

# Environmental drivers of freshwater macrophyte diversity and community composition in calcareous warm-water rivers of America and Africa

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Author post-print (accepted) deposited by Coventry University's Repository

## Original citation & hyperlink:

Tapia Grimaldo, J, O'Hare, MT, Kennedy, MP, Davidson, TA, Bonilla-Barbosa, J, Santamaría-Araúz, B, Gettys, L, Varandas Martins, S, Thomaz, SM & Murphy, KJ 2017, 'Environmental drivers of freshwater macrophyte diversity and community composition in calcareous warm-water rivers of America and Africa' *Freshwater Biology*, vol 62, no. 9, pp. 1511-1527

<https://dx.doi.org/10.1111/fwb.12962>

DOI 10.1111/fwb.12962

ISSN 0046-5070

ESSN 1365-2427

Publisher: Wiley

**This is the peer reviewed version of the following article: Tapia Grimaldo, J, O'Hare, MT, Kennedy, MP, Davidson, TA, Bonilla-Barbosa, J, Santamaría-Araúz, B, Gettys, L, Varandas Martins, S, Thomaz, SM & Murphy, KJ 2017, 'Environmental drivers of freshwater macrophyte diversity and community composition in calcareous warm-water rivers of America and Africa' *Freshwater Biology*, vol 62, no. 9, pp. 1511-1527, which has been published in final form at**

<https://dx.doi.org/10.1111/fwb.12962>

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1 **Environmental drivers of freshwater macrophyte diversity and community**  
2 **composition in calcareous warm-water rivers of America and Africa**

3

4 Julissa Tapia Grimaldo<sup>1,2</sup>, Matthew T. O'Hare<sup>2</sup>, Michael P. Kennedy<sup>3\*</sup>, Thomas A. Davidson<sup>4</sup>,  
5 Jaime Bonilla-Barbosa<sup>5</sup>, Betzy Santamaría-Araúz<sup>5</sup>, Lyn Gettys<sup>6</sup>, Sara Varandas Martins<sup>1</sup>,  
6 Sidinei M. Thomaz<sup>7</sup> and Kevin J. Murphy<sup>1</sup>

7

8 <sup>1</sup>Institute of Biodiversity, Animal Health and Comparative Medicine, University of Glasgow,  
9 Glasgow G12 8QQ, Scotland

10 <sup>2</sup>Centre for Ecology and Hydrology, Bush Estate, Penicuik, Midlothian EH26 0QB, Scotland

11 <sup>3</sup>School of Energy, Construction and Environment, Coventry University, Priory Street,  
12 Coventry CV1 5FB, England

13 <sup>4</sup>Department of Bioscience, 25 Vejlsøvej, 8600, Silkeborg, Aarhus University, Denmark

14 <sup>5</sup>Laboratorio de Hidrobotánica, Departamento de Biología Vegetal, Centro de  
15 Investigaciones Biológicas, Universidad Autónoma del Estado de Morelos, Av. Universidad  
16 1001, Col. Chamilpa, 62209 Cuernavaca, Morelos, México

17 <sup>6</sup>University of Florida, Fort Lauderdale Research and Education Center, 3205 College Ave,  
18 Davie, Florida, FL 33314, USA

19 <sup>7</sup>Nupélia, Universidade Estadual de Maringá, Maringá, PR, Brasil

20

21 Keywords: alpha-diversity, beta-diversity, freshwater plants, hardwater rivers, latitude

22

23 Running-title: Macrophytes in tropical hardwater rivers

24 \*Correspondence: Michael Kennedy, School of Energy, Construction and Environment,  
25 Coventry University, Priory Street, Coventry CV1 5FB, England; [ab9280@cov.ac.uk](mailto:ab9280@cov.ac.uk)

26 **SUMMARY**

27 1. This study assessed the hypothesis that spatial and environmental drivers of river  
28 macrophyte diversity and community composition differ in relative importance in calcareous  
29 river systems located in warm regions of America versus Africa.

30 2. We collected aquatic vegetation and spatio-environmental data, during 2006 – 2011, from  
31 >200 hardwater rivers, and associated floodplain waterbodies, located up to 30° North or  
32 South of the Equator, in México, Trinidad, Brasil, Argentina, USA (Florida), South Africa,  
33 Botswana, and Zambia.

34 3. Species rarefaction procedures were used to assess the impacts of differing sampling  
35 effort in the two continents upon estimation of  $\gamma$ -diversity (“species pool”). We then used a  
36 cluster analysis approach (Two-Way Indicator Species Analysis: TWINSpan) to classify  
37 samples into groups based upon species composition. Variation in species richness,  
38 community composition and six spatial and environmental variables, among samples making  
39 up these groups, were compared using ANOVA and Kruskal-Wallis procedures. Regression  
40 trees and redundancy analysis were used to infer the relative importance of spatial and  
41 environmental drivers in explaining variation in local species richness and species  
42 community composition between the two continents. Sorensen’s index ( $C_s$ ) was calculated to  
43 estimate species turnover ( $\beta$ -diversity) between African and American samples.

44 4. In total 378 macrophyte taxa were recorded, with no significant difference in mean  
45 macrophyte  $\alpha$ -diversity between African and American sites, but with evidence for high  
46 species turnover between the two continents ( $C_s = 0.17$ ). Rarefaction analysis confirmed the  
47 existence of a larger macrophyte species pool in the hardwater rivers sampled in Africa  
48 compared to America. TWINSpan classification identified seven sample end-groups, only  
49 one of which contained a mix of sites from both continents. PERMANOVA and nMDS  
50 ordination analysis confirmed significant differences in community composition present in

51 these sample groups. There were substantial differences between the sample-groups for  $\alpha$ -  
52 diversity, and for spatial and environmental variables.

53 5. The high species turnover between Africa and America may be accounted for by  
54 geographical segregation, along with differences in aquatic habitat characteristics, and  
55 varying long-distance dispersal capacities of individual species. The relative importance of  
56 spatial and physico-chemical drivers (latitude, pH, altitude, alkalinity and electrical  
57 conductivity; but not flow) differed between the continents in influencing variation in both  
58 macrophyte diversity and community composition. Latitude was a significant,  
59 though non-linear and rather complex, spatial driver of macrophyte  $\alpha$ -diversity in both  
60 American and African hardwater rivers. Water chemistry variables varied in relative  
61 importance as drivers of macrophyte  $\alpha$ -diversity for African and American sites individually,  
62 and for all sites combined, but pH and/or electrical conductivity were more important than  
63 alkalinity in each case. In all three cases, altitude was consistently the third most important  
64 driver of  $\alpha$ -diversity. Spatial and environmental variables played important roles in structuring  
65 macrophyte community composition in warm-water calcareous rivers in both America and  
66 Africa, with latitude being the strongest individual driver. Thus, this spatial variable, which is  
67 a surrogate for numerous enviro-climatic variables, appears to be of importance in  
68 determining macrophyte distributions at large spatial scales, for the ecosystem type  
69 examined here.

70

## 71 **Introduction**

72 Recently, there has been a major effort to improve understanding of the drivers of  
73 biogeographic distributions and diversity of freshwater macrophyte species, some (but by no  
74 means all) of which have broad planetary distributions (e.g. Bornette *et al.*, 1998;  
75 Santamaría, 2002; Murphy *et al.*, 2003; Makkay *et al.*, 2008; Carvalho *et al.*, 2009; Heikkinen  
76 *et al.*, 2009; Lang & Murphy, 2012; Chappuis *et al.*, 2012, 2014; Kennedy *et al.*, 2015, 2017;  
77 Morandeira & Kandus, 2015; Ranieri *et al.*, 2015; Tapia Grimaldo *et al.*, 2016; Redekop *et*  
78 *al.*, 2016; Alahuhta *et al.*, 2017). Most of these studies have examined macrophyte diversity  
79 and distributions in cool-temperate river and lake systems, with least attention being paid to  
80 warm-water river macrophyte communities. Even fewer studies have directly compared Old  
81 and New World freshwater macrophyte ecology: a rare example is Jacobsen & Terneus  
82 (2001), on stream vegetation in Ecuador and Denmark. Examples of environmental drivers  
83 variously reported to be important, at differing geographical scales, include enviro-climatic  
84 factors associated with variation in latitude (e.g. evapotranspiration regime), and  
85 environmental heterogeneity associated with a range of physico-chemical factors. Altitude,  
86 water and substrate chemistry, flow regime and human-related habitat alteration are often  
87 considered relevant in this context.

