_{val} , etc.) was only possible when
603 removing and making no prediction on validation samples poorly represented by the
604 calibration database. It seems relevant to not use a spectral library for making prediction on
605 samples it poorly represents; but having to make prediction on less samples to improve
606 overall prediction accuracy may be somewhat disturbing. Actually, making no prediction on
607 target samples poorly represented by the calibration set is not specific to local calibration, and
608 may occur in global calibration too: a cut-off value of a distance parameter between each
609 target sample and the calibration set can also be used for removing target samples that are
610 considered outliers (e.g. Castaldi et al., 2018, who used the H distance). We must admit the
611 incomplete ability of MIRS to quantify SIC and other soil properties on certain samples as
612 long as soil spectral libraries are incomplete. It is however predictable that the
613 representativeness of soil spectral libraries will become more and more complete with time.

614

615 *4.4. Local calibration with spiking*

616 Few authors have performed local calibration with spiking. Gogé et al. (2014) performed
617 locally weighted calibration with spiking for VNIRS prediction of SIC in a 24-km² target area
618 using the French national library. This approach had little benefit when compared to locally
619 weighted calibration without spiking and to global calibration with and even without spiking,
620 which again was attributed to the strong spectral features of carbonates. Limited effect of
621 spiking might also be attributed to random selection of spiking samples. Seidel et al. (2019)
622 used VNIRS and PLS for predicting SOC in two fields in Germany. Spectral neighbours of
623 target samples (50 per sample) were selected from a national library then all used for building
624 global calibration. Selecting spectral neighbours in the national library reminds of local
625 calibration, though prediction was not made for each target sample individually. However,
626 there was no threshold below which a calibration sample was not considered as neighbour;

627 and as target samples were poorly represented by the national library (as specified by PCA),
628 predictions were poor without spiking ($RPD_{val} \leq 1.3$ vs. ≤ 1.2 with the full national library).
629 This underlines the interest of such threshold (e.g. cut-off R), which ensures that samples used
630 for calibration are really neighbours, and that if neighbours are lacking, there will be no
631 prediction for the target samples considered, thus no risk of poor prediction. Then Seidel et al.
632 (2019) observed that spiking improved predictions markedly ($RPD_{val} \geq 2.5$, ≥ 2.9 and ≥ 3.3
633 with 10, 15 and 20 spiking samples, respectively).
634 In the present study, prediction using local calibration became possible for all validation
635 samples through spiking, when either few calibration neighbours were required or when
636 spiking samples were heavily extra-weighted. Prediction accuracy was similar in both cases,
637 indicating that extra-weighting had little interest in local calibration, which contrasted with
638 global calibration. When compared with global calibration with spiking, the benefit of local
639 calibration with spiking resulted often from not predicting validation samples poorly
640 represented by the calibration database, as already observed without spiking. But in contrast
641 to what was observed without spiking, noticeable benefit could also result from calibration by
642 neighbours, when the spiking samples were moderately extra-weighted (i.e. global calibration
643 was little improved by spiking) and few calibration neighbours were required (i.e. prediction
644 using local calibration was possible for all validation samples).
645 The optimisation of parameters proposed here could not be generalised, as it clearly depended
646 on the size, spectral diversity and spectral proximity of the calibration database and validation
647 set. Values proposed for number and extra-weight of spiking samples and number of
648 calibration neighbours could be seen as orders of magnitude.

649

650 **5. Conclusion**

651 Global calibration yielded accurate prediction ($RPD_{val} \approx 4$), which could be noticeably
652 improved when the library was completed with extra-weighted spiking samples (optimally
653 10 samples \times 40 times; $RPD_{val} \approx 6$). Prediction was more accurate using local calibration
654 without spiking, but on a validation set that was reduced from 134 to 115 samples
655 ($RPD_{val} \approx 8$; on the same reduced set, global calibration without spiking yielded $RPD_{val} \approx 6$).
656 Local calibration with spiking (optimally 10 samples without extra-weight) yielded
657 moderately less accurate prediction but for the full validation set when ≥ 4 calibration
658 neighbours were required ($RPD_{val} \approx 7$), or higher accuracy on 115-116 samples when
659 ≥ 50 neighbours were required ($RPD_{val} \approx 9$).

