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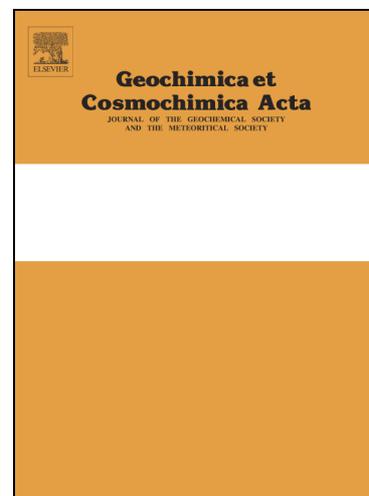
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1 **A hierarchical Bayesian approach to the classification of C₃ and C₄ grass pollen based on**
2 **SPIRAL $\delta^{13}\text{C}$ data**

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24 **Abstract**

25 Differentiating C₃ and C₄ grass pollen in the paleorecord is difficult because of their
26 morphological similarity. Using a spooling wire microcombustion device interfaced with an
27 isotope ratio mass spectrometer, Single Pollen Isotope Ratio AnaLysis (SPIRAL) enables
28 classification of grass pollen as C₃ or C₄ based upon $\delta^{13}\text{C}$ values. To address several limitations
29 of this novel technique, we expanded an existing SPIRAL training dataset of pollen $\delta^{13}\text{C}$ data
30 from 8 to 31 grass species. For field validation, we analyzed $\delta^{13}\text{C}$ of individual grains of grass
31 pollen from the surface sediments of 15 lakes in Africa and Australia, added these results to a
32 prior dataset of 10 lakes from North America, and compared C₄-pollen abundance in surface
33 sediments with C₄-grass abundance on the surrounding landscape. We also developed and tested
34 a hierarchical Bayesian model to estimate the relative abundance of C₃- and C₄-grass pollen in
35 unknown samples, including an estimation of the likelihood that either pollen type is present in a
36 sample. The mean (\pm SD) $\delta^{13}\text{C}$ values for the C₃ and C₄ grasses in the training dataset were -29.6
37 $\pm 9.5\text{‰}$ and $-13.8 \pm 9.5\text{‰}$, respectively. Across a range of % C₄ in samples of known
38 composition, the average bias of the Bayesian model was <3% for C₄ in samples of at least 50
39 grains, indicating that the model accurately predicted the relative abundance of C₄ grass pollen.
40 The hierarchical framework of the model resulted in less bias than a previous threshold-based
41 C₃/C₄ classification method, especially near the high or low extremes of C₄ abundance. In
42 addition, the percent of C₄ grass pollen in surface-sediment samples estimated using the model
43 was strongly related to the abundance of C₄ grasses on the landscape ($n= 24$, $p< 0.001$, $r^2= 0.65$).
44 These results improve $\delta^{13}\text{C}$ -based quantitative reconstructions of grass community composition
45 in the paleorecord and demonstrate the utility of the Bayesian framework to aid the interpretation
46 of stable isotope data.

47 **1. INTRODUCTION**

48 Knowledge of biotic responses to past climatic variability is important for anticipating
49 future change (Flessa et al., 2005). Fossil pollen assemblages are a valuable indicator of
50 spatiotemporal variation in plant community composition on the landscape (Williams et al.,
51 2004). However, grass (Poaceae) pollen is typically morphologically indistinct below the family
52 level (Fægri et al., 1989), rendering pollen analysis a blunt instrument for investigating past
53 changes in grassland ecosystems. This problem hampers our understanding of the ecology and
54 evolution of grasslands, which today cover a major portion of Earth's land surface and regulate
55 key biogeochemical cycles (Saugier and Roy, 2000).

56 Carbon isotopic analysis of grass pollen offers an important tool for distinguishing C₃ and
57 C₄ grasses in the paleorecord (Amundson et al., 1997; Descolas-Gros and Scholzel, 2007; Nelson
58 et al., 2006). Recent technical advances include Single Pollen Isotope Ratio AnaLysis (SPIRAL),
59 which involves the use of a spooling-wire microcombustion device interfaced with an isotope-
60 ratio mass spectrometer (SWiM-IRMS) for the $\delta^{13}\text{C}$ analysis of individual grass pollen grains
61 (Nelson et al., 2007). Nelson et al. (2007) showed that $\delta^{13}\text{C}$ values of pollen from known C₃ and
62 C₄ grasses could be distinguished based on their distribution around a threshold $\delta^{13}\text{C}$ value of -
63 19.2‰. Although high variability and overlapping ranges of $\delta^{13}\text{C}$ values for C₃ and C₄ grasses
64 prevent perfect classification, a significant correlation was found between $\delta^{13}\text{C}$ -based estimates
65 of % C₄-grass pollen in surface-sediment samples and the abundance of C₄ grasses on the
66 landscape at ten sites in North America (Nelson et al., 2008).

67 Despite the useful paleoenvironmental information obtained from SPIRAL, the existing
68 technique has several inherent limitations. First, SPIRAL was developed (Nelson et al., 2007)
69 and validated (Nelson et al., 2008) with a small amount of data from North American grasses and

70 grasslands. For example, only four C₄ grasses and four C₃ grasses were used to identify the
71 threshold $\delta^{13}\text{C}$ value separating C₃ and C₄ (Nelson et al., 2007). Thus the applicability of this
72 technique to a broader range of grassland ecosystems remains uncertain. Additionally, a fixed
73 $\delta^{13}\text{C}$ threshold was selected to differentiate C₃ and C₄ grasses, which may be problematic
74 because $\delta^{13}\text{C}$ values vary both within and among species (Cerling, 1999). Finally, there is no
75 formal propagation of uncertainty for SPIRAL, which means that the precision of the technique
76 is not well constrained. In this study, we address these problems by (1) expanding the reference
77 $\delta^{13}\text{C}$ dataset for distinguishing C₃- from C₄-grass pollen, (2) improving the validation dataset
78 from North America (Nelson et al., 2008) by adding new surface-sediment samples from lakes in
79 Africa and Australia, and (3) developing and evaluating a hierarchical Bayesian model to
80 estimate the percent of C₃- and C₄-grass pollen based on SPIRAL $\delta^{13}\text{C}$ data.