88 Freshwater macrophytes are “aquatic photosynthetic organisms large enough to see with the  
89 naked eye, that actively grow permanently or periodically submerged below, floating on, or  
90 growing up through the water surface” of freshwater systems (Chambers *et al.*, 2008). In this  
91 study we deal only with vascular freshwater macrophytes, not considering bryophytes or  
92 macroalgae.

93 There is good evidence that the Neotropical biogeographic region, comprising South and  
94 Central America, plus a small area of North America, namely part of Texas and most of  
95 Florida (Escalante *et al.*, 2010), is a global hotspot for vascular freshwater macrophyte  
96 biodiversity, with a recorded  $\gamma$ -diversity (species regional pool) of 984 macrophyte species,

97 according to Chambers *et al.* (2008). In contrast the Afrotropical region (Africa and the  
98 Arabian Peninsula, south of the Tropic of Cancer) has a lower macrophyte  $\gamma$ -diversity, with  
99 614 species, while the Nearctic (Greenland and North America, excluding parts of Texas and  
100 Florida) has a macrophyte  $\gamma$ -diversity slightly higher than the value for the Afrotropics, at 644  
101 species (Chambers *et al.*, 2008).

102 It is not known whether these differences in diversity occur because of natural causes (e.g.,  
103 habitat limitations, or for evolutionary reasons), or are due to differences in sampling effort,  
104 or both. Afrotropical freshwaters are probably under-recorded for aquatic plant species  
105 (examples of, usually quite local, surveys include: Denny, 1973, 1985; Simpson, 1975;  
106 Chabwela & Siwale, 1986; Machena, 1988; Sarr *et al.* 2001; Adesina *et al.*, 2011; Achieng'  
107 *et al.*, 2014). A recent survey of 228 sites in Zambian rivers (including both hard- and  
108 softwater systems: Kennedy *et al.*, 2015) recorded 335 macrophyte taxa, but the cumulative  
109 sequential records curve for the dataset showed little sign of reaching an asymptote. It is  
110 hence likely that many additional macrophyte species remain to be found in Zambian rivers,  
111 and the situation is probably the same for other tropical African countries.

112 In contrast there has been quite a substantial macrophyte survey effort in the Neotropics,  
113 particularly in South American freshwater systems (e.g., Bertoli, 1996; Murphy *et al.*, 2003;  
114 Thomaz *et al.*, 2009; Rolon & Matchik, 2006; Amaral *et al.*, 2008; Sousa *et al.*, 2010, 2011;  
115 Varandas Martins *et al.*, 2013; Bottino *et al.*, 2014; Neiff *et al.*, 2014; Bando *et al.*, 2015;  
116 Schneider *et al.*, 2015), though less so in Central America (e.g., Crow, 1993; Philbrick *et al.*,  
117 1995; Anonymous, 1999; Bonilla-Barbosa, 2004). Compared to Africa, the macrophyte flora  
118 of the Neotropics is probably reasonably well known, although there is evidence that the  
119 asymptote of the species-sampling effort curve (for all freshwater habitats combined) has not  
120 been reached in this region either (e.g., Ferreira *et al.*, 2011).

121 In the Nearctic the survey effort for aquatic macrophyte vegetation has been very  
122 substantial, with >2000 publications on the macrophyte ecology (of both the Nearctic and

123 Neotropical parts) of Florida alone held, for example, by the Center for Aquatic and Invasive  
124 Plants Aquatic Plant Information Retrieval System ([www.plants.ifas.ufl.edu/apirs](http://www.plants.ifas.ufl.edu/apirs)). It is  
125 probable that the freshwater macrophyte  $\gamma$ -diversity of the Nearctic is nearly completely  
126 described.

127 In this study we examined variation in river vascular freshwater macrophyte community  
128 characteristics, and their potential spatio-environmental drivers, on a broad intercontinental  
129 scale, comparing warm regions of the New and Old World. Specifically, we targeted one  
130 widespread type of river ecosystem, namely calcareous (“hardwater”) rivers and their  
131 associated high-connectivity riverine static or slow-flowing waterbodies, occurring in warm-  
132 temperate to tropical regions of America and Africa.

133 We define “hardwater systems” as minimally having a mean calcium carbonate  
134 concentration ( $\text{CaCO}_3$ )  $>10 \text{ mg L}^{-1}$  (approximately  $>200 \mu\text{Eq L}^{-1}$ ), or bicarbonate  
135 concentration (alkalinity:  $\text{HCO}_3^-$ )  $>12.2 \text{ mg L}^{-1}$  (approximately  $>200 \mu\text{Eq L}^{-1}$ ): following Moyle  
136 (1945) and Tapia Grimaldo (2013). Calcareous rivers may have much greater hardness than  
137 these minimal values; bicarbonate concentrations  $>4000 \mu\text{Eq L}^{-1}$  were recorded at several  
138 sites in our study. Hardwater rivers arise on a range of catchment geologies, including  
139 karstic limestone, softer calcareous rocks such as chalk, gypsum and certain types of  
140 sandstone, and calcium-rich alluvial soils (Tapia Grimaldo, 2013). All of these geologies  
141 occurred within the set of sites examined here.

142 In this study we tested the hypothesis that significant differences in macrophyte community  
143 structure exist between calcareous warm-water rivers (and their associated high-connectivity  
144 waterbodies) located in warm-temperate to tropical regions of the New World, versus those  
145 in the Old World, taking the Afrotropics as the target Old World region. Specifically, we  
146 examined differences in macrophyte diversity and community composition between these  
147 regions, in relation to a spatial variable (latitude) and a set of physico-chemical factors  
148 (altitude, pH, electrical conductivity, alkalinity and water flow regime) potentially influencing

149 these differences. No previous study has examined this issue, which is of added interest in  
150 the context of establishing baseline data to assess potential changes in river floras  
151 associated with global climate change and other human stressors. We expected to see  
152 differences between these macro-regions primarily because of differences in their physico-  
153 chemical characteristics (e.g., Payne, 1986). Historic geographical segregation between the  
154 regions, and variation in relevant biotic factors, were also considered likely to influence  
155 differences in macrophyte diversity and community composition when comparing African and  
156 American warm-water calcareous rivers.

157

## 158 **Methods**

### 159 *Study area*

160 A dataset consisting of 292 samples, from Africa (n = 208 samples) and America (n = 84),  
161 was collected from sites located on rivers, and associated waterbodies with high connectivity  
162 to the river system. Sites were primarily located in flowing river channels. These included  
163 main river channels, tributaries, and distributaries (channels which flow into or out of the  
164 main river, within its floodplain, depending on main river channel water level: an example  
165 within our dataset being the Baia River in the floodplain of the Upper Rio Paraná in Brasil:  
166 Varandas Martins *et al.*, 2013). There was a smaller component of sites in static to slow-  
167 flowing water channels closely associated with rivers (e.g. backwaters and spring runs); and  
168 floodplain riverine lakes, oxbows, and cenotes (sinkholes, produced from the collapse of  
169 limestone-bedrock, filled with groundwater derived from underground rivers), which are lentic  
170 but closely connected to the river channel.

171 Sites were selected which had hardwater conditions; macrophyte communities present;  
172 reflected the range of environmental conditions occurring across each target area; and were  
173 reasonably accessible. For safety reasons, some otherwise suitable sites were excluded in



174 Africa because dangerous animals were present. Within the boundaries of these criteria  
175 sampling sites were selected at random along rivers and their associated waterbodies.