660 Of course, parameter optimisation depended on the size, diversity and proximity of the
661 calibration database and validation set, and the optimums proposed here could not be
662 generalised (number and extra-weight of spiking samples, number of calibration neighbours).
663 Nevertheless, the results of the present study confirmed that MIRS prediction of SIC is
664 accurate, and demonstrated that the usefulness of large soil spectral libraries can be improved
665 by the combination of representative spiking and local calibration, while extra-weighting of
666 spiking samples has no additional effect.

667 Local calibration raised the question of either making prediction, or not, on validation samples
668 poorly represented by the calibration database, as long as the database is incomplete; and the
669 choice depends on user's priority (e.g. most accurate prediction or prediction on all samples).
670 Anyway, identifying such samples is crucial, at least to know that prediction is not fully
671 reliable for them. Moreover, it is predictable that spectral libraries will become more and
672 more complete with time, and that accurate prediction will become possible for all samples. In
673 such conditions, calcimetry could be increasingly replaced by MIRS for SIC analysis.

674

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691

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883 **Table 1.** Presentation of the sites of the target set.
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Town or village	Number of plots	Longitude and latitude	Elevation (m a.s.l.)	Soil type ^a	Mean clay content (%)	Mean sand content (%)	SIC range and mean (g kg ⁻¹)
Terrats	11	42° 36' 27" N, 02° 46' 14" E	135	Luvisol and Cambisol	27	41	0-3, 0
Lesquerde	19	42° 48' 01" N, 02° 31' 47" E	358	Arenosol	7	81	0-1, 0
Montagnac	21	43° 28' 50" N, 03° 29' 02" E	54	Calcisol	38	32	23-84, 50
Faugères	21	43° 33' 57" N, 03° 11' 19" E	284	Cambisol	16	52	0-12, 1
Aigues-Mortes	18	43° 34' 02" N, 04° 11' 33" E	5	Arenosol	7	88	17-25, 23
Vergèze	17	43° 44' 37" N, 04° 13' 14" E	32	Cambisol	36	18	2-50, 28
Jonquières Saint-Vincent	19	43° 49' 38" N, 04° 33' 48" E	37	Rhodic Luvisol	18	43	0-6, 1
Saint-Hippolyte- du-Fort	15	43° 57' 56" N, 03° 51' 28" E	170	Calcisol	42	22	7-86, 43
Saint-Victor- la-Coste	23	44° 03' 38" N, 04° 38' 29" E	143	Calcisol	16	54	1-43, 20
Whole target set	164	-	-	-	-	-	0-86, 19
National library	2178	-	-	-	-	-	0-104, 6

885 ^aIUSS Working Group WRB (2014).

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Table 2. Prediction results after global calibration without or with spiking.