81

82 **2. METHODOLOGY**

83

84 **2.1 Herbarium and surface-sediment samples**

85 We performed $\delta^{13}\text{C}$ analyses on pollen from herbarium specimens of 28 grass species,
86 including additional pollen from five of the eight species previously analyzed in Nelson et al.
87 (2007) (Electronic Annex EA-1). Our expanded training dataset includes these new results and
88 all of the $\delta^{13}\text{C}$ data reported in Nelson et al. (2007). These specimens were collected between
89 1927 and 1995 from Africa, Australia, and North America.

90 As a step to develop a global relationship between C₄ grass abundance and SPIRAL data,
91 we performed $\delta^{13}\text{C}$ analysis of pollen in surface sediments from Africa and Australia to
92 supplement the published $\delta^{13}\text{C}$ dataset from North America (Nelson et al., 2008). All of the
93 surface-sediment samples from Africa and Australia come from lakes, with the exception of
94 Rumuiku Swamp in Africa (Electronic Annex EA-2). The samples typically represent the upper
95 ~5 cm of sediment, which likely accumulated during the past several decades. We do not have
96 data on the composition and abundance of grasses around our African and Australian sites.
97 Therefore, we estimated the relative abundance or productivity of C₄ grasses around each site
98 based on the relationships of C₄ grasses with various environmental factors reported in the
99 literature (Electronic Annex EA-2). In equatorial East Africa, C₄-grass abundance (Livingstone
100 and Clayton, 1980) and productivity (Tieszen et al., 1979) are negatively correlated with
101 elevation, with C₄ grasses predominating below ~1500 m. We used the relationship in Tieszen et
102 al. (1979) to estimate C₄ grass abundance around each of our African sites. In Australian
103 grasslands, minimum January temperatures (JANT; °C) and median August rainfall (AURF; cm)
104 are strong predictors of C₄ grass abundance in the regional grass flora (Hattersley, 1983). We
105 obtained JANT and AURF data from the Australian Bureau of Meteorology (www.bom.gov.au)
106 and used the relationship in Hattersley (1983) to calculate C₄ grass abundance around each of our
107 Australian sites. For each North American site the percent contribution of C₄ grasses to the total
108 potential production of grasses was determined using the relationship between latitude and C₄-
109 grass productivity (Tieszen et al., 1997).

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113 2.2 Sample treatment and isotopic analysis

114 All samples were treated using standard pollen preparation techniques modified to
115 exclude carbon-containing compounds (Nelson et al., 2006), except that hydrofluoric acid was
116 not used for the herbarium specimens, which has little influence on pollen $\delta^{13}\text{C}$ (Jahren, 2004).
117 Grass pollen grains were isolated in Nanopure water on a microscope slide at 200x magnification
118 using an Eppendorf Transferman micromanipulation device. Individual grains were transferred
119 to ~0.4 μL drops of Nanopure water and applied to a SWiM device interfaced with a
120 ThermoFinnigan Delta V IRMS using a steel and glass syringe (Nelson et al., 2007; Nelson et
121 al., 2008). Sample data were normalized to VPDB using a two-point normalization curve with
122 in-house 2.5 nmol C standards of leucine (true $\delta^{13}\text{C} = -32.1\text{‰}$), sorbitol (true $\delta^{13}\text{C} = -16.2\text{‰}$),
123 serine (true $\delta^{13}\text{C} = -25.7\text{‰}$), and/or glycine (true $\delta^{13}\text{C} = -37.9\text{‰}$) that were calibrated against the
124 USGS40 and USGS41 glutamic acid standards.

125 The number of individual grains of grass pollen applied to the SWiM device ranged from
126 88 to 239 per sample for the herbarium and surface-sediment samples. We followed Nelson et al.
127 (2007; 2008) for the $\delta^{13}\text{C}$ analysis of individual pollen grains. Briefly, along with each sample,
128 we analyzed blanks of Nanopure water to which a single pollen grain was added and then
129 removed. The mean plus 2 standard deviations of blank CO_2 yields was set as a minimum size
130 threshold; grains below this threshold were excluded. The final $\delta^{13}\text{C}$ data were corrected for
131 blank ^{13}C content using isotopic mass balance. The $\delta^{13}\text{C}$ values of herbarium specimens were
132 corrected to a pre-industrial $\delta^{13}\text{C}$ value of atmospheric CO_2 (-6.3‰ ; Friedli et al., 1986).

133

134

135

136 2.3 Statistical model

137 We chose a Bayesian approach for our statistical analysis. Bayesian methods differ
138 theoretically from more widely-used frequentist approaches primarily in that Bayesian methods
139 include *prior* distributions for all unknown parameters to be estimated. Following a fundamental
140 theorem of probability known as Bayes' theorem, prior distributions can be combined with the
141 likelihood of a given dataset (i.e., the probability of observing the dataset, given as a function of
142 unknown parameters) to yield *posterior* parameter distributions. Formally and conceptually, a
143 posterior distribution represents a prior notion of an unknown parameter value, updated with
144 available data according to the proposed model. In many cases (e.g. linear regression), Bayesian
145 and frequentist approaches yield essentially equivalent results when the prior distributions
146 selected are uninformative (i.e. provide little constraint on the unknown parameters), and/or
147 when the dataset is sufficiently large to overwhelm the priors. In other cases, however, the choice
148 of priors can be influential, and the inherent subjectivity in assigning priors has been central to
149 arguments for and against the use of Bayesian methods. For a summary of these theoretical
150 considerations, see Savage (1962).