176 In Africa study sites were located in:

177 (i) Zambia: 176 samples from 130 individual sites throughout the country. Tropical: centred  
178 on 13°S, 29°E (latitude range: 8.89090 - 17.8875°S), sampled 2006 – 2011. In Zambia only  
179 some sites were repeat-sampled in wet and dry seasons of a single year, or in different  
180 years during the study period (for more on this see Kennedy *et al.*, 2015, 2016);

181 (ii) Botswana: tropical: 21 sites in the Okavango Delta, centred on 18.8°S, 22.5°E (latitude  
182 range: 18.33908 - 19.57003°S); sampled 2006; and

183 (iii) South Africa: warm-temperate: 11 sites, in the Highveld area of the Vaal River, centred  
184 on 26.5°S, 29.5°E (latitude range: 26.36711 – 26.97082°S); sampled 2009 - 2010.

185 In America the study areas were in:

186 (i) USA (northern Florida): subtropical to warm-temperate: 27 sites centred on 29.5°N, 82°W  
187 (latitude range: 29.08102 - 30.83998°N); sampled 2011;

188 (ii) México (Yucatan Peninsula): tropical: 18 sites centred on 19°N, 88.5°W (latitude range:  
189 18.44031 – 21.56547°N); sampled 2011;

190 (iii) Trinidad: tropical: 17 sites centred on 10.6°N, 61.5°W (latitude range: 10.57670 –  
191 10.71050°N); sampled 2011;

192 (iv) Argentina, located near the confluence of the Río Paraguay and Middle Río Paraná, in  
193 the Provinces of Chaco and Corrientes (warm-temperate, with three sites centred on 27.4°S,  
194 58.7°W (latitude range: 27.245 – 27.460°S); sampled 2010); and

195 (v) three areas of Brasil (all sampled 2010): (a) Chapada Diamantina in the State of Bahia:  
196 tropical, with two sites centred on 12.4°S, 41°W; (b) the Upper Rio Paraná and its floodplain,  
197 in the States of Paraná and Mato Grosso do Sul: subtropical, with 17 sites centred on

198 23.5°S, 54°W; and (c) the Bonito/ Southern Pantanal area of the State of Mato Grosso do  
199 Sul: subtropical, with 11 sites centred on 21°S, 56.5°W (total latitude range for Brasil sites:  
200 12.4000 – 25.85909°S).

201 In the Northern Hemisphere, the total latitudinal range for sample sites was 20.26328°  
202 (ranging from a site in Trinidad at 10.57670°N, to a site in northern Florida at 30.83998°N).

203 In the Southern Hemisphere, the site closest to the Equator was located in northern Zambia  
204 (8.89088°S), and the furthest-south was a site in Argentina (27.45996°S), giving a latitudinal  
205 range of 18.56908°.

206 Sampling was typically conducted during periods when rivers were experiencing baseflow  
207 conditions, during the dry season. This was partly to facilitate access to sites. Dry-season  
208 sampling also minimised the possibility of post-flood changes in water chemistry skewing  
209 results. Some sites in Zambia were sampled during both wet and dry-seasons and  
210 substantial changes to water chemistry were observed following flood events (Kennedy *et*  
211 *al.*, 2008; Kennedy *et al.*, 2015, 2016). Individual samples from these repeat-sampled sites  
212 were, however, treated as discrete units, hence the effects of wet season conditions on  
213 analytical results were identifiable.

214

#### 215 *Biological and environmental data*

216 Data on macrophyte species presence (vascular species only were included in the study)  
217 and environmental parameters were collected by field survey, and supporting laboratory  
218 analysis of water samples, during 2006-2011, from standard 100 m stretches of each target  
219 waterbody. All survey data were personally collected by the authors, to ensure a robust level  
220 of standardised quality control for species identification and other field data collection.

221 Macrophyte surveys broadly followed the international standard EN 14184 (European  
222 Committee for Standardization, 2003), to collect qualitative data for macrophyte taxa

223 occurrence (submerged, floating and emergent: Chambers *et al.*, 2008) within each survey  
224 stretch. A standard macrophyte-sampling grapnel (attached to a 5 m long cord, and thrown  
225 from bank or boat as appropriate) was used where necessary as an aid to collection of  
226 submerged species. Nomenclature follows The Plant List ([www.theplantlist.org](http://www.theplantlist.org)). Herbarium  
227 voucher specimens were deposited with Coventry University (UK) and the Herbarium of the  
228 University of Morelos (HUMO), Universidad Autónoma del Estado de Morelos (México).  
229 Plants were identified to species level except where a lack of flowers, or other diagnostic  
230 structures, permitted identification only to genus or family level. All macrophyte taxa present  
231 at a site were used to calculate  $\alpha$ -diversity (S: number of taxa present per sample), but for  
232 other data-analysis purposes, only records identified to species level were utilised.  
233 Information on the distributional status of each species as endemic, native/ naturalised, or  
234 introduced/ invasive, within the Afrotropical and one or both of the Nearctic/ Neotropical  
235 biogeographic regions was obtained from various sources. These included e-Monocot  
236 (<http://e-monocot.org>); Flora Zambesiaca: (<http://apps.kew.org/efloras>); Flora of Zambia:  
237 [www.zambiaflora.com](http://www.zambiaflora.com); Flora of Botswana: [www.botswanaflora.com](http://www.botswanaflora.com)); GBIF (Global  
238 Biodiversity Information Facility): <http://www.gbif.org/species>; Flora acuática vascular del  
239 área focal Felipe Carrillo Puerto, Corredor Biológico Sian Ka'an-Calakmul, Quintana Roo,  
240 México (Bonilla-Barbosa, 2004): [http://www.gbif.org/dataset/7f7f1342-f762-11e1-a439-  
241 00145eb45e9a](http://www.gbif.org/dataset/7f7f1342-f762-11e1-a439-00145eb45e9a); MEXU/Colección de Plantas Acuáticas: [www.gbif.org/dataset/9606752e-  
242 f762-11e1-a439-00145eb45e9a](http://www.gbif.org/dataset/9606752e-f762-11e1-a439-00145eb45e9a); Amaral *et al.* (1998), Scremin-Dias *et al.* (1999), Pott & Pott  
243 (2000), Gerber *et al.* (2004), and Cook (2004).

244 Spatial and environmental variables used for this study included latitude: absolute ° (N or S  
245 of the Equator); and altitude (m above sea level, a.s.l.), recorded using a hand-held Garmin  
246 Etrex (or similar) Global Positioning System (GPS) instrument, and supplemented where  
247 necessary by reference to Global Earth or other large scale maps. A subjective assessment  
248 of flow (flow categories and approximate corresponding flow velocity intervals follow Lang &  
249 Murphy, 2012) was made on a four-point scale: 0 = static: (0 m s<sup>-1</sup>); 1 = slow flow

250 (approximately  $<0.2 \text{ m s}^{-1}$ ); 2 = moderate flow (approximately  $0.2 - 0.4 \text{ m s}^{-1}$ ); 3 = fast flow:  
251 “riffle” or white-water showing (approximately  $>0.4 \text{ m s}^{-1}$ ). Electrical conductivity (EC:  $\mu\text{S cm}^{-1}$ ) and pH were measured on-site, using a Schott 178 Handylab 264 meter, or similar  
252 instrument. Water samples were collected at each site (in an undisturbed sediment area) for  
253 subsequent laboratory measurement of alkalinity ( $\mu\text{Eq L}^{-1}$  bicarbonate), using the Gran  
254 alkalinity titration method (Neal, 2001).

256

### 257 *Statistical methods*

258 Two strategies were used in order to minimise sampling effects and make  $\gamma$ -diversity  
259 (“species pool”) comparable between continents (Melo *et al.*, 2007). The first was  
260 construction of rarefaction curves for American and African sites, and the second utilised the  
261 incidence-based Chao2 estimator (Chao, 1987; Colwell, 2013). R was used to carry out both  
262 analyses.

263 In order to assess species turnover between sites located in America and those in Africa,  $\beta$ -  
264 diversity (Koleff *et al.*, 2003) was measured using the Sorensen index ( $C_s$ ):

$$265 \quad C_s = 2j/(a + b)$$

266 where  $a$  = number of species present in samples surveyed in area  $a$ ;  $b$  = number of species  
267 present in samples surveyed in area  $b$ ; and  $j$  = number of species present in common in  
268 areas  $a$  and  $b$ . Low values for this index imply low commonality between the regional  
269 species-sets compared.