Spiking samples and weight	Best pre-treatment	Calibration							Validation								
		N_{cal}^a	$Mean_{cal}^b$	SD_{cal}^b	LV^c	$SECV^d$	$R^2_{cv}^e$	RPD_{cv}^f	N_{val}^a	$Mean_{val}^b$	SD_{val}^b	SEP^d	bias	slope	$R^2_{val}^e$	RPD_{val}^f	
0	None	1155	2178	6.4	16.0	11	2.7	0.97	6.0	134	17.8	19.3	5.2	-2.8	1.10	0.96	3.7
10×1	D1	1155	10 spiking samples only	22.7	30.2	3	5.3	0.97	5.7	134	17.8	19.3	3.6	0.4	1.04	0.97	5.3
30×1	D1	205	30 spiking samples only	23.3	29.0	3	5.0	0.97	5.8	134	17.8	19.3	3.8	-0.6	1.10	0.97	5.1
10×1	D1	1155	2188	6.5	16.1	11	2.7	0.97	6.0	134	17.8	19.3	4.9	-2.5	1.09	0.96	3.9
10×10	D1	205	2278	7.1	17.1	13	2.9	0.97	5.9	134	17.8	19.3	3.9	-1.2	1.07	0.97	5.0
10×40	D1	205	2578	8.9	19.5	14	3.2	0.97	6.1	134	17.8	19.3	3.3	-0.1	1.04	0.97	5.9
10×50	D1	1155	2678	9.4	20.0	14	3.2	0.97	6.3	134	17.8	19.3	3.3	0.2	1.03	0.97	5.9
30×10	None	1155	2478	8.4	18.8	11	2.8	0.98	6.7	134	17.8	19.3	4.0	-1.6	1.08	0.97	4.8
30×200	D1	1155	8178	18.8	26.8	14	3.1	0.99	8.7	134	17.8	19.3	3.0	-0.2	1.04	0.98	6.4

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^a N_{cal} and N_{val} are the number of calibration samples and of validation samples predicted, respectively.

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^b $Mean_{cal}$, SD_{cal} , $Mean_{val}$ and SD_{val} are the mean and standard deviation over the calibration and validation set, respectively (in g kg⁻¹).

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^c LV is the number of PLS latent variables (or terms).

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^d SECV and SEP are the standard error of cross-validation and prediction, respectively (in g kg⁻¹).

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^e R^2_{cv} and R^2_{val} are the determination coefficient for cross-validation and validation, respectively.

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^f RPD_{cv} is the ratio of SD_{cal} to SECV, RPD_{val} is the ratio of SD_{val} to SEP.

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Table 3. Prediction results after local calibration without or with spiking (and comparison with global calibration on the same validation samples).

Spiking samples and weight	Procedure	Best pre-treatment	Validation							
			$N_{\text{val}}^{\text{a}}$	$\text{Mean}_{\text{val}}^{\text{b}}$	$\text{SD}_{\text{val}}^{\text{b}}$	SEP^{c}	bias	slope	$R^2_{\text{val}}^{\text{d}}$	$\text{RPD}_{\text{val}}^{\text{e}}$
0	Local calibration using 4 to possibly 2178 calibration neighbours	None001	133	17.8	19.4	4.8	-2.0	1.07	0.95	4.1
0	Global calibration with 2178 samples and the same 133 validation samples	None1155	133	17.8	19.4	4.9	-2.7	1.09	0.96	4.0
0	Local calibration using 50 to possibly 2178 calibration neighbours	None001	115	17.0	20.7	2.7	-0.8	1.02	0.99	7.6
0	Global calibration with 2178 samples and the same 115 validation samples	None1155	115	17.0	20.7	3.3	-1.6	1.05	0.98	6.4
10×1	Local calibration using 4 to possibly 2188 calibration neighbours	SNV001	134	17.8	19.3	2.7	-0.0	1.01	0.98	7.3
10×1	Local calibration using 50 to possibly 2188 calibration neighbours	SNVD001	115	17.0	20.7	2.3	-0.5	1.03	0.99	8.9
10×1	Global calibration with 2188 samples and the same 115 validation samples	SNV1155	115	17.0	20.7	2.7	-0.2	1.00	0.98	7.8
10×10	Local calibration using 4 to possibly 2278 calibration neighbours	SNV001	134	17.8	19.3	2.7	-0.5	1.01	0.98	7.2
10×10	Local calibration using 50 to possibly 2278 calibration neighbours	SNVD001	115	17.0	20.7	2.3	-0.4	1.03	0.99	9.1
10×10	Global calibration with 2278 samples and the same 115 validation samples	D1205	115	17.0	20.7	2.7	-0.5	1.04	0.98	7.5
10×50	Local calibration using 4 to possibly 2678 calibration neighbours	SNV001	134	17.8	19.3	2.7	-0.5	1.01	0.98	7.2
10×50	Local calibration using 50 to possibly 2678 calibration neighbours	D001	134	17.8	19.3	2.7	-0.8	1.03	0.98	6.8
10×50	Global calibration with 2678 samples and all 134 validation samples	D1155	134	17.8	19.3	3.3	0.2	1.03	0.97	5.9
30×1	Local calibration using 4 to possibly 2208 calibration neighbours	SNV001	134	17.8	19.3	2.7	-0.5	1.03	0.98	7.2
30×1	Local calibration using 50 to possibly 2208 calibration neighbours	SNVD001	115	17.0	20.7	2.3	-0.5	1.03	0.99	9.2
30×1	Global calibration with 2208 samples and the same 115 validation samples	SNV1155	115	17.0	20.7	2.6	-0.4	1.00	0.98	7.9
30×10	Local calibration using 4 to possibly 2478 calibration neighbours	SNV001	134	17.8	19.3	2.7	-0.5	1.04	0.98	7.2
30×10	Local calibration using 50 to possibly 2478 calibration neighbours	SNV001	116	17.0	20.6	2.3	-0.5	1.04	0.99	8.8
30×10	Global calibration with 2478 samples and the same 116 validation samples	SNVD1155	116	17.0	20.6	2.5	-0.2	1.03	0.99	8.2

