

1 Estimating uncertainty in pooled proxy time-series, including stable isotopes in tree-
2 rings

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27

28 **Abstract** (198 words)

29

30 Stable carbon isotope time-series ($\delta^{13}\text{C}$) from tree-rings are capable of providing

31 valuable palaeoclimatic information, but analysis of individual tree-rings is time

32 consuming and expensive. Pooling material from several tree-rings prior to isotopic

33 analysis reduces costs, but does not allow the magnitude of uncertainty in the mean

34 $\delta^{13}\text{C}$ chronology to be calculated unless the pool is broken and each tree-ring

35 measured individually at regular intervals. Here we use a comparison of pooled and

36 mean individual (the arithmetic mean of isotopic data from tree series measured

37 individually) $\delta^{13}\text{C}$ records to determine whether the true error structure of the time

38 series is better captured by using the overall mean error estimate for the entire time

39 series or by linear interpolation between the equally spaced measurements. We
40 conclude that where autocorrelation exists within the error structure of a chronology,
41 annual estimates of 95% confidence intervals, developed through linear interpolation
42 at 5-year or 10-year intervals, are preferable to using the overall mean uncertainty.
43 The method outlined increases the viability of pooled $\delta^{13}\text{C}$ records for palaeoclimatic
44 research by retaining error structure whilst reducing analytical time and costs. The
45 method is applied here using tree-ring data, but could theoretically be applied to any
46 well-replicated time-series.

47

48 **Keywords:** tree rings; climate reconstruction; stable isotopes; pooling; proxy time-
49 series; error

50

51 **1. Introduction**

52

53 The stable carbon isotope composition ($\delta^{13}\text{C}$) of tree-ring cellulose has been used
54 to produce valuable palaeoclimate reconstructions from many trees species,
55 growing within a range of climatic regimes (McCarroll and Pawellek, 2001;
56 Masson-Delmotte et al., 2005; Poussart and Schrag, 2005; Kirilyanov et al., 2008;
57 Kress et al., 2010; Gagen et al., 2011; McCarroll et al., 2011; Seftigen et al. In
58 press). The construction of well-replicated, multi-centennial chronologies through
59 the analysis of individual tree-rings (Gagen et al., 2007; Young et al., 2010)
60 permits annual assessment of isotopic variability, allowing confidence intervals to
61 be placed around the mean isotope value and the resulting climate reconstructions
62 (McCarroll and Pawellek, 1998; McCarroll and Loader, 2004, 2006). This
63 method also allows identification and reduction of non-climatic trends in

64 individual time-series (Loader et al. 2007), such as juvenile effects (Gagen et al.
65 2008) and physiological responses to increasing atmospheric CO₂ concentrations
66 (Gagen et al., 2007, 2011).

67

68 The construction of long stable isotope chronologies comprising individual tree
69 series is, however, time consuming and relatively expensive. Researchers have
70 attempted to overcome these limitations by pooling (combining) the material from
71 sampled trees for each year prior to isotopic analysis. Pooling of raw wood, prior
72 to the isolation of α -cellulose, is the most commonly adopted approach, leading to
73 a large reduction in the number of samples that have to be prepared, and this
74 method has been successfully employed to extract climatic information from tree-
75 rings (Rebetez et al., 2003; Treydte et al., 2007; Loader et al., 2008; Tardif et al.,
76 2008; Hiltavuori et al., 2009; Rinne et al. 2010). Although the same weighting of
77 each constituent tree is only guaranteed by pooling of equal amounts of well
78 homogenised (powdered) raw wood, it has been reported that the bias from
79 differing mass contributions of raw wood towards a pool appears to be negligible
80 (Borella et al., 1998; Leavitt, 2008).

81

82 Pooled $\delta^{13}\text{C}$ chronologies can also be constructed by isolating the α -cellulose from
83 individual tree-rings and creating an annual pool using equal masses from each
84 constituent tree. Whilst this approach results in increased sample preparation
85 time, it requires the same number of isotopic analyses as the methods outlined
86 above and permits retention of sample material from individual tree-rings if
87 desired. Combination of equal masses of isotopically homogenous α -cellulose
88 ensured an equal weighting of each tree within the chronology. This method of

89 pooling should, therefore, produce a mean isotope chronology equivalent to that
90 obtained by calculating the mean $\delta^{13}\text{C}$ of individual trees within a chronology and
91 is the approach adopted in this study.