270 To assess variation in macrophyte community composition an ecologically-relevant  
271 classification of samples was generated, in terms of species present at each site, using the  
272 divisive clustering procedure TWINSpan (Hill, 1979). A matrix of samples x species for the  
273 full dataset was used, including only taxa identified to species level. This produced a set of  
274 end-groups of samples (stop-criterion for clustering sample division: division eigenvalue

275 >0.300), for which spatial, environmental and diversity variables were further compared  
276 using inferential statistics.

277 For inferential statistical testing, to compare mean values of response variables ( $\alpha$ -diversity,  
278 S; latitude; pH; altitude; alkalinity; electrical conductivity, EC; flow) measured at sites,  
279 between TWINSPAN sample-groups, variables were first assessed for normality using  
280 Ryan–Joiner testing, and all proved to meet the conditions of normality. Homogeneity of  
281 variance was then assessed using Levene’s Medians test, and only two variables (pH and  $\alpha$ -  
282 diversity, S) met the assumption of no significant difference in homogeneity of variance  
283 between datasets included in the test. For these two variables, one-factor analysis of  
284 variance (ANOVA), with *post-hoc* mean-separation using Tukey’s Least Significant  
285 Difference test was utilised. The remaining variables were assessed using the non-  
286 parametric Kruskal-Wallis procedure. Permutational multivariate ANOVA (PERMANOVA:  
287 Anderson, 2001) was used to test for significant differences in community composition  
288 composition across the TWINSPAN groups. In order to investigate the relative importance of  
289 latitude versus the measured environmental data in influencing community composition a  
290 variance partitioning exercise was carried out on the species presence-absence data, using  
291 distance-based redundancy analysis (db-RDA) based on Bray-Curtis distance (Anderson &  
292 McArdle, 2001). . Variance partitioning is a standard procedure (Borcard *et al.*,1992) used to  
293 determine the relative influence of different variables in shaping community composition. A  
294 number of db-RDA analyses were carried out: first a full model incorporating latitude and the  
295 available environmental data (pH, EC, alkalinity and altitude); second, the model was rerun  
296 with latitude as the covariable; then, third, another run with the environmental variables as  
297 covariables. By comparing the fractions of variance explained by each model it is possible to  
298 calculate the relative influence of latitude versus environment. Permutation tests were  
299 applied to assess the significance of the various models.

300 The above tests are multivariate and investigated the community composition data. In order  
301 to investigate the response in the univariate species richness data we used Boosted

302 Regression Trees (BRTs), which can cope with a combination of categorical and continuous  
303 data (De'ath, 2007). BRTs were employed to determine the factors that best predict variation  
304 in species richness across the full dataset, and for each continent separately. The approach  
305 of Elith *et al.* (2008) was employed to find the optimal number of trees. Tree complexity was  
306 set at three with a learning rate of 0.001, and with the bag fraction set at 0.75, meaning each  
307 individual tree was constructed using 75% of the data, with its predictive ability tested on the  
308 remaining 25% (Elith *et al.*, 2008). BRTs are excellent tools for finding patterns in large  
309 complex data sets, using thousands of small trees to find variables that (in this case) best  
310 predict species richness, but they do not provide a good means to visualise the data. Thus,  
311 we used a single univariate regression tree (De'ath, 2002), pruned using a cost-complexity  
312 measure, to show how the different explanatory variables relate to patterns in species  
313 richness. Indirect gradient analysis ordination, using non-metric multidimensional scaling  
314 (nMDS: with Bray-Curtis distance measures), and t-tests were also used in analyses of the  
315 dataset. Inferential tests were conducted using Excel (with the Real Statistics add-in  
316 package: [www.real-statistics.com/free-download/real-statistics-resource-pack](http://www.real-statistics.com/free-download/real-statistics-resource-pack)), and Minitab  
317 version 15.1.0. PERMANOVA, nMDS, and BRTs and regression tree analysis were all  
318 carried out in R (R Core Team, 2015). The vegan package was used for PERMANOVA,  
319 dbRDA, variance partitioning and nMDS (Oksanen *et al.*, 2016); the gbm package  
320 (Ridgeway, 2015: [www.cran.r-project.org/web/packages/gbm/gbm.pdf](http://www.cran.r-project.org/web/packages/gbm/gbm.pdf)) with additional code  
321 from Elith *et al.* (2008) for the BRTs; and rpart (Therneau *et al.*, 2015) for the regression  
322 trees. Where appropriate, outcomes were considered significant at  $p < 0.05$ .

323

## 324 **Results**

325 In total 378 individual macrophyte taxa were recorded: 154 from America and 242 from  
326 Africa (for full macrophyte records see Appendix S1; and for associated geositional and  
327 environmental data recorded for sampling sites see Appendix S2; both attached to the on-

328 line version of this article). Taxonomic resolution varied between individual countries (the  
329 strongest being Florida, Argentina and Zambia, with México and Trinidad the weakest,  
330 largely reflecting the availability of local literature and expertise available for aquatic  
331 macrophyte identification). Overall the total broke down to 291 taxa fully identified to species  
332 level, 49 identified to genus, and 38 only to family level.

333 The distinctness of the species composition of the New and Old World floras was compared  
334 using both our field data and pre-existing species records from all types of freshwater  
335 habitats (see sources for these additional records in Methods). While pre-existing species  
336 records from all freshwater habitats indicate that 144 (49.5%) of the 291 species that we  
337 found, co-occur in both the New and Old Worlds, our results suggest that there is a much  
338 greater degree of macrophyte species separation between the continents for calcareous  
339 river habitats surveyed in this study. At our sites, just 25 species (8.6%) occur at both  
340 American and African sites.

341 From our field data 156 species (53.6%) were found in African samples only, while for all  
342 freshwater habitats 80 species (27.5% of the total found in our survey) are recorded from the  
343 Afrotropics but not the Neotropical/ Nearctic regions. From our field data 110 species  
344 (37.8%) were found only in American samples, while for all freshwater habitats 67 (23.0%)  
345 species are recorded from the Neotropical/ Nearctic but not the Afrotropical region. The  
346 commonest species, in terms of number of samples in which they were recorded, were  
347 mostly those typical of African sites (Fig. 1). Only seven of these 25 common species co-  
348 occurred at sites in both continents.

349 Rarefaction plots of cumulative species records collected from the two continents (Fig. 2)  
350 approached an asymptote in both cases, and no further increments of the number of species  
351 were found, even when doubling the sampling effort by extrapolation. The estimated values  
352 and confidence intervals (CI) for total species richness ( $\gamma$ -diversity) produced using the  
353 Chao2 estimator were 208.6 (CI 95%: 208.06 – 213.52) and 86.0 (CI 95%: 84.28 – 98.33),

354 respectively for Africa and America. Taken together, these results provide evidence that the  
355 sampling effort in both continents was adequate to estimate values for the species pool, and  
356 in both cases were close to the real measured values for  $\gamma$ -diversity.

357 Mean  $\alpha$ -diversity, directly measured as number of taxa recorded at each sample-site  
358 (including taxa not identified fully to species level for each sample), did not significantly differ  
359 between America and Africa, with an average of about eight taxa per sample. Endemic  
360 species showed fairly similar proportional occurrences in both continents, but there was a  
361 higher proportion of introduced/ invasive species at sites in America, compared with Africa  
362 (Table 1).

363 Because there was no significant difference in  $\alpha$ -diversity between the two regions  
364 compared (and also because the data collected were qualitative records), a simple measure  
365 of  $\beta$ -diversity was appropriate for use with this dataset (Jost, 2007). The value of the  
366 Sorensen coefficient calculated for comparison of macrophyte species turnover between the  
367 two sample sets was low at  $C_s = 0.17$ , emphasising the dissimilarity between the floras  
368 present in warm-water calcareous river systems in Africa and America.

369 TWINSpan classification of the dataset gave seven end-groups of samples, labelled Groups  
370 A – G. These were produced with division eigenvalues in the range 0.347 - 0.780,  
371 suggesting reasonable to strong separation of groups based on the macrophyte species  
372 composition of their component samples. There were substantial differences in the primary  
373 floristic characteristics of the seven sample-groups (Table 2), and also for mean values of  
374 the six spatial and environmental variables measured, as well as for  $\alpha$ -diversity (Fig. 3).  
375 There was strong segregation between groups of samples located in Africa and in America,  
376 with only one sample-group (Group D) containing samples from sites located in both  
377 continents.