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^a N_{val} is the number of validation samples predicted.

^b Mean_{val} and SD_{val} are the mean and standard deviation over the validation set (in g kg^{-1}).

^c SEP is the standard error of prediction (in g kg^{-1}).

^d R^2_{val} is the determination coefficient for validation.

^e RPD_{val} is the ratio of SD_{val} to SEP.

Figure captions

Fig. 1. Distribution of soil inorganic carbon (SIC) content in the national library (2178 samples, 0–30 cm depth) and in the target set (164 samples, 0–15 cm depth).

Fig. 2. Comparison between reference measurements and MIRS predictions of SIC content on the validation set using (a) global calibration without spiking, (b) local calibration using 4 to 2188 spectral neighbours from the national library enriched with 10 spiking samples without extra-weighting, (c) local calibration using 50 to 2278 neighbours from the national library enriched with 10 spiking samples extra-weighted 10 times, and (d) global calibration with similar spiking conditions and the same validation samples.

Fig. 3. Projection of the validation samples on the PCA built with the national library.

Fig. 4. Standard error of prediction (SEP) of SIC content when performing global MIRS calibration with the national library and spiking samples, as affected by the number and weight of spiking samples, or with spiking samples only (i.e. without the national library).

Fig. 5 Distribution of validation samples according to their number of spectral neighbours in the calibration set (a) without spiking; (b) with 10 spiking samples replicated 1 time, (c) replicated 10 times, and (d) replicated 50 times; and (e) with 30 spiking samples replicated 1 time, (f) replicated 10 times, and (g) replicated 50 times.

Fig. 1. Distribution of soil inorganic carbon (SIC) content in the national library (2178 samples, 0–30 cm depth) and in the target set (164 samples, 0–15 cm depth).