92

93 A major limitation of pooled chronologies is the inability to calculate the standard
94 deviation between the constituent trees and therefore, to assign confidence limits
95 around $\delta^{13}\text{C}$ values. Quantification of uncertainty is an essential requirement for
96 climate reconstructions (Jansen et al., 2007), particularly if data are to be used to
97 test the veracity of climate model retrodictions (McCarroll, 2010). A potential
98 solution to this problem may be to split the pooled chronology at regular intervals
99 (eg: every 5th, 10th, 15th or 20th year) and to analyse each tree individually to
100 calculate the standard deviation. This method was successfully applied to a
101 pooled $\delta^{13}\text{C}$ chronology constructed using white spruce (*Picea glauca* (Moench)
102 Voss) in subarctic Manitoba, Canada (Tardif et al., 2008), and allowed
103 quantification of uncertainty (confidence intervals around the mean) for every
104 fifth year.

105

106 Given a measure of uncertainty at regular intervals, there are two options for
107 extrapolating those values so that they apply to the whole chronology: either apply
108 the overall mean uncertainty to every annual value, or interpolate between the
109 values in series-order to capture temporal changes in the error structure. This
110 study aims to test whether interpolation provides a better estimate of the true error
111 structure than simply using the mean.

112

113

114 **2. Methods**

115

116 Long-lived Scots pine (*Pinus sylvestris* L.) trees growing at Southern Glens (057°N;
117 005°W), western Highlands of Scotland, were sampled using a 10mm Haglöf
118 increment borer. Cores from twenty one trees, growing on north-facing slopes (0-30°
119 inclination) between 80 and 380m.a.s.l., were air dried for two weeks, sanded using
120 progressively finer grades of abrasive paper and crossdated using a binocular
121 microscope and Velmex measuring stage interfaced with a computer. TSAPWin™
122 and COFECHA (Holmes, 1999; Grissino-Mayer, 2001; Rinn, 2003) were used to
123 absolutely date each tree series and a detrended ring-width chronology for Southern
124 Glens was constructed using the computer program ARSTAN (Holmes, 1983). Under
125 the magnification of a binocular microscope, annual whole-ring (early- plus late-
126 wood) increments were cut into thin slivers using a scalpel. α -cellulose was isolated
127 from annual raw wood samples using standard techniques (Loader et al., 1997; Rinne
128 et al., 2005). In order to produce isotopically homogenous sample material, the
129 resulting α -cellulose was placed into 2mm micro-centrifuge tubes with deionised
130 water and homogenised using a Hielscher UP 200S ultrasonic probe (Loader et al.,
131 2008; Laumer et al., 2009). Following freeze drying (ModulyoD Thermo Savant), an
132 equal quantity of alpha-cellulose (0.5mg \pm 0.05mg) was removed from each annual
133 individual cellulose sample and combined to produce an annual pool. Pooled samples
134 were then placed in deionised water and homogenised again, before being freeze
135 dried. Between 0.30 and 0.35mg of alpha-cellulose were weighed into tin foil
136 capsules, crimped and placed into a sample tray. Stable carbon isotope analysis was
137 conducted on ANCA and SerCon GSL elemental analysers (1000°C combustion
138 temperature), interfaced with 20-20 Isotope Ratio Mass Spectrometers (IRMS) (PDZ-

139 Europa). Stable isotope results are expressed as per mille (‰), relative to the
140 international standard Vienna Pee Dee Belemnite (VPDB) standard (Coplen, 1995,
141 2006).