378 Analysis of the species data using PERMANOVA confirmed that the TWINSpan groups had  
379 significantly different community compositions. The results for all-sites combined were F:



380 16.54,  $R^2$ : 0.26,  $p < 0.001$ ; for African sites alone the corresponding outcome was  $F$ : 13.22,  
381  $R^2$ : 0.16,  $p < 0.001$ ; and for American sites alone:  $F$ : 10.08,  $R^2$ : 0.27,  $p < 0.001$ . A clear  
382 separation of sample-groups in nMDS ordination space was also apparent (see ordination  
383 plots provided as Appendix S3 in Supporting File 3, attached to the on-line version of this  
384 article) for all-sites, African sites, and American sites, but particularly so for America, which  
385 further emphasises the differences in species-set supported by each group of samples.

386 The results suggest that both spatial and environmental factors may act as drivers of  
387 macrophyte community composition present at the seven sets of survey sites making up the  
388 TWINSPAN sample-groups, given the significant and often substantial differences observed  
389 between TWINSPAN sample-groups for all six variables measured (Fig. 3a - f). The least  
390 variation was, however, seen for flow class, suggesting that this may be weaker than the  
391 other variables in driving differences between TWINSPAN sample-groups. Significant  
392 variation in  $\alpha$ -diversity also occurred between the seven sample-groups (Fig. 3g).

393 The outcome of the partial db-RDA analysis is in good agreement with these results. For the  
394 full data set of environmental variables (pH, conductivity, alkalinity and altitude) and latitude  
395 12.4% of the variation in the community composition data was explained. The variance  
396 uniquely attributable to the environmental variables was 6.4%, whereas the variance unique  
397 to latitude was only 2.9%, the remainder being shared. The corresponding values for %  
398 variance explained by the models for Africa alone were 10.4, 6.6 and 2.7% for full-model,  
399 unique to environment, and unique to latitude, respectively. For America the corresponding  
400 values were 22.1, 11.6 and 9.8% (alkalinity was not significant within the environmental data  
401 for America, but all other outcomes in these analyses were significant at  $p < 0.05$ ). This  
402 suggests that both latitude and environmental variables influence the community  
403 composition, with environment exerting a greater influence in this case.

404 The highest mean value for  $\alpha$ -diversity was seen in Group B, a small all-African sample-  
405 group, dominated by a set of samples from the Okavango Delta in Botswana. Indicators for

406 the group were a diverse set of Afrotropical native and endemic species (Table 2). Fig. 3  
407 shows that this set of samples was (within the range of values covered by this study)  
408 characterised by low pH and conductivity, intermediate latitude, high altitude, moderate flow,  
409 and fairly low alkalinity (similar to that of three other groups, with a mean of c. 1000  $\mu\text{Eq L}^{-1}$ ,  
410 indicative of intermediate-hardwater conditions, as defined by Tapia Grimaldo, 2013).

411 In contrast the sample-group with lowest  $\alpha$ -diversity, Group A, only contained Neotropical  
412 samples, all from Brasil. Indicators for this group (Table 2) consisted of one species native in  
413 America, one endemic to the Neotropics, and one invasive in the Neotropics. This group was  
414 characterised (Fig. 3) by low conductivity but quite high pH, and had the second highest  
415 mean latitude of the seven sample-groups. Sites in this group tended to be located at fairly  
416 low altitude. Flow was usually moderate to fast, and the group average for alkalinity was  
417 higher than for Group B, at 1500 – 2000  $\mu\text{Eq L}^{-1}$ , though this still suggests that most sites  
418 were of intermediate-hardwater status (Tapia Grimaldo, 2013).

419 The remaining groups, of intermediate  $\alpha$ -diversity, showed quite substantial variability in  
420 mean environmental characteristics. For example, Groups F and G were made up of mainly  
421 low-lying sites with high mean conductivity (in some cases impacted by marine saline  
422 influences, producing very high conductivity values), and rather high pH, located around the  
423 Caribbean, together with a few sites further south in South America (Fig. 3). These groups  
424 had a quite different macrophyte community from the other five sample-groups, with a mix, in  
425 both cases, of samples from three or four Neotropical/ Nearctic countries. The indicator  
426 species, however (Table 2), suggest a clear difference in vegetation between the two  
427 sample-groups, with Group G (dominated by Florida sites, with sub-tropical to warm-  
428 temperate conditions) being indicated by a pair of species native to both the Nearctic and  
429 Neotropics. In contrast, lower-latitude tropical American sites made up Group F, mostly from  
430 México and Trinidad, and was indicated by four species different from those of Group G: two  
431 endemic to the Neotropics/ Nearctic, plus two grass species, one native and the other  
432 invasive in both American bioregions.

433 Single regression tree dendrograms (Fig. 4) for all-sites combined, show average species  
434 richness and the number of sites (n) in the “leaves” (end member-groups), with and without  
435 latitude included (as a spatial variable, latitude summarises the influence of many other  
436 factors, which may have a direct effect on plants, acting across the latitudinal range). When  
437 latitude is included it dominates the tree, explaining a high proportion of the variance, and  
438 tending to mask the influence of the environmental variables in driving species richness.  
439 When the spatial variable is excluded, the principal environmental variables seen to drive  $\alpha$ -  
440 diversity in this classification are pH, altitude, electrical conductivity and alkalinity. In keeping  
441 with the outcome hinted at by the inferential statistical analysis exercise, above, flow was not  
442 shown by the regression analysis to be of importance as an environmental driver of  
443 macrophyte  $\alpha$ -diversity in this dataset.

444 Partial dependence plots show, in detail, the effect of predictors on the response variable,  
445 after taking into account the average effects of all other predictors in the model. So these  
446 plots should describe variation unique to the variable in question, though where strong  
447 interactions or correlations exist this is less reliable (Elith *et al.* 2008). The outcomes of the  
448 BRT analyses (Figs. 5 – 7) provide information to permit determination of the best predictors  
449 of macrophyte  $\alpha$ -diversity (species richness: S) respectively for the all-sites, American, and  
450 African datasets (% deviance explained: 19% for the whole dataset; 26% for the African  
451 sites, and 24% for the American sites). The plots show that there are different numbers of  
452 influential predictors of species richness for the three datasets, and that their relative  
453 importance and the “shape” of their influence (across the gradient-range covered by each  
454 individual predictor variable) both vary. The plots also show the proportion of the explained  
455 variance that each variable accounts for in the data, and the shape of the relationship -  
456 smoothed with the dashed line in the diagrams. The models performed well in terms of  
457 observed vs. predicted outcomes (0.70 - 0.80 correlation), with cross validated correlation  
458 scores (which compare model predictions with observations left out when building the  
459 model) of 0.48 - 0.50 for the three models.

460 The same variables (latitude, EC and altitude) are the most influential for both the African  
461 and American data sets, with pH and alkalinity also both influential in the African dataset.  
462 What is clear from the plots presented here (Figs. 5 – 7) is that latitude is a powerful  
463 predictor of species richness, but there were observable differences in response between  
464 the continents. For Africa, species richness increases gradually with distance from the  
465 Equator, starting from a relatively low latitude, whereas in America there was no discernible  
466 latitude effect until 20°, whereupon there was a rapid increase in richness. There is also a  
467 large difference in the response curve for altitude. The largest change in diversity for  
468 America occurs below 300 m a.s.l., whilst the lowest-altitude site in the African dataset is  
469 around 400m a.s.l. Electrical conductivity is an additional important factor shaping  
470 macrophyte species richness in this dataset. The partial dependence plots show that in  
471 America it is at the low end of the EC gradient that the influence on richness is greatest, with  
472 rising EC corresponding to higher macrophyte species richness. For the African data the  
473 pattern is less clear, but an increase is evident in the non-smoothed data.