151 From a pragmatic standpoint, advances in computational methods have provided a
152 consistent and convenient framework for fitting complex models from a Bayesian perspective,
153 where a frequentist approach would be infeasible or impossible. This practical advantage is the
154 motivation for our Bayesian model. The model we propose below is relatively straightforward,
155 and is closely related to model-based clustering methods (Fraley and Raftery, 2002).

156 Nevertheless, the exact model structure is specific to our context and goals, i.e. estimating C_4
157 grass abundance in unknown samples and the likelihood that they contain C_4 grass pollen. We
158 know of no frequentist approach that would suffice to fit such a model, whereas in a Bayesian

159 context it can be solved using generic numerical methods. For a practical introduction to such
 160 methods, we recommend Clark (2007) and Hoff (2009).

161 We designed a hierarchical Bayesian model to predict the percent of C₄ grains in samples
 162 of unknown composition based on the δ¹³C values of individual grass pollen grains (Fig. 1). At
 163 the basis of the model is the likelihood function

$$164 \quad y_i \sim \begin{cases} N(m_{C_3}, S_{C_3}^2), & x_i = 0 \\ N(m_{C_4}, S_{C_4}^2), & x_i = 1 \end{cases}$$

165 in which, for the i^{th} grain in the sample, y_i is the measured δ¹³C of the grain, x_i is a binary
 166 variable identifying the grain as C₃ ($x_i = 0$) or C₄ ($x_i = 1$), μ and σ^2 represent the population
 167 means and variances (respectively) for C₃ and C₄ grains as indicated by subscripts, and $N(\mu, \sigma^2)$
 168 denotes the normal (Gaussian) distribution with mean μ and variance σ^2 . In other words, the
 169 likelihood is the conditional probability of observing the δ¹³C value of an individual grain, given
 170 the classification of the grain and assuming normally-distributed δ¹³C values for both C₃ and C₄.
 171 We calculated m_{C_3} , m_{C_4} , $S_{C_3}^2$, and $S_{C_4}^2$ from the herbarium dataset described above, and
 172 subsequently treated these variables as fixed in our predictive model.

173 Because the C₃/C₄ identity of the pollen grains in sediment samples is unknown, we
 174 added a second hierarchical level to model x , the indicator variable for C₄ presence, based on the
 175 unknown proportion of C₄ grains in the population, θ :

$$176 \quad x_i \sim \text{Bernoulli}(\theta)$$

177 i.e.,

$$x_i = \begin{cases} 1 & \text{with probability } q \\ 0 & \text{with probability } (1 - q) \end{cases}$$

The unknown parameter θ requires a prior distribution as well. In defining this prior, we introduced a final hierarchical level in the model to accommodate samples composed of (1) purely C_3 , (2) purely C_4 , or (3) both C_3 and C_4 pollen grains. We refer to these sample types as “ C_3 -only”, “ C_4 -only”, and “mixed”, respectively, and define the prior distribution of θ separately for each:

$$q \sim \begin{cases} 0 & \text{for } C_3\text{-only samples} \\ \text{Uniform}(0,1) & \text{for mixed samples} \\ 1 & \text{for } C_4\text{-only samples} \end{cases}$$

In other words, if a sample is identified as C_3 -only or C_4 -only, then θ is assigned a constant value of 0 or 1 (respectively). For mixed samples, θ must be estimated based on the data. In this case, the uniform prior represents our lack of knowledge of the true proportion of C_4 in the sample by assuming *a priori* that all values of θ are equally likely.

The compound prior on θ effectively defines three distinct sub-models. In a Bayesian framework, these models can be fit simultaneously to formally compare their ability to describe a given dataset. This simple form of Bayesian model selection (Dellaportas et al., 2002) treats the choice of model itself as an unknown parameter, which therefore requires its own prior distribution. We assumed that the sub-models were equally likely *a priori*, and thus assigned each a prior probability of 1/3. The posterior estimate of the model-selection parameter then yields “posterior model probabilities” representing the relative probability that each candidate model (i.e. sample type) is the true model. This allows for hypothesis testing analogous to the

197 use of p -values (e.g. rejecting a candidate model if it has a posterior probability <0.05 ; Marden,
198 2000).

199 The division of the main hierarchy into three possible submodels serves two purposes.
200 First, for samples that truly contain only one pollen type, the corresponding monotypic model is
201 conceptually correct, and generally provides a better fit than if only the “mixed” model is
202 allowed (data not shown). Second, fitting this model produces a posterior estimate of θ while
203 simultaneously calculating the posterior probability of each sample type. In applications aimed
204 primarily at assessing the relative abundance of C_4 grains in a sample (e.g. to compare C_4
205 abundance across space or time), θ will be of primary interest. However, in some cases the goal
206 of SPIRAL may be to identify whether one pollen type is present or absent in a sample (e.g.
207 Urban et al., 2010). For that purpose, the posterior model probabilities allow explicit
208 quantification of the probability that either or both types are present.

209 We fit the model by Markov Chain Monte Carlo (MCMC) sampling using the software
210 package JAGS (version 3.1.0; Plummer, 2011) interfaced through R (R Development Core
211 Team, 2010) with the library *rjags* (Plummer, 2012). Briefly, the software uses a variety of
212 MCMC algorithms to sample over possible values of the unknown parameters. For each
213 parameter, the resulting posterior distribution (i.e. histogram of all values sampled during the
214 MCMC sequence) is an approximation of the true probability density function of the parameter
215 given the dataset of observations. Any population statistic of interest can then be estimated from
216 the corresponding sample statistic for the MCMC sample. For example, we summarize θ by its
217 posterior median, calculated as the sample median across the entire MCMC sequence.