142

143 The $\delta^{13}\text{C}$ chronologies presented here were produced for the purpose of palaeoclimate
144 reconstruction, so they have been corrected both for changes in the isotopic ratio of
145 atmospheric carbon dioxide, by simple addition using (and extrapolating) the values
146 provided by McCarroll and Loader (2004), and for changes in intrinsic water-use
147 efficiency in response to increased atmospheric carbon dioxide (CO_2) concentration,
148 using the Pre-INDustrial (PIN) correction proposed by McCarroll et al. (2009). Both
149 pooled and mean individual $\delta^{13}\text{C}$ chronologies comprise a minimum of seven trees
150 between AD 1650 and 2007. The pooled record incorporates one specimen which
151 was omitted from the mean individual chronology. Tree 50B (AD 1715-1820) does
152 not demonstrate a common signal with the mean individual chronology ($r = -0.10$,
153 AD 1715-1820). This is beneficial, as it provides the opportunity to assess whether
154 incorporation of a “noisy” series significantly affects the isotopic signal of the pooled
155 $\delta^{13}\text{C}$ chronology at this level of replication.

156

157 Annual 95% confidence intervals were calculated for the mean individual chronology
158 using the equation given below, where n is the number of trees in a given year, SD is
159 the standard deviation of those trees, and t is the t distribution value for n in that year.

160

161
$$95\% \text{ confidence interval} = t \cdot \left(\frac{SD}{\sqrt{n}} \right)$$

162

163 Annual confidence intervals were calculated through linear interpolation between
164 equidistant pairs of data points (observed confidence intervals). Interpolated datasets
165 were developed for intervals of 5, 10, 15 and 20 years and for all possible
166 combinations of years within these categories (e.g. AD 1650, 1651....1654 for 5-year
167 intervals). Therefore, the sensitivity of interpolation to choice of years can be
168 assessed. The reduction of error (RE) statistic (National Research Council 2007) is
169 used to assess whether the interpolated values have greater predictive skill than
170 simply using the overall mean. This should permit the identification of an optimum
171 sampling resolution, whereby the cost of interpolation remains advantageous and still
172 retains the error structure of a time series better than the mean. The equation for RE
173 is given below, but the terms of the equation have been adjusted for this study.

174

175

$$\text{RE} = 1 - \left[\frac{\sum_{i=1}^n (x_i - \hat{x}_i)^2}{\sum_{i=1}^n (x_i - \bar{x}_c)^2} \right],$$

176

177 where x_i and \hat{x}_i are the observed and interpolated 95% confidence intervals in year i
178 and \bar{x}_c is the mean of the observed confidence intervals for the entire time series.

179 Years which were used in the interpolation equation were omitted prior to the
180 calculation of RE.

181

182 **3. Results**

183

184 **Pooled and mean individual isotope chronologies**

185

186 A strong relationship ($r = 0.85$; $P < 0.01$, $n = 358$) exists between mean individual and
187 pooled $\delta^{13}\text{C}$ chronologies for Southern Glens from AD 1650-2007 (Figure 1). Both
188 chronologies have very similar mean values (pooled = -24.14‰ ; mean individual =
189 -24.18‰) and variances (pooled = 0.08‰ ; mean individual = 0.09‰) throughout this
190 period and exhibit near identical trends at high and low frequency timescales. An F -
191 test, followed by a z -test (Fowler et al., 1998), demonstrates that there are no
192 significant differences ($P = 0.05$) between the variances ($F = 1.09$; F -critical = 1.19)
193 or means ($z = 1.77$; z -critical = 1.95) of the pooled and mean individual $\delta^{13}\text{C}$
194 chronologies from AD 1650-2007.

195

196 *Figure 1*

197

198 The incorporation into the pooled chronology of a noisy series (tree 50B, AD 1715-
199 1820) does not result in any significant difference in either variance ($F = 1.29$; F
200 critical = 1.38) or mean value ($z = 0.19$; z critical = 1.95) between the two
201 chronologies during this period and they remain highly correlated ($r = 0.74$; $P < 0.01$).