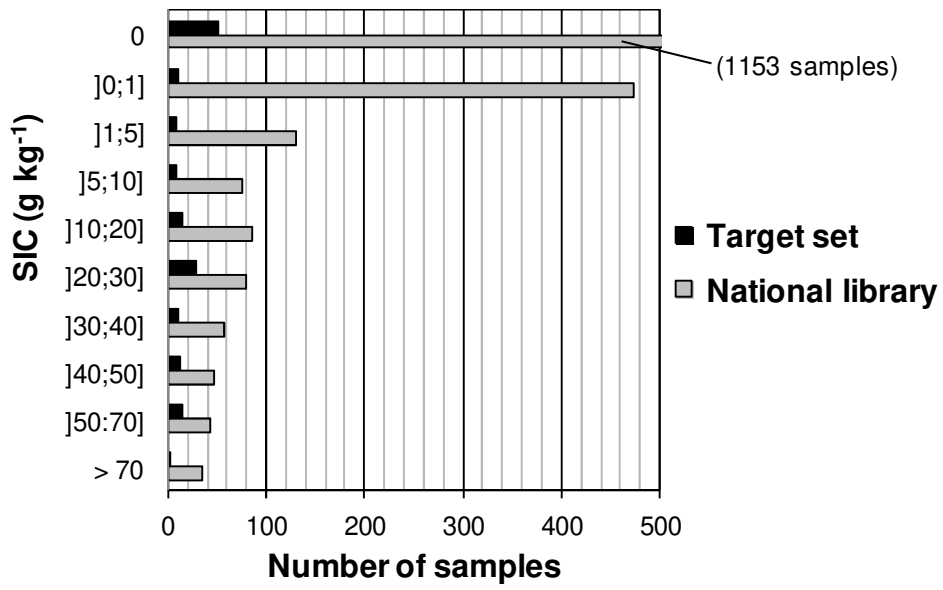


Fig. 2. Comparison between reference measurements and MIRS predictions of SIC content on the validation set using (a) global calibration without spiking, (b) local calibration using 4 to 2188 spectral neighbours from the national library enriched with 10 spiking samples without extra-weighting, (c) local calibration using 50 to 2278 neighbours from the national library enriched with 10 spiking samples extra-weighted 10 times, and (d) global calibration with similar spiking conditions and the same validation samples.