218 We used pseudodata from the herbarium samples to verify the model. We produced
219 samples with known composition of 0 to 100% C_4 in 10% increments, and sample sizes of 50,

220 100, and 150 grains. We randomly generated 1000 replicates of each % C₄ X sample-size
221 combination, and fit the model to each replicate sample to generate posterior estimates of θ and
222 posterior probabilities for each sample type (C₃-only, C₄-only, or mixed). For comparison, we
223 also estimated % C₄ for each sample using the threshold-based classification method (i.e. Nelson
224 et al., 2007), but with the threshold value (the midpoint between m_{C_3} and m_{C_4}) updated to reflect
225 the expanded herbarium dataset. Finally, we used the model to estimate the percent of C₄ grains
226 in the surface sediments of sites in Africa, Australia, and North America. For comparison of
227 these estimates with the relative abundance of C₄ grasses on the landscape, we used reduced
228 major axis regression because of symmetry in the variables on the x and y axes (Smith, 2009),
229 and because both the x and y variables contain uncertainty (McArdle, 1988). The fit of this
230 regression was compared with a 1:1 relationship following equations outlined in McArdle
231 (1988). These regression analyses were performed in R (R Development Core Team, 2010).

232

233 3. RESULTS AND DISCUSSION

234

235 3.1 $\delta^{13}\text{C}$ of C₃ and C₄ grass pollen: an expanded training set

236 The expanded training set is based on pollen from 31 herbarium specimens. The number
237 of grass pollen grains applied to the moving wire with peak areas exceeding the 2σ threshold of
238 blanks ranges between 21 and 130 grains per sample, with an average of 62 grains per sample
239 (Electronic Annex EA-1). The expanded training set therefore includes 1,921 $\delta^{13}\text{C}$ values, 1,402
240 of which were obtained as part of the present study. An average of 32% of applications of pollen
241 from herbarium samples yield a peak area above the blank threshold, which is lower than results
242 from surface-sediment samples from North American lakes (47%, Nelson et al., 2008) and

243 Miocene/Oligocene sediment samples (45%, Urban et al., 2010). The mean $\delta^{13}\text{C}$ values of grass
244 pollen range between -42.7 and -24.0‰ for C_3 species and between -17.2 and -10.5‰ for C_4
245 species (Electronic Annex EA-1). A majority of the pollen $\delta^{13}\text{C}$ values fall within the typical
246 $\delta^{13}\text{C}$ ranges for C_3 (-34 to -22‰) and C_4 (-15 to -10‰) plants (Fig. 2; Electronic Annex EA-1).
247 However, similar to previous results, the $\delta^{13}\text{C}$ variation is large, with many individual data points
248 exceeding these ranges, likely because of variability in the magnitude and composition of the
249 analytical blank (Nelson et al., 2007).

250 The updated herbarium dataset yields somewhat different parameter estimates than those
251 reported by Nelson et al. (2007). Estimates of $m_{\text{C}_3} = -29.6\text{‰}$ and $m_{\text{C}_4} = -13.8\text{‰}$ are more
252 negative than previously determined values (-26.9‰ and -11.5‰, respectively), leading to an
253 estimated threshold value of -21.7‰ that is also more negative than the original value (-19.2‰).
254 Variability of $\delta^{13}\text{C}$ in the new dataset is similar between C_3 and C_4 grains (standard deviation =
255 9.5‰ for each), which is greater than previously determined for C_3 (6.3‰), but similar for C_4
256 (9.6‰). Based on the updated values, the probability of an individual grain being identified as C_4
257 by the Bayesian model varies smoothly over the range of possible $\delta^{13}\text{C}$ values (Fig. 2).

258 In terms of estimating the overall composition of unknown samples, the pseudodata
259 experiments show a striking improvement of the Bayesian approach. Overall, results from
260 samples of pseudodata randomly generated from the herbarium dataset illustrate that Bayesian
261 estimates of % C_4 grass pollen are highly accurate (Fig. 3). For all sample sizes tested, bias (i.e.,
262 the mean deviation between the estimated and true % C_4) is $\leq 5.5\%$, with largest biases when true
263 C_4 composition is 80% ($n=50$) or 10% ($n \geq 100$). Average biases across all true % C_4 values are
264 only 2.9% for sample size $n=50$, and 2.4% for $n=100$ and $n=150$. By contrast, the original
265 threshold-based methodology of Nelson et al. (2007) produces accurate estimates of sample

266 composition when true composition is near 50%, but becomes increasingly biased towards
267 underestimation (overestimation) as true % C_4 increases (decreases). Maximum bias of ~16% for
268 the threshold-based approach occurs for purely C_3 or C_4 samples, and average bias across all true
269 % C_4 values is 8.2%.

270 The improved accuracy of the Bayesian model for samples with low and high abundances
271 of C_4 grass pollen is a function of its hierarchical structure. The model explicitly incorporates θ ,
272 the estimated relative abundance of C_4 grains in the population, as well as a model-selection
273 parameter representing the possibility that either C_3 or C_4 can be entirely absent from a sample.
274 The MCMC approach then solves for these parameters simultaneously while accounting for the
275 fact that they both influence the likelihood of an individual grain being identified as C_3 or C_4 . By
276 contrast, the threshold method relies on a fixed threshold value with classification accuracies for
277 C_3 and C_4 grains that are independent of sample composition. In practice, the threshold method
278 misclassifies approximately the same percent C_3 and C_4 grains. Thus, near 50% true C_4
279 abundance, the number of misclassification errors for C_3 and C_4 are similar, which results in
280 offsetting effects on estimated % C_4 and small net bias. However, when % C_4 is far from 50%
281 the misclassification errors are imbalanced, which results in a biased estimate of % C_4 .