202

203 **Assessment of error structure**

204

205 Normally, one would assume that the inter-annual error is independent. However in
206 many proxy timeseries based upon biological systems this may not always be the
207 case. The non-parametric ‘number of runs test’ demonstrates that the true uncertainty
208 values, at annual resolution, are not randomly arranged through time ($z = -8.54$,
209 $P < 0.0001$, $n = 358$). This is confirmed by the presence of significant autocorrelation
210 within the time series ($r_1 = 0.61$, $P < 0.01$, $n = 357$). The error structure of the annual

211 time series is thus far from white noise, and the uncertainty for a given year is not
212 independent of the uncertainty for the surrounding years. This means that there is at
213 least the potential for interpolation to capture some of the real error structure of the
214 time series.

215

216 The value of interpolation is assessed here by using the results from the individual
217 tree-rings to produce artificial pooled chronologies. As values are available for every
218 tree in every year, we can therefore assess the effect of varying the interval at which
219 the artificial pool is broken and also assess the sensitivity of the results to the
220 individual years on which the interpolation is based.

221

222 When the pool is broken every five years, it is clear that linear interpolation follows
223 the true error structure of the data very well (Figure 2A). Almost 74% of the
224 interpolated values fall within 0.1‰ of the true values, compared with only 59% when
225 the overall mean is used. The superiority of interpolation over the mean is
226 demonstrated by a positive RE value (0.30). Splitting the pool every ten years still
227 produces interpolated values that follow the true error structure very well, with almost
228 70% of interpolated values falling within 0.1‰ and a positive RE value (0.20). A 10-
229 year interpolation interval thus provides an 11.7% increase in data that are distributed
230 within 0.1‰ of observed values, relative to using the mean (Table 1).

231

232 As the sampling interval exceeds 10 years, more individual series fail the RE test
233 because the coincidence of interpolation intervals with extreme values in the error
234 structure affect a greater proportion of the dataset. Interpolation is not a “perfect”
235 solution; this is highlighted by the 10-year interpolated series starting in AD 1656,

236 which is influenced by a relatively extreme isotopic value. Removal of a single
237 extremity (AD 1766) results in an increase in RE from -0.10 to 0.20. Even with a
238 spacing of 15 years, interpolation performs better than the overall average. A
239 reasonable guide to the likely advantage of using interpolation is provided by the first
240 order autocorrelation of the error estimates (Table 1), since as this approaches zero
241 there can be no advantage over using the overall average error.

242

243 Figure 2 shows examples of 5, 10, 15 and 20 year interpolated series, starting with the
244 year AD 1650. The histograms and line graphs (A-D) confirm that interpolation is
245 most effective when the distance between sampling intervals is equal to, or less than,
246 10 years. Beyond this length of interval, less of the trend in the error structure is
247 captured.

248

249 *Table 1*

250

251 *Figure 2*

252

253 Addition of the observed, interpolated (10-year intervals starting AD 1650) and mean
254 (10-year intervals) 95% confidence intervals to the pooled $\delta^{13}\text{C}$ chronology (Southern
255 Glens) appear to yield very similar levels of uncertainty (Figure 3, graph C), a result
256 of the larger variance in the $\delta^{13}\text{C}$ data (0.08‰), relative to the error time series
257 (0.01‰). A section of the chronology presented in graphs A and B (Figure 3)
258 demonstrate how interpolation at 10-year intervals is more capable of retaining the
259 error structure of the chronology, relative to the mean.

260

261 *Figure 3*

262

263 **4. Discussion and Conclusions**

264

265 Prior to this study, a number of pooling methods had been proposed for stable isotope
266 dendroclimatology (Leavitt and Long, 1984; Boettger and Friedrich, 2009), but there
267 was a lack of knowledge regarding the relationship between pooled and mean
268 individual chronologies and no practical approach to quantifying uncertainty. The
269 large-scale comparison presented here confirms that equally-weighted, pooled (α -
270 cellulose) $\delta^{13}\text{C}$ chronologies developed from Scots pine trees growing at Southern
271 Glens are equivalent to chronologies constructed by taking the mean $\delta^{13}\text{C}$ of
272 individual trees. Therefore, this method is presented as a viable means of constructing
273 a pooled $\delta^{13}\text{C}$ chronology, with a significant reduction (typically 60-80%) in the
274 number of required analyses. Pooling raw wood prior to chemical treatment to isolate
275 α -cellulose would result in further savings.