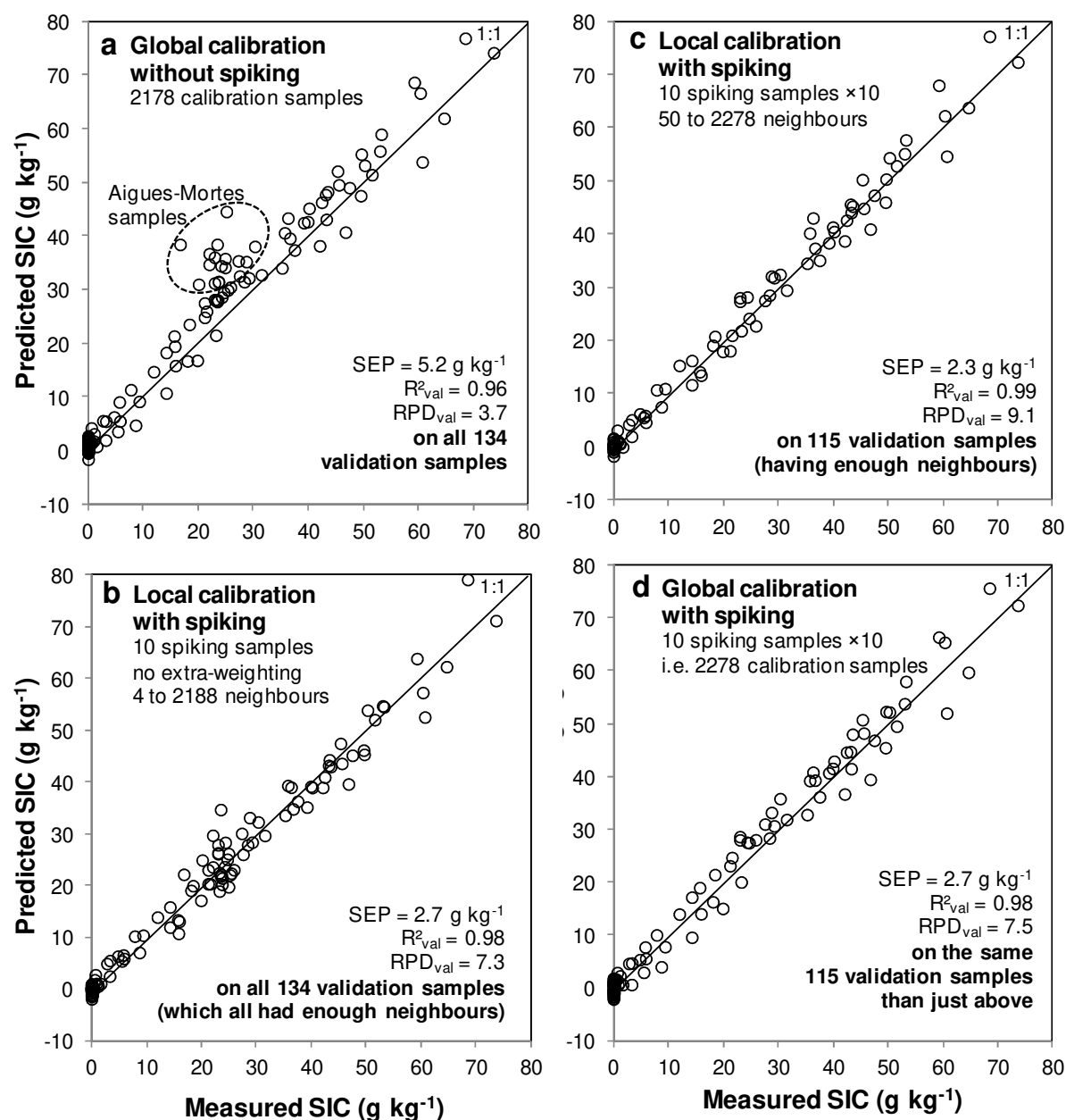


Fig.3. Projection of the validation samples on the PCA built with the national library.

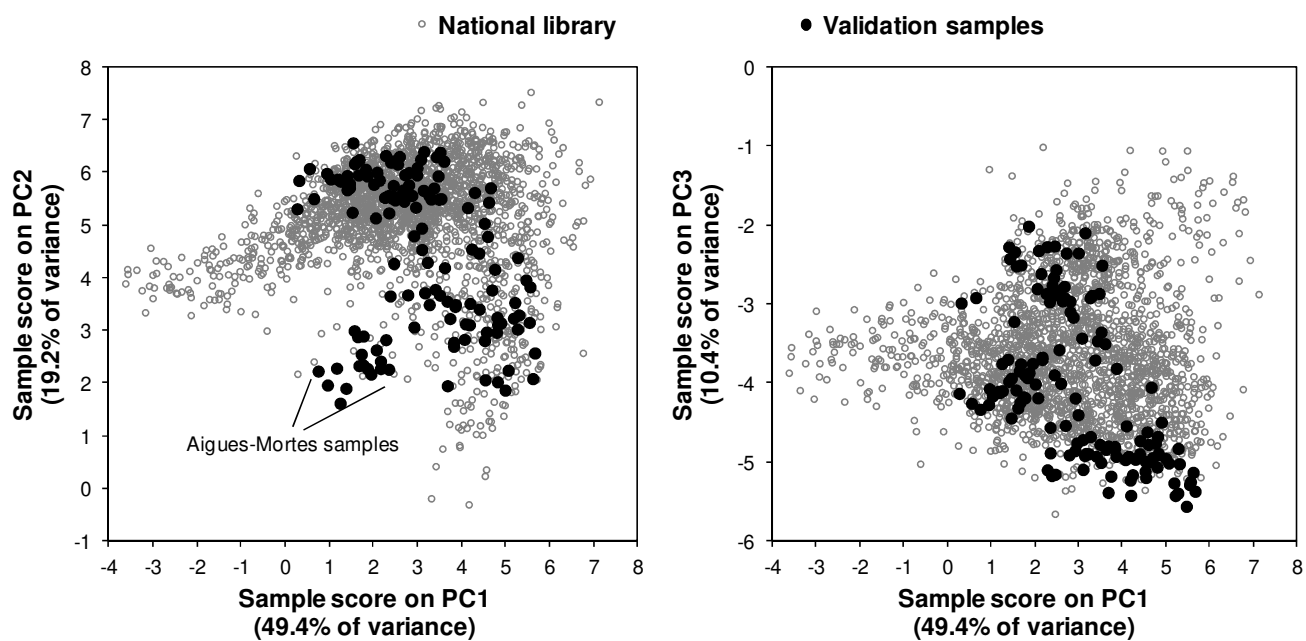


Fig. 4. Standard error of prediction (SEP) of SIC content when performing global MIRS calibration with the national library and spiking samples, as affected by the number and weight of spiking samples, or with spiking samples only (i.e. without the national library).