282 To illustrate how the hierarchical Bayesian model overcomes this limitation, here we
283 consider a hypothetical sample with low (<50%) C_4 abundance, and we note that the opposite
284 rationale applies for samples with high C_4 abundance. For a low- C_4 sample, the data favor a
285 correspondingly low estimate of θ . Consequently, the likelihood of any grain being identified as
286 C_4 in the model is diminished, reflecting the reduced probability of a C_4 grain being found in a
287 sample when the true abundance of C_4 grains is low. This in turn causes fewer C_3 grains with
288 ambiguous $\delta^{13}\text{C}$ values to be misclassified as C_4 . As the true percent of C_4 in the hypothetical

289 sample approaches 0, the data will begin to favor selection of the C₃-only model, which prevents
290 misidentification of C₃ grains. These same mechanisms lead to an increased proportion of C₄
291 grains misclassified as C₃ in low-C₄ samples. However, since a sample with low C₄ abundance
292 has fewer C₄ than C₃ grains by definition, the net effect is an improvement in accuracy relative to
293 the threshold-based method.

294 Our Bayesian model can also be used to assess the presence or absence of C₄ grasses on
295 the landscape (Fig. 4). For example, for pseudodata samples containing 0% C₄, the posterior
296 probability of the C₃-only model [P(C₃-only)] has a median value of >0.95, indicating strong
297 preference for the correct model most of the time. Similarly, for pseudodata samples containing
298 100% C₄, median P(C₄-only) is ~0.94 indicating strong preference for the C₄-only model.
299 Furthermore, our results suggest that the method has substantial power to reject the C₃-only
300 model when C₄ grains are in fact present. For example, with a sample size of 100 grains, median
301 P(C₃-only) is <0.01 for samples with only 20% C₄. Samples with C₄ present in lower abundance
302 are more ambiguous. Among samples with 10% C₄, for instance, median P(C₃-only) of a 100-
303 grain sample is 0.54. The ability to identify C₄ presence improves with sample size. For example,
304 for a sample with 10% C₄, median P(C₃-only) is 0.23 with n=150 grains, compared to 0.73 with
305 n=50 grains. Thus, for samples of relatively large size (≥ 100 grains) the practical detection limit
306 for reliably identifying the presence of C₄ grains in a sample is between 10-20% C₄.

307

308 **3.2. Field validation of grass-pollen $\delta^{13}\text{C}$ as a proxy indicator of C₃/C₄ abundance**

309 For the surface-sediment samples from Africa and Australia, the number of grass pollen
310 grains with peak areas exceeding the 2σ threshold of blanks ranges between 30 and 142 grains
311 per sample, with an average of 52 grains per sample (Electronic Annex EA-2). The total surface-

312 sediment dataset therefore includes 1,522 $\delta^{13}\text{C}$ values, 773 of which were obtained as part of the
313 present study. On average, 48% of applications of pollen from sediment samples yield a peak
314 area above the blank threshold. A majority of the pollen $\delta^{13}\text{C}$ values fall within or between
315 typical $\delta^{13}\text{C}$ ranges for C_3 and C_4 plants (Electronic Annex EA-2, EA-3, and EA-4). However, as
316 with the expanded herbarium dataset, the $\delta^{13}\text{C}$ variation is large.

317 Bayesian estimates of the median % C_4 grass pollen from the surface-sediment samples
318 range between 0 and 99% (Fig 5; Electronic Annex EA-2). Across the large spatial and
319 environmental gradients represented by our surface-sediment sites, we expected that the
320 abundance of C_3 and C_4 grass pollen in surface sediments would be overall similar to the
321 abundance of C_3 and C_4 grasses on the landscape. Consistent with this expectation, there was a
322 significant relationship between the Bayesian estimates of % C_4 grass pollen in the surface-
323 sediment samples from Africa, Australia, and North America and C_4 -grass abundance around
324 these sites (Fig. 5; $n=24$, $p<0.001$, $r^2=0.65$). Furthermore, this relationship does not differ from
325 a 1:1 relationship ($p=0.45$), indicating no consistent bias in the representation of C_3 and C_4
326 grasses that may be associated with factors such as pollen productivities or preservation in
327 sediments. We excluded one site, Rumuiku Swamp, from the regression because it had unusually
328 low % C_4 grass pollen for its elevation, probably because the local swamp environment
329 supported a greater abundance of C_3 grasses. However, the regression remains significant even if
330 Rumuiku swamp is included ($n=25$, $p<0.001$, $r^2=0.54$). Nelson et al. (2008) found a similar
331 relationship in North America using the original (-19.2‰) threshold method, but lacked data
332 from sites with $<20\%$ C_4 grass abundance on the landscape. The additional data in the present
333 study helps to extend this range and further validates SPIRAL as a tool for paleoenvironmental
334 reconstruction.

335

336 **3.3 Application to the paleorecord: interpreting SPIRAL $\delta^{13}\text{C}$ data in the Bayesian**
337 **framework**

338 The improved estimates of C_4 -grass abundance from incorporation of SPIRAL data into
339 the Bayesian model can help to assess factors (e.g. atmospheric CO_2 concentrations) controlling
340 the origin, expansion, and variations in abundance of C_4 grasses in Earth's history. To
341 demonstrate the application of the model to the paleorecord, we reevaluated a published SPIRAL
342 dataset (Urban et al., 2010). Briefly, Urban et al. (2010) measured $\delta^{13}\text{C}$ of grass pollen grains in
343 sediments spanning the early-Oligocene to middle-Miocene from sites in southwestern Europe
344 and used a threshold value of -19.2‰ (before modification for variations in $\delta^{13}\text{C}$ of atmospheric
345 CO_2 and aridity) to detect the presence of pollen from C_4 grasses. The samples in that study
346 contained between 63 and 100 grains. Results indicated that C_4 grasses appeared on the
347 landscape of southwestern Europe no later than the early Oligocene, which suggests that low
348 $p\text{CO}_2$ may not have been the main driver and/or precondition for the development of C_4
349 photosynthesis in the grass family.