276

277 Splitting the pool at regular intervals, and measuring the isotopic ratio of each ring
278 individually, provides equally spaced measurements of between-tree variability and
279 therefore of the uncertainty in the estimate of the mean. By using a well replicated
280 $\delta^{13}\text{C}$ chronology, where every tree-ring was measured individually, we were able to
281 simulate pooled chronologies where the position and spacing of the splitting could be
282 varied. The error structure of the time series, at annual resolution, was far from white
283 noise, showing significant autocorrelation. Consequently, we found that given a
284 spacing of 5 or 10 years, linear interpolation between adjacent split pool

285 measurements provided a better approximation to the true error structure than simply
286 applying the overall mean uncertainty to every year.

287

288 The optimum period to split a pooled chronology is a function of the degree of
289 autocorrelation in the error structure (not the isotope values) and will therefore vary
290 with each data set. Appropriately replicated tree-ring stable isotope chronologies
291 constructed using a varying numbers of trees, which enter and leave the chronology at
292 different times, are very likely to have error structures that differ significantly from
293 white noise and in these situations, the interpolation approach would lead to an
294 improvement in error estimation. When a chronology is produced by pooling alone it
295 is not possible to measure the true error structure, but our results indicate that the first
296 order autocorrelation observed in the split pool samples (Table 1) provides a
297 reasonable guide to the relative merits of either interpolating the errors or just using
298 the overall mean. When a pooled sampling approach is being applied to a new site,
299 where the error structure of the data is completely unknown, we suggest that splitting
300 the pool every ten years and interpolating the uncertainty is a reasonable compromise
301 between the cost of analyses and the need to quantify the uncertainty around the
302 mean.

303

304 This research has increased the viability of pooled $\delta^{13}\text{C}$ chronologies through
305 proposing and testing a method of estimating annual confidence intervals. Whilst this
306 approach may not always identify individual years where confidence intervals vary
307 significantly from adjacent years, it has been identified as an effective approach for
308 assessing and maintaining temporal variability in uncertainty around a chronology.

309 The method outlined significantly reduces the analytical time and expense of

310 constructing $\delta^{13}\text{C}$ chronologies and is likely to be applicable to other stable isotope
311 (oxygen and hydrogen) series from tree-rings or to other well-replicated proxy
312 records.