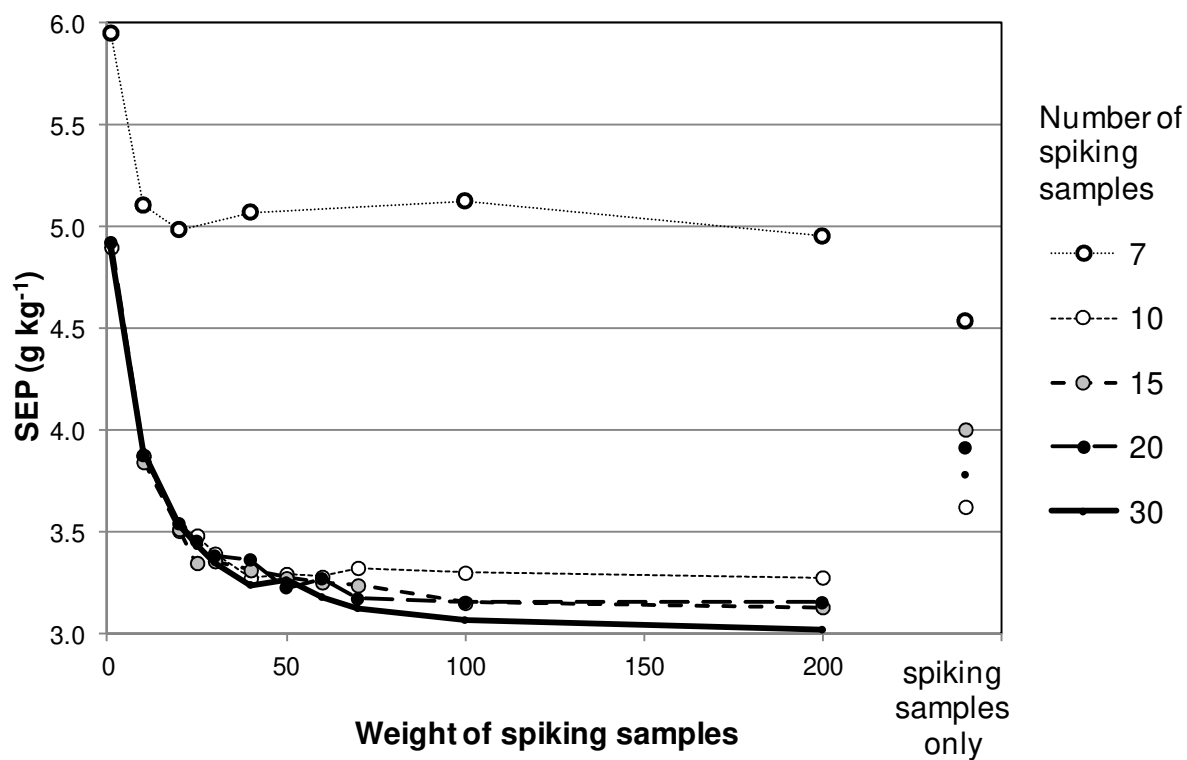


Fig. 5. Distribution of validation samples according to their number of spectral neighbours in the calibration set (a) without spiking; (b) with 10 spiking samples replicated 1 time, (c) replicated 10 times, and (d) replicated 50 times; and (e) with 30 spiking samples replicated 1 time, (f) replicated 10 times, and (g) replicated 50 times.