474

## 475 **Discussion**

476 Comparisons of diversity and community composition of macrophytes in warm-water  
477 calcareous river systems within the two continents provided evidence that Africa and  
478 America differ in several ways. For macrophyte diversity, scale of analysis is important.  
479 Large scale diversity ( $\gamma$ -diversity) is, on our current evidence (though we think that may  
480 change when additional sites, outwith the envelope of site-conditions examined here, are  
481 sampled in the future) much higher in these systems in Africa than in America, and this  
482 difference cannot be accounted for by sampling effects. However, at local scale ( $\alpha$ -diversity)  
483 there is similarity between the two regions (Table 1), and this was an unexpected result,  
484 given the substantial differences in physical and chemical characteristics of hardwater river  
485 habitats sampled in the two continents (e.g., Payne, 1986; also our data presented here).

486 For example, we observed much greater variation in range of electrical conductivity within  
487 the American sites, compared with Africa; while altitude showed generally higher values  
488 within African sites, compared with America. While surprising, this result for  $\alpha$ -diversity is  
489 robust, given the consistency of both sampling strategy and sampling team, across the  
490 survey sites in both continents.

491 There is quite strong evidence for significant variation in  $\alpha$ -diversity between the main  
492 macrophyte community-types indicated by the TWINSPAN sample-classification (Fig. 3),  
493 while the results of PERMANOVA, nMDS sample ordination, and partial db-RDA analysis  
494 confirmed the observed species compositional variation across the TWINSPAN groups. The  
495 variation in macrophyte community between the two continents was substantial. Only one  
496 sample-group (TWINSPAN Group D), contained samples from both Africa and America.

497 Evidence from the species distribution literature, and online distributional databases, for the  
498 species found in our survey (see Methods for sources utilised), indicates substantial overlap  
499 for their distributions (between all freshwater habitats combined) in the Afrotropics and  
500 Neotropics/ Nearctic. However, we found that this was not the case for these plants in warm-  
501 water calcareous river habitats in the two continents, with most of the species recorded  
502 being found at sites in only one or the other continent, and with only a small proportion of  
503 species in common between them. There is, of course, no *a priori* reason why we should  
504 expect the same  $\gamma$ -diversity pattern to occur in all individual types of freshwater habitat, and  
505 on our evidence warm-water calcareous rivers show substantial differences in species pool  
506 (both in diversity and species presence), between America and Africa. This small proportion  
507 of species co-occurring in both continents suggests that, at least for the type of ecosystem  
508 studied here, it is perhaps not the case that “aquatic vascular plants generally show broad  
509 distributional ranges”, as was suggested by Santamaría (2002).

510 Our thoughts on this are further supported by the level of endemism (at regional level)  
511 observed in the dataset, which offers a partial explanation for our results. Approximately one  
512 third of the species that we recorded from each continent were endemic either to the

513 Afrotropics or the Neotropics/ Nearctic (Table 1), and so by definition do not occur in both  
514 continents.

515 Differences in species niche-breadth may provide a second clue. Niche-breadth values have  
516 been calculated (from data collected in Zambia) for 44 of the species found in our survey  
517 (Kennedy *et al.*, 2017). Excluding the endemics, it is notable that none of the eight species  
518 present in our dataset which were considered by these authors to have narrow niche-  
519 breadth (e.g. *Thelypteris confluens* (Thunb.) C.V. Morton, *Tristicha trifaria* (Bory ex Willd.)  
520 Spreng.) co-occurred at our survey sites in both Africa and America. In contrast, nine of 31  
521 species that were allocated by Kennedy *et al.* (2017) to intermediate/ broad niche-breadth  
522 status (e.g. *Ceratophyllum demersum* L., *Cyperus difformis* L.) were found at our survey  
523 sites in both continents. It is possible that generalist species, with greater niche-breadth  
524 (implying a wide tolerance of habitat conditions, and relatively good dispersal abilities, for  
525 example via long-distance endo- and exozoochory, utilising migratory waterfowl: e.g. Agami  
526 & Waisel, 1986; Clausen *et al.*, 2002; Santamaría, 2002; Coughlan *et al.*, 2017) are likely to  
527 have a reasonably high chance of finding suitable conditions for colonization in warm-water  
528 calcareous river habitats in both continents. In contrast, narrow-niche species by definition  
529 tend to have more specialist survival strategies (Grime, 1979) and narrower ecological  
530 tolerances, likely including traits influencing reproductive and dispersal capability, and  
531 potentially limiting range size. Recent evidence for this in freshwater organisms, including  
532 macrophytes, is provided by Slatyer *et al.* (2013) and Kennedy *et al.* (2017). Such species  
533 may, for example, utilise specialised reproductive and dispersal strategies (e.g. underwater  
534 pollination; vegetative propagule dispersal mechanisms: Sculthorpe, 1967; Smits *et al.*,  
535 1989; Barrat-Segretain, 1996; Donald, 1996; Wingfield & Murphy, 2006; Akasaka &  
536 Takamura, 2011; Redekop *et al.*, 2016), of possible low efficiency in promoting long-distance  
537 dispersal on a broad-scale planetary basis. In turn, this makes it less likely that these  
538 species will be present at geographically widely-separated locations. It logically follows that  
539 these specialist-strategy species may have more difficulty than generalists in finding

540 appropriately-similar locations for colonisation in both African and American warm-water  
541 calcareous rivers.

542 Although we do not consider here vicariance factors associated with ancestral  
543 phytogeographical influences on current macrophyte distributions (such as those associated  
544 with impacts of glaciation events etc.), which are certainly important, but well covered  
545 elsewhere in the literature (e.g., Santamaría, 2002; Les *et al.* 2003; Nies & Reusch, 2005;  
546 Chen *et al.*, 2012a, b; Zhu *et al.*, 2015), we do think that more recent actions related to  
547 human activities may be relevant. For example, the proportion of invasive/ introduced  
548 species in American rivers was substantially higher (at around 9% of total  $\gamma$ -diversity) than in  
549 Africa (Table 1), and this is a further likely contribution to explaining the observed community  
550 composition differences between the continents. A good example is invasive *Hydrilla*  
551 *verticillata* (L.f.) Royle (thought to be native to the Palearctic/ Oriental bioregions (Zhu *et al.*,  
552 2015), though there are also some possibly-native records from Africa:  
553 <http://www.cabi.org/isc/datasheet/28170>). This plant was found at 14 sites in our survey, all  
554 in Florida (though it has also recently been recorded as invasive in hardwater river sites in  
555 one of the areas of Brasil (the Upper Rio Parana) that we sampled: e.g., Sousa *et al.*, 2010).  
556 This species was not present at any of the hardwater river sites sampled in Africa during the  
557 study period, though there is a single previous record from a Zambian calcareous river, the  
558 Kafue River in 1981: <http://www.gbif.org/occurrence/1140612468>.

559 For macrophytes, introductions are frequently related to aquarist activity, remediation,  
560 intentional release, and escape from managed environments, such as Botanic Gardens  
561 (Brundu, 2015). A recent worldwide survey (Crafton, 2015) also suggested that international  
562 trade is a further determinant of invasive success. We suspect that these human-related  
563 activities are less intense in Africa than in America, possibly partially explaining differences  
564 in invasive species presence between calcareous rivers in the two continents.

565 At many sites, in all the countries of America and Africa examined here, macrophyte  $\alpha$ -  
566 diversity was low. Our results support previous studies which suggest that, in freshwater  
567 systems, local driving factors (chemical, physical or biological) seem to be of overriding  
568 importance in determining whether or not macrophyte diversity at an individual site is  
569 depressed below the optimal level within a given geographical region (e.g. Baattrup-  
570 Pedersen *et al.* 2006; Rolon & Matchik, 2006; Chappuis *et al.*, 2012; Lang & Murphy, 2012;  
571 Bando *et al.*, 2015; Kennedy *et al.*, 2015; Morandeira & Kandus, 2015; Schneider *et al.*,  
572 2015; Tapia Grimaldo *et al.*, 2016). Physical size of individual rivers, however, seemed to be  
573 of little importance in influencing  $\alpha$ -diversity. Despite their apparently-large potential area for  
574 colonisation, often large rivers are too deep, or their discharge is too great, or they are too  
575 turbid, to allow macrophytes to colonise further out into the channel than the marginal zone  
576 (e.g. Murphy *et al.*, 2003; Sousa *et al.*, 2011; Varandas Martins *et al.*, 2013).