350 We evaluated the probability that the $\delta^{13}\text{C}$ data in samples from Urban et al. (2010)
351 support the C_3 -only model in our Bayesian analysis. We adjusted the $\delta^{13}\text{C}$ values of the Urban et
352 al. (2010) samples to that of pre-industrial $\delta^{13}\text{C}$ of atmospheric CO_2 (-6.3‰) using estimated
353 values of $\delta^{13}\text{C}$ of atmospheric CO_2 during the Cenozoic based on benthic foraminifera $\delta^{13}\text{C}$ data
354 (Tippie et al., 2010). The probability of a C_3 -only model was <0.01 (indicating $>99\%$ probability
355 that at least some C_4 grains were present) for all samples (Electronic Annex EA-5). However,
356 low water availability may have caused the $\delta^{13}\text{C}$ values of C_3 plants to shift in the positive
357 direction (Ehleringer and Cooper, 1988). To account for the potential influence of aridity we

358 shifted the mean $\delta^{13}\text{C}$ value of our C_3 training set by 1-3‰ in the positive direction, as in Urban
359 et al. (2010). All but one sample had a $P(\text{C}_3\text{-only})$ of <0.01 after addition of 1‰ to the mean $\delta^{13}\text{C}$
360 value of the C_3 training dataset. Six of the eight samples, including the oldest two, had a $P(\text{C}_3\text{-}$
361 $\text{only})$ of <0.05 after addition of 3‰ to the mean $\delta^{13}\text{C}$ value of the C_3 training dataset (Electronic
362 Annex EA-5). The mean Bayesian estimates of % C_4 grass pollen are particularly high in the
363 oldest two samples, consistent with the identification of plant communities in regions where
364 today C_4 grasses are dominant as the closest analogs for the corresponding pollen assemblages
365 (Suc, 1984). Therefore, our Bayesian estimates of % C_4 grass pollen confirm the prior conclusion
366 of Urban et al. (2010) that C_4 grasses occurred on the landscape of southwestern Europe by at
367 least the early Oligocene. The main advantage of the Bayesian model over the threshold
368 approach used the context of the Urban et al. (2010) study is that the former allows for an
369 explicit estimate of the probability of C_4 grasses being present on the landscape, which is
370 essential for quantitatively assessing the timing of C_4 -grass origin in geological history.

371 Overall, our new $\delta^{13}\text{C}$ data along with the Bayesian framework improve quantitative
372 reconstructions of variation in the relative abundance of C_3 and C_4 grasses in response to
373 environmental changes in the paleorecord. The flexible and hierarchical nature of the Bayesian
374 model yields more accurate estimation of the abundance of C_4 grass pollen than the simpler, but
375 biased, threshold approach, and also provides posterior model probabilities that enable
376 hypothesis testing. Thus we recommend that future estimates of C_3 and C_4 grass abundances
377 should, when possible, be made using Bayesian methods rather than threshold-based counting
378 approaches. Bayesian analyses have begun to have important applications in the interpretations
379 of geochemical isotope data. For example, recent studies have used Bayesian analysis to develop
380 probabilistic region-of-origin assignments in wildlife and human forensics (Kennedy et al., 2011;

381 Wunder, 2010), enhance radiocarbon-age modeling for sediment records (Blaauw et al., 2007;
382 Blaauw and Christen, 2011), and enable detection of climate-related shifts in elemental and
383 isotopic abundances in peat cores (Gallagher et al., 2011). The increased use of Bayesian
384 approaches promises to transform the environmental interpretations of geochemical data,
385 especially in cases where small samples are involved. We expect that Bayesian analyses will
386 become a mainstay of geochemistry.

387

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495 **Figure legends**496
497

498 Figure 1. Conceptual diagram of the hierarchical Bayesian model used in this study. The
499 likelihood function describes the probability distribution of $\delta^{13}\text{C}$ values for each pollen grain in a
500 sample (y_i), given its classification as C_3 or C_4 ($x_i = 0$ or $x_i = 1$, respectively). The distribution of
501 x_i in turn depends on θ , the proportion of C_4 grains in the population. Finally, the prior
502 distribution of θ varies among sub-models representing three possible sample types (C_3 -only,
503 mixed, C_4 -only). See Section 2.3 for details.

504

505 Figure 2. Histograms of $\delta^{13}\text{C}$ values from individual grains of grass pollen (1‰ bins). The
506 dashed grey line represents data from C_3 grasses and the black line data from C_4 grasses (y-axis
507 on left). The solid grey line represents the calculated probability of individual grains being
508 classified as C_4 as a function of $\delta^{13}\text{C}$ (y-axis on right).

509

510 Figure 3. Estimated vs. true % of C_4 grains in samples of pseudodata derived from the herbarium
511 training dataset. Columns correspond to three sample sizes ($n=50, 100,$ and 150 grains). Rows
512 correspond to results from Bayesian (top) and threshold (bottom) methods. For each panel, the
513 mean (thick black line) and 95% confidence intervals (thin black lines) of estimates from 1000
514 random samples are plotted. The solid grey lines represent 1:1 relationships.

515

516 Figure 4. Probability that each candidate model (rows: C_3 -only, mixed, and C_4 -only) is the true
517 model for pseudodata samples of known size (columns: 50, 100, or 150 grains) and composition
518 (x-axis: 0-100% C_4). The dashed grey horizontal lines represent $p = 0.05$. For each set of

519 pseudodata samples, the box represents the 25-75th percentiles of posterior probabilities, with
520 median indicated by a heavy black line. The whiskers encompass all remaining points within 1.5
521 times the interquartile range of the box, and points outside this range are plotted individually.

522
523 Figure 5. Estimated C₄ coverage (%) on the landscape around lakes in Africa (diamonds),
524 Australia (X symbol), and North America (triangles), compared to the abundance of C₄ grass
525 pollen (%) in the surface-sediments of these sites, as estimated from $\delta^{13}\text{C}$ of individual grains of
526 grass pollen using the Bayesian model. The major axis slope is 0.97 and 95% confidence interval
527 of the slope is 0.75 - 1.24. The data point with an asterisk is excluded from the regression, as
528 explained in section 3.2. The 1:1 line is the solid grey line; the regression line is represented by
529 the black dashed line. Error bars on each data point represent 95% confidence intervals.