313

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323

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Figure

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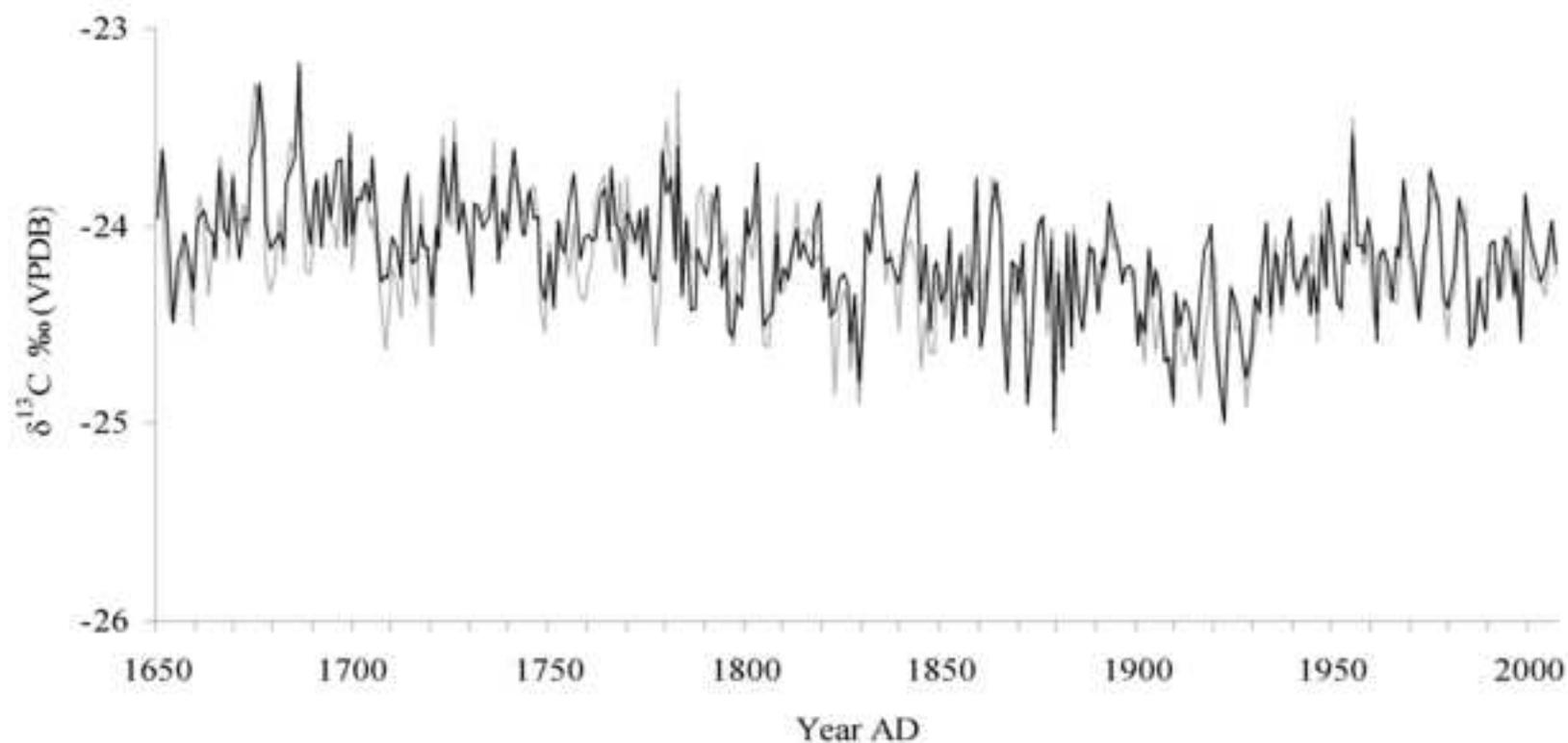


Figure 1: Comparison of pooled (black line) and mean individual (grey line) $\delta^{13}\text{C}$ chronologies for Southern Glens between AD 1650 and 2007. The two chronologies are highly correlated over this time period ($r = 0.85$, $P < 0.01$, $n = 358$).