577 The outcomes of our study emphasised the role of the spatial variable latitude in driving  
578 macrophyte diversity and community composition, despite the fairly limited latitudinal range  
579 (a band 18 - 20° of latitude wide, commencing about 8 - 10°N or S of the Equator, and  
580 running up to about 30°N or S) covered by our study. Latitude integrates a number of enviro-  
581 climatic variables, such as maximum and minimum annual temperature, precipitation, and  
582 evapotranspiration, which have individually previously been found to be good predictors of  
583 large-scale freshwater macrophyte diversity (e.g. Chappuis *et al.*, 2012; Tapia Grimaldo *et*  
584 *al.*, 2016).

585 A question remaining to be addressed is whether the variation in environmental  
586 heterogeneity seen between sampling sites located in the two continents might be  
587 influencing the observed findings of this study. For example, most of the African sites were  
588 located at high altitude whilst most sites in American river systems were located at low  
589 altitude (although for both altitude and all the other spatio-environmental variables studied  
590 there was an overlap in the range of values observed, when comparing sites from the two  
591 continents). This apparent sampling bias (at least in the case of altitude) is of course a



592 product of the differences in geography between the areas sampled. Florida and Yucatan,  
593 for example, have no high ground at all, whilst Zambia, Botswana and the South African  
594 Highveld are all upland regions. The question is whether results obtained from within the  
595 envelope of spatial (latitude) and physico-chemical (altitude, pH, conductivity, alkalinity, flow)  
596 conditions encompassing our sites apply only within that envelope, or are more widely  
597 applicable. Further work is clearly required to address that question, for example by  
598 attempting to find and sample low-lying calcareous rivers in Africa, and high-altitude  
599 calcareous rivers in America. At present, we conclude that our findings should be considered  
600 as being primarily applicable within the environmental envelopes encompassing the river  
601 systems studied, pending further research.

602 Our results support the findings of some, but not all, of the relevant previous studies in the  
603 literature which have examined large-scale drivers of freshwater macrophyte diversity. For  
604 example, Chappuis *et al.* (2012) found that latitude was a major driver of macrophyte  
605 diversity (in their case, country  $\gamma$ -diversity) across cool to warm-temperate Palearctic regions  
606 of Europe and North Africa. On a broader world scale, evidence is similarly provided by  
607 Crow (1993), from Central and North America, and Tapia Grimaldo *et al.* (2016), working  
608 with data from Africa and the British Isles, to suggest that both latitude and environmental  
609 factors play a role in predicting macrophyte diversity in freshwater systems. However, Viana  
610 *et al.* (2014) concluded that environmental and biogeographical factors, rather than latitude  
611 *per se*, drive aquatic plant species richness across Europe. Similarly, Alahuhta *et al.* (2017)  
612 suggested that, at a global scale, environmental heterogeneity (notably variability in altitude  
613 range within a region) plays the main role in driving macrophyte  $\beta$ -diversity, between lakes  
614 located in 21 different regions of the world. In our study, altitude was, in every case, third in  
615 importance (always behind latitude), in predictive value in this context (Figs. 5 – 7). Other  
616 physico-chemistry variables (pH, EC, alkalinity) showed less consistency across the  
617 analyses as being useful predictors of macrophyte  $\alpha$ -diversity in warm-water calcareous river  
618 systems.

619 The findings of all these studies, including our own, clearly emphasise the need for further  
620 work in this field (not just in warm-water calcareous rivers, but in freshwater habitats as a  
621 whole, planet-wide) to resolve the relative importance of spatial and environmental drivers in  
622 influencing macrophyte diversity. The importance of gaining improved baseline  
623 understanding of how such factors may affect freshwater macrophyte distributions and  
624 diversity can hardly be over-emphasised in the current context of global climate change.

625 A criticism of our study is that the snapshot environmental data mainly utilised here are  
626 unlikely to represent the longer-term mean values of individual variables at each site. Clearly  
627 it would be useful in any follow-up studies to include repeat-sampling to address this issue  
628 further. It is also likely that further work may show that other environmental factors, such as  
629 nutrient status (e.g., Kennedy *et al.*, 2016), as well as biotic interactions, including  
630 competition from non-native species (e.g. Michelan *et al.*, 2010; Sousa *et al.*, 2010), might  
631 be of importance in driving both diversity and community composition of warm-water  
632 calcareous river plants.

633 The evidence from the regression tree and boosted regression tree analysis here leads us to  
634 conclude that latitude is a significant, though non-linear and rather complex, spatial driver of  
635 hardwater river macrophyte  $\alpha$ -diversity, within the latitudinal range encompassed by this  
636 study. Altitude, pH, conductivity and alkalinity were also of importance in driving diversity,  
637 though varying in individual importance between Africa and America. The importance of  
638 latitude, even within a narrow range encompassing only low-latitude ecosystems, raises the  
639 possibility (see also Tapia Grimaldo *et al.*, 2016) that this factor may prove to be a driver of  
640 calcareous river macrophyte diversity across larger latitudinal gradients.

641

## 642 **Acknowledgements**

643 For funding we thank CONACYT (México), CEH (UK), the EC/ACP Science & Technology  
644 Programme (AFS/2009/219013), the UK DfID DeLPHE Programme, the Carnegie Trust for

645 the Universities of Scotland, and the UK DEFRA Darwin Programme. S.M.T. acknowledges  
646 the Brazilian National Council of Technological and Scientific Development (CNPq) for  
647 continuous funding through a Research Productivity Grant. We thank Eduardo Ribeiro  
648 Cunha (Universidade de Maringá) for undertaking the rarefaction analyses. We also thank all  
649 those who facilitated our fieldwork and other aspects of the study. Krzysztof Szoszkiewicz  
650 kindly criticised the ms prior to submission.

651

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908

909 **Supporting Information**

910 The original datasets collected for this study are available as additional Supporting Files  
911 attached to the on-line version of this article.

912

	Afrotropics	America
Total spp. ( $\gamma$ -diversity)	181	135
Native/ naturalised spp.	171	123
% native/ naturalised (of total spp.)	94.5	91.1
Endemic spp. (to Afrotropics or America, respectively)	57	47
% endemic spp. (of total spp.)	31.5	34.8
Introduced/invasive spp. (to Afrotropics or America, respectively)	10	12
% introduced/ invasive spp. (of total spp.)	5.5	8.9
Mean $\alpha$ -diversity (S: mean number of taxa recorded per site) $\pm$ standard error	$8.8 \pm 0.35^{\text{NS}}$	$7.9 \pm 0.50^{\text{NS}}$
Maximum S (recorded number of macrophyte taxa per site)	23	27

914

915 Table 1. Total macrophyte  $\gamma$ -, and mean and maximum  $\alpha$ -diversity recorded for sites  
916 surveyed in African (Afrotropical), and American (Neotropical/ Nearctic) countries, showing  
917 data for native/naturalised species (with percentages of endemic species for each region)  
918 and introduced/ invasive species. Comparison of mean S by t-test: not significant (NS:  $p$   
919  $>0.05$ )