530

531

Figure 1

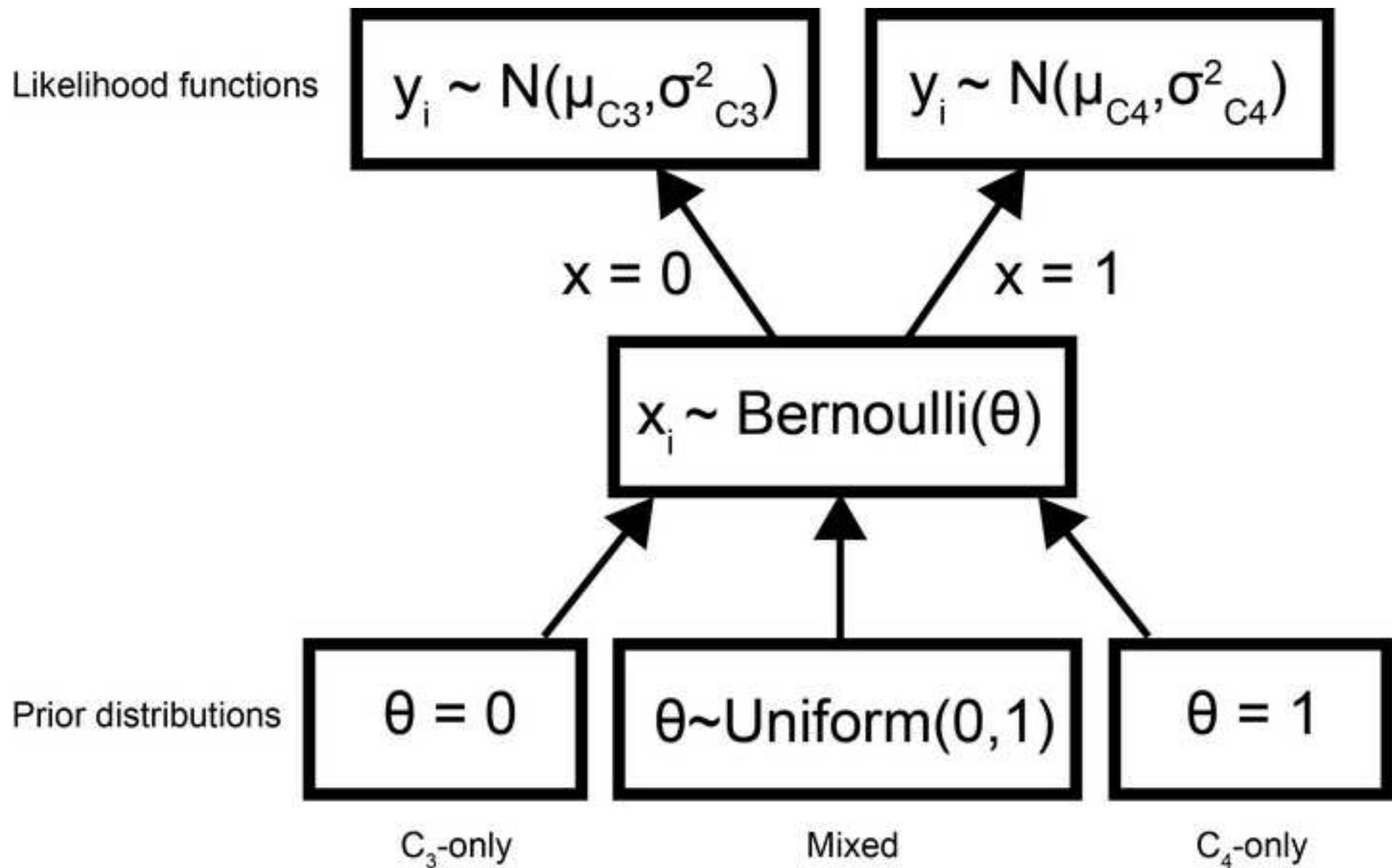
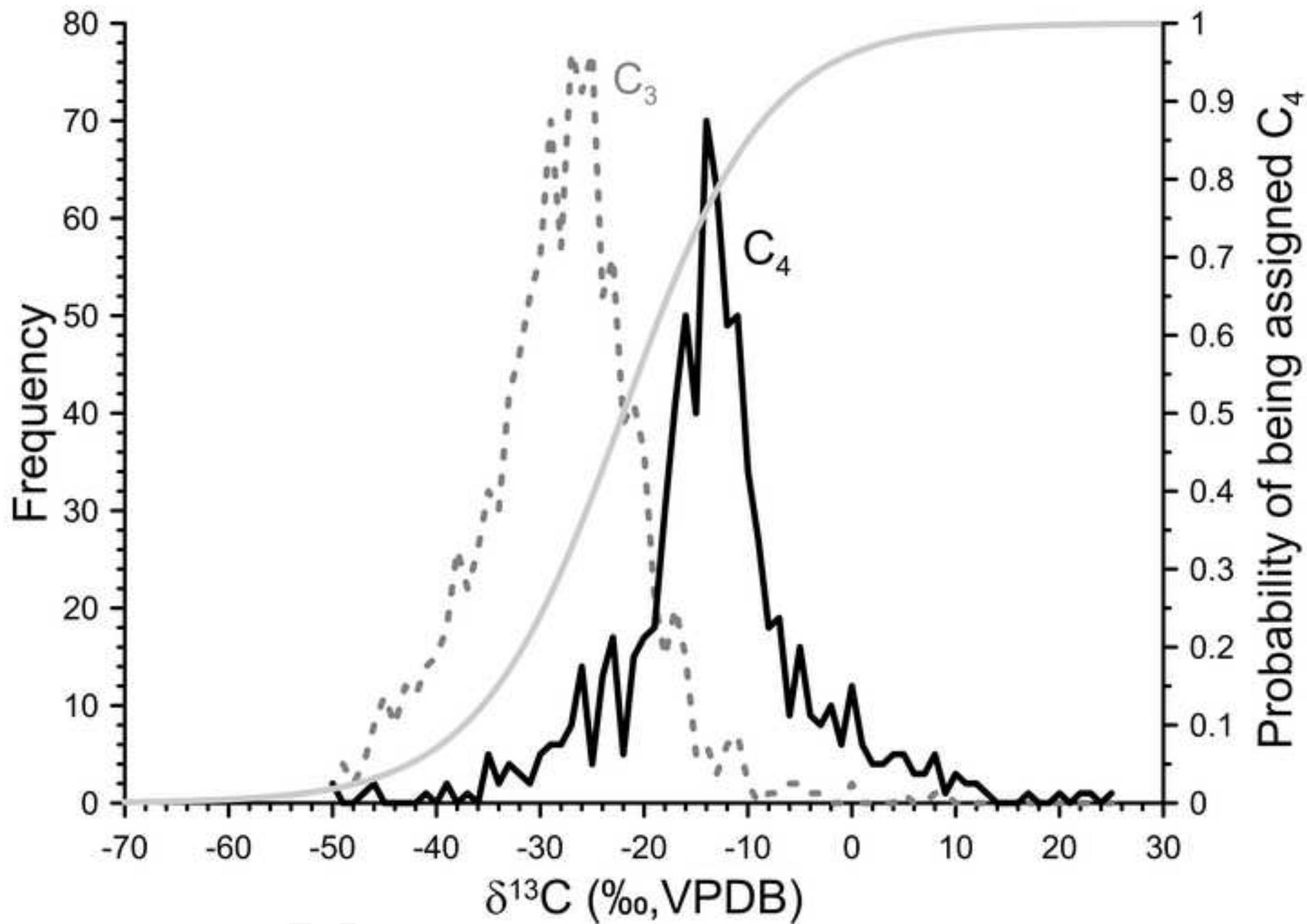