Figure

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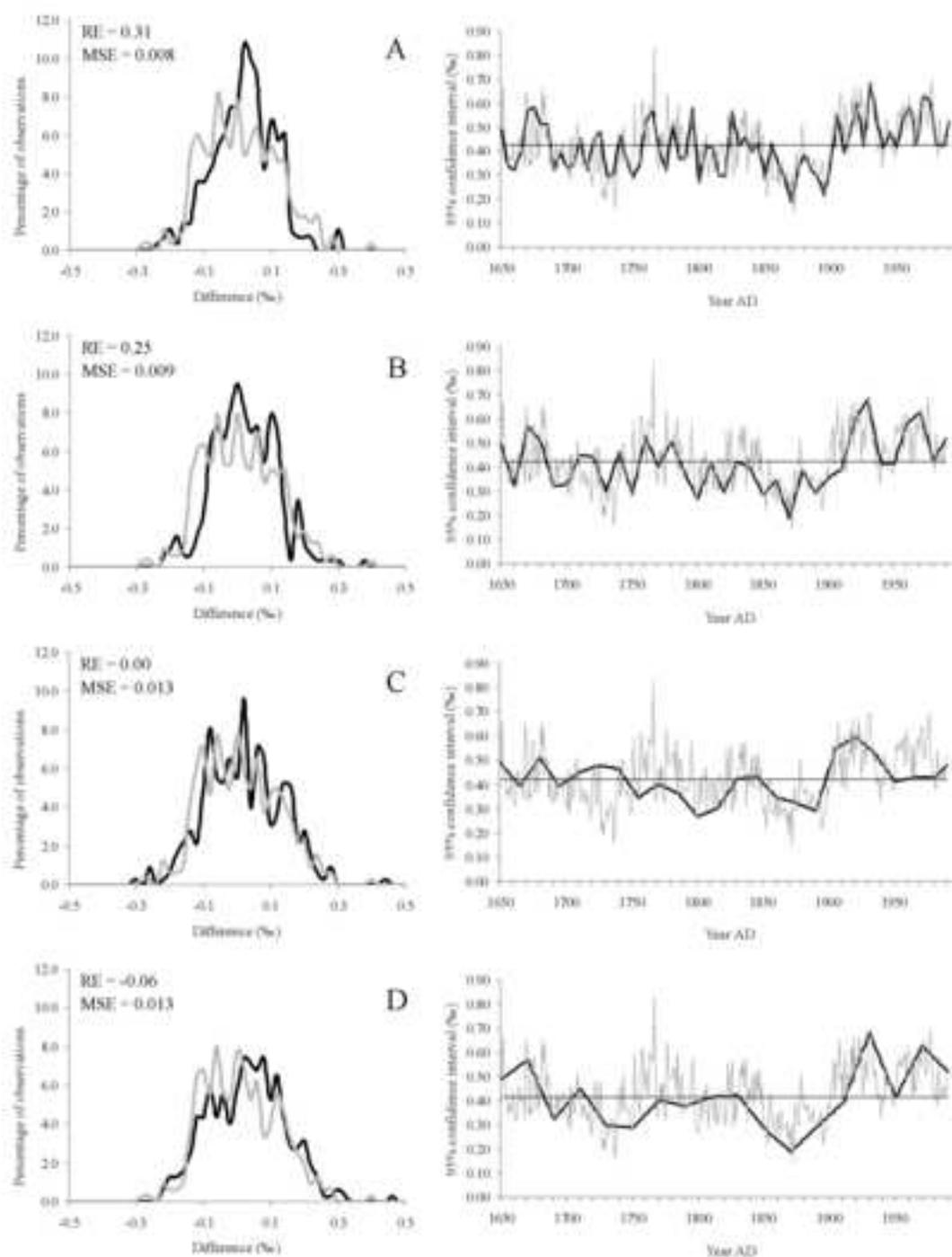


Figure 2: Histograms showing the difference (%) between observed minus interpolated confidence intervals (black line) and observed minus mean confidence intervals (grey line). The line graphs show the observed confidence intervals (grey line) and the interpolated confidence intervals (black line) for the Southern Glens chronology. The horizontal black line represents the mean observed uncertainty for the interpolation intervals throughout the chronology. Comparisons of observed and interpolated data are presented (starting with the year AD 1650) for 5 (A), 10 (B), 15 (C) and 20-year intervals (D).

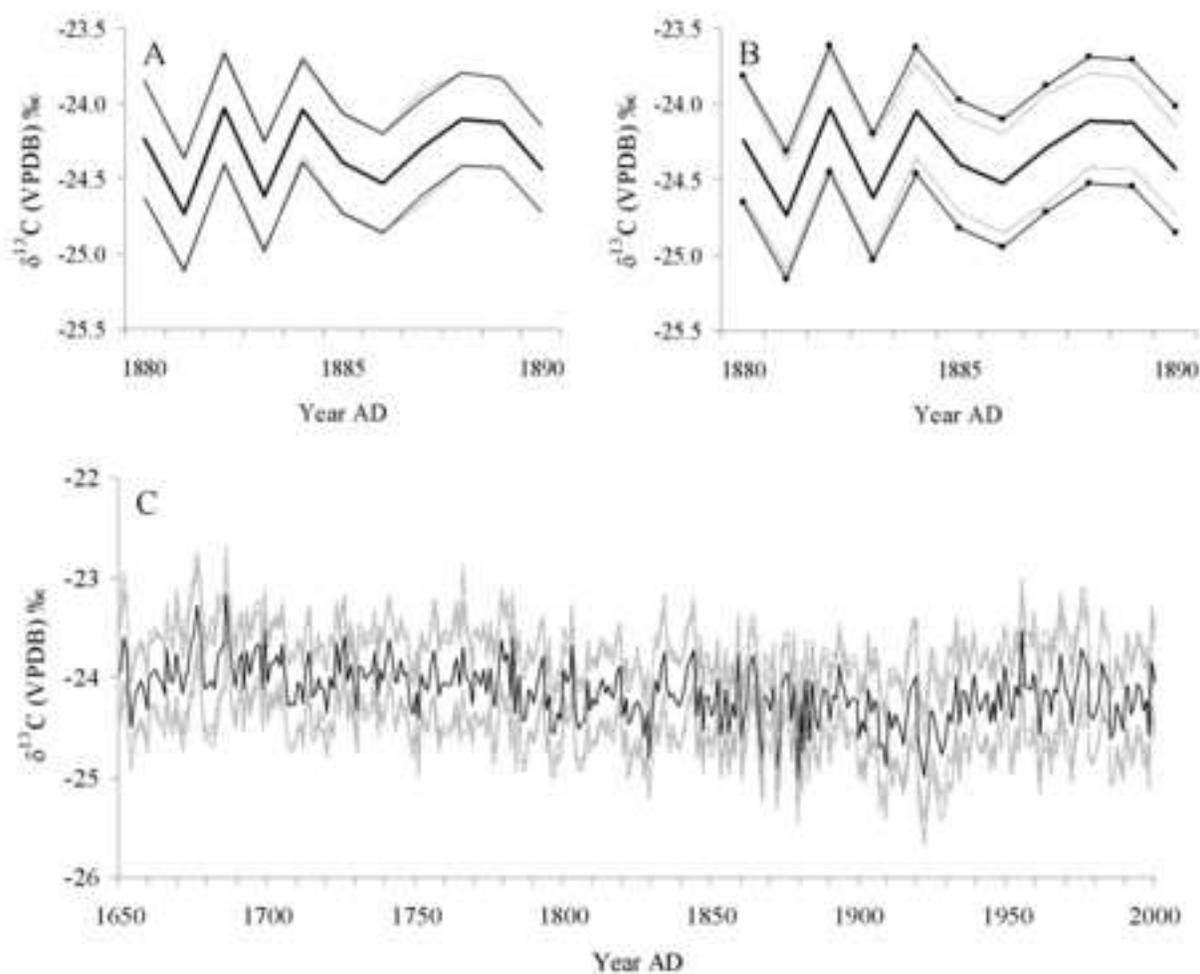
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Figure 3: The addition of 95% confidence intervals to the pooled $\delta^{13}\text{C}$ chronology (Southern Glens). The difference between observed (grey line) and interpolated (black line) (A) and observed and mean confidence intervals (black circles) (B) is shown for a short section (AD 1880-1890) of the chronology (thick black line). The differences between observed, interpolated and mean confidence intervals (grey lines) appear insignificant when applied to the pooled $\delta^{13}\text{C}$ chronology (black line) (C), a result of higher variance in $\delta^{13}\text{C}$ values (0.08‰) relative to the observed error structure (0.01‰) between AD 1650 and 2007.

Table 1: Comparison of mean first order autocorrelation, mean MSE, mean RE and the relationship between observed, interpolated and mean confidence intervals for 5, 10, 15 and 20-year intervals.

| | Interpolation interval | | | |
|-----------------------------|------------------------|----------|----------|----------|
| | 5-years | 10-years | 15-years | 20-years |
| Observed - interpolated | | | | |
| Mean MSE | 0.008 | 0.010 | 0.011 | 0.011 |
| % of data within 0.1‰ | 73.7 | 69.9 | 67.3 | 65.8 |
| % of data within 0.2‰ | 96.7 | 95.5 | 94.7 | 79.5 |
| Observed - mean | | | | |
| Mean MSE | 0.012 | 0.012 | 0.012 | 0.012 |
| % of data within 0.1‰ | 58.8 | 58.2 | 58.4 | 58.0 |
| % of data within 0.2‰ | 94.1 | 93.9 | 94.0 | 93.9 |
| | | | | |
| First order autocorrelation | 0.41 | 0.26 | 0.20 | 0.08 |
| Mean RE | 0.30 | 0.20 | 0.11 | 0.09 |