Figure 2

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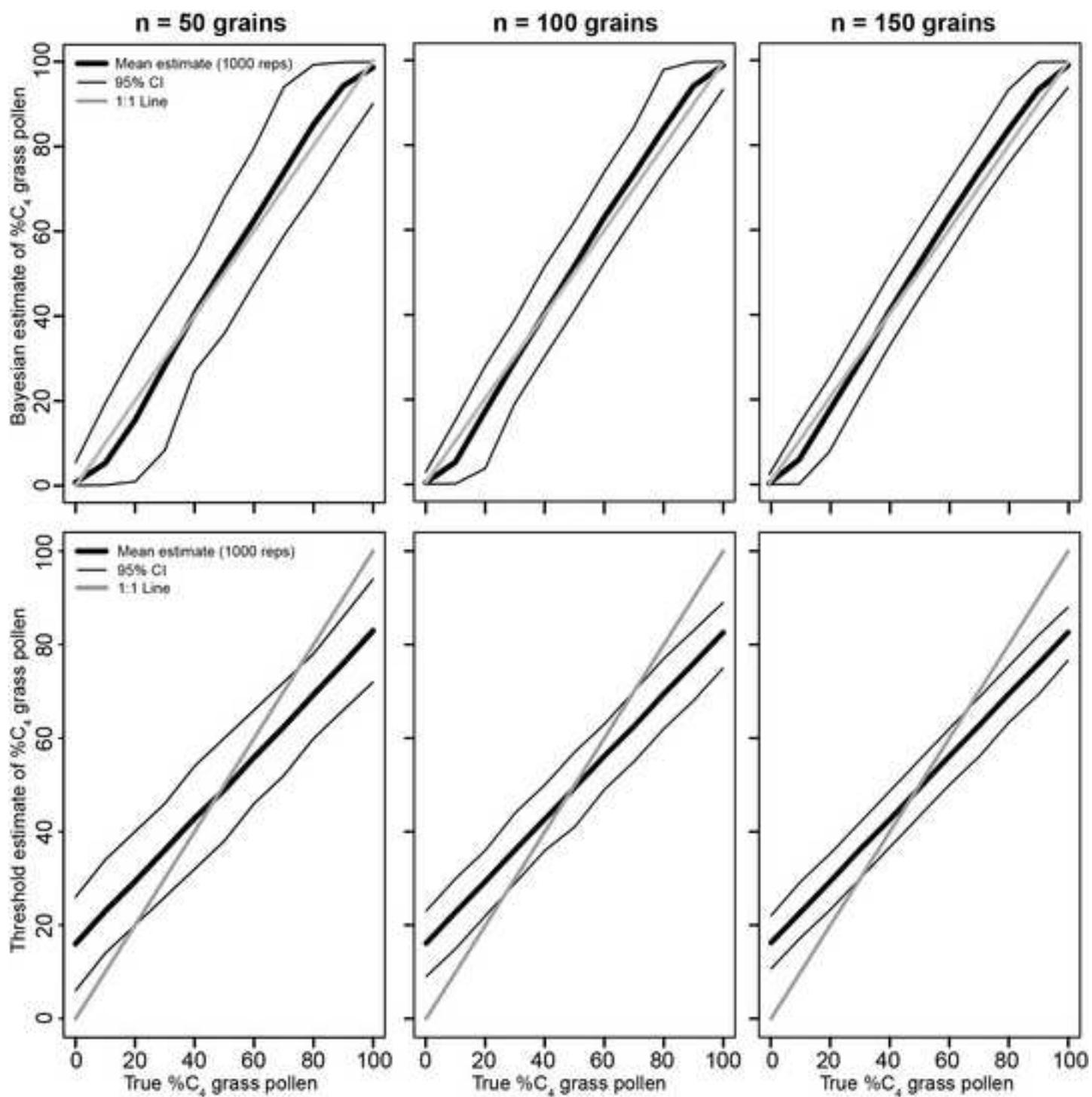


Figure 4

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