920

TWINSPAN sample-group							
	A	B	C	D	E	F	G
Samples per group (n)	18	18	32	88	77	26	33
Eigenvalue for group production	0.780	0.519	0.519	0.347	0.347	0.681	0.681
Number of samples per country represented in group	Brasil (18)	Zambia (2), Botswana (15), South Africa (1)	Botswana (5), South Africa (2), Zambia (25)	Zambia (78), Botswana (1), South Africa (1), Trinidad (7)	Zambia (70), South Africa (7)	Trinidad (8), México (17), Brasil (1)	Florida (27), Trinidad (2), México (1), Argentina (3)
Indicator species	IA, IN <i>Brachiaria arrecta</i> (= <i>Urochloa arrecta</i> ) EN <i>Hymenachne pernambucensis</i> NaA, NaN <i>Oxycaryum cubense</i>	NaA, IN <i>Cyperus articulatus</i> EA <i>Cyperus pectinatus</i> NaA <i>Eleocharis dulcis</i> EA <i>Miscanthus junceus</i> NaA, IN <i>Nymphoides indica</i> subsp. <i>occidentalis</i> NaA, NaN <i>Oxycaryum cubense</i> NaA <i>Schoenoplectus corymbosus</i> NaA, NaN <i>Utricularia foliosa</i>	EA <i>Panicum subalbidum</i> NaA <i>Phragmites mauritianus</i>	NaA, IN <i>Nymphaea nouchali</i> var. <i>caerulea</i> NaA, IN <i>Panicum repens</i>	NaA, NaN <i>Commelina diffusa</i> NaA, IN <i>Cyperus alopecuroides</i> NaA, NaN <i>Cyperus involucratus</i> EA <i>Panicum subalbidum</i> NaA, IN <i>Pennisetum macrourum</i> NaA <i>Persicaria attenuata</i> subsp. <i>africana</i> NaA <i>Persicaria decipiens</i> NaA <i>Schoenoplectus corymbosus</i>	EN <i>Eleocharis cellulosa</i> EN <i>Fuirena simplex</i> IA, NaN <i>Paspalum notatum</i> IA, IN <i>Brachiaria arrecta</i> (= <i>Urochloa arrecta</i> )	NaA, NaN <i>Lemna aequinoctialis</i> IA, NN <i>Pontederia cordata</i>

Table 2. Characteristics of seven sample end-groups produced by TWINSPAN classification of 292 samples, using only fully-identified species. Indicator species for each group are shown together with information on distributional status of each species in Africa (Afrotropics) and America (Neotropics/



Nearctic combined): <sup>IA, IN</sup> introduced/ invasive to <sup>IA</sup> Afrotropics or <sup>IN</sup> Nearctic/ Neotropics; <sup>NaA, NaN</sup> native/naturalised to <sup>NaA</sup> Afrotropics or <sup>NaN</sup> Nearctic/  
Neotropics; <sup>EA, EN</sup> endemic to <sup>EA</sup> Afrotropics or <sup>EN</sup> Nearctic/ Neotropics

## Figure Legends

Figure 1. Percentage of (a) African and (b) American samples with records for each of 25 commonest species in the dataset ( $\geq 20$  records): Bra arr: *Brachiaria arrecta* (Poaceae) = *Urochloa arrecta*; Cer dem: *Ceratophyllum demersum* (Ceratophyllaceae); Com dif: *Commelina diffusa* (Commelinaceae); Cyp alo: *Cyperus alopecuroides* (Cyperaceae); Cyp art: *Cyperus articulatus* (Cyperaceae); Cyp pap: *Cyperus papyrus* (Cyperaceae); Eic cra: *Eichhornia crassipes* (Hydrocharitaceae); Ele dul: *Eleocharis dulcis* (Cyperaceae); Hyd umb: *Hydrocotyle umbellata* (Araliaceae); Lag ili: *Lagarosiphon ilicifolius* (Hydrocharitaceae); Lem aeq: *Lemna aequinoctialis* (Araceae); Lud ads: *Ludwigia adscendens* (Onagraceae); Naj hor: *Najas horrida* (Hydrocharticaceae); Nym noc: *Nymphaea nouchali* var. *caerulea* (Nymphaeaceae); Pan rep: *Panicum repens* (Poaceae); Pan sub: *Panicum subalbidum* (Poaceae); Per att: *Persicaria attenuata* (Polygonaceae); Per dec: *Persicaria decipiens* (Polygonaceae); Per hyd: *Persicaria hydropiper* (Polygonaceae); Per sen: *Persicaria senegalensis* (Polygonaceae); Phr mau: *Phragmites mauritianus* (Poaceae); Pot sch: *Potamogeton schweinfurthii* (Potamogetonaceae); Sal mol: *Salvinia molesta* (Salviniaceae) = *Salvinia adnata*; Sch cor: *Schoenoplectus corymbosus* (Cyperaceae); Typ dom: *Typha domingensis* (Typhaceae); Val ame: *Vallisneria americana* (Hydrocharitaceae).

Figure 2. Rarefaction plots estimating  $\gamma$  – diversity, using macrophyte taxa records from rivers and associated water bodies for 84 samples collected from 5 countries in America (2010 – 2011), and 208 samples from 3 countries in Africa (2006 – 2011). Black: Africa; grey: America.

Figure 3. Variation in mean values ( $\pm$  standard error) between TWINSpan sample-groups for spatial and environmental variables measured, and for  $\alpha$ -diversity: (a) electrical conductivity (EC:  $\mu\text{S cm}^{-1}$ ):  $p < 0.001$ ; (b) latitude (absolute  $^{\circ}$  north or south of Equator):  $p$

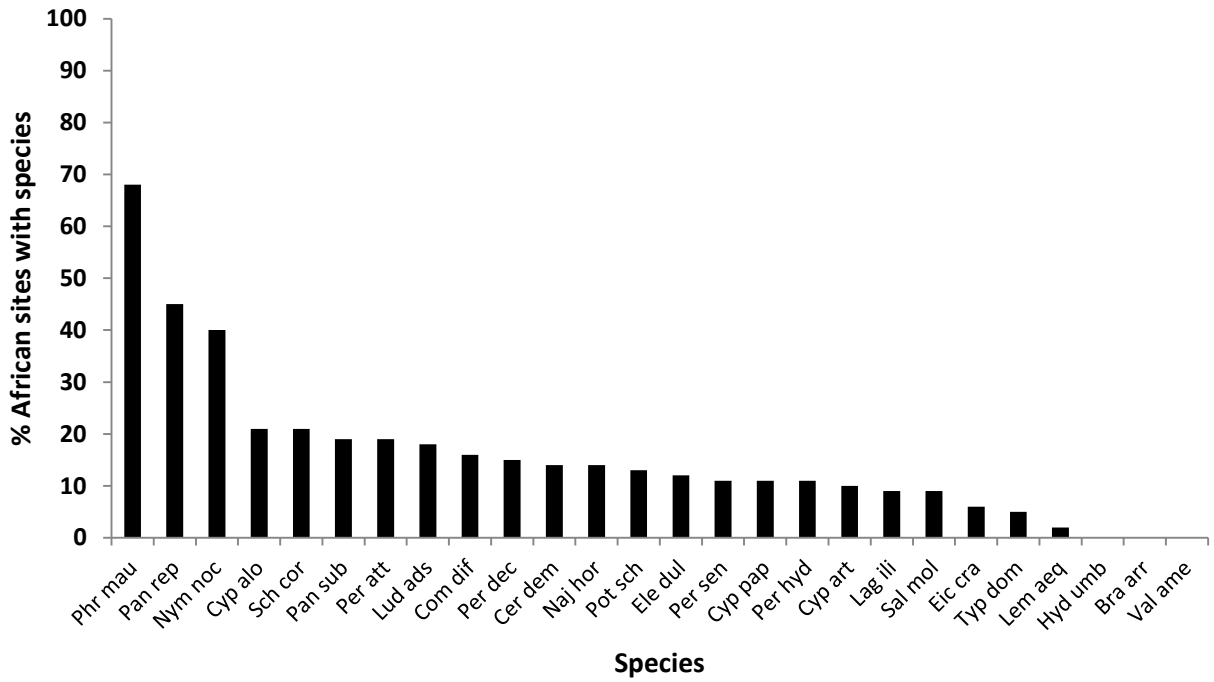
<0.001; (c) altitude (m above mean sea level: a.s.l.):  $p < 0.001$ ; (d) flow class (0: still - 3: fast-flowing):  $p < 0.05$ ; (e) pH:  $p < 0.001$ ,  $F$ : 8.513; (f) alkalinity ( $\mu\text{Eq L}^{-1}$ ):  $p < 0.001$ ; (g) S ( $\alpha$ -diversity: number of macrophyte taxa recorded per site):  $p < 0.001$ ,  $F$ : 15.371. Means for pH and S labelled with a letter in common do not significantly differ (ANOVA outcome with *a-posteriori* Tukey's mean separation test, significant at a minimum of  $p < 0.05$ ). Other variables analysed using Kruskal-Wallis test procedure: overall significance shown for outcome.

Figure 4. Regression tree dendrograms, for all-sites combined dataset, showing average species richness (S) and number of sites (n) in dendrogram end-groups ("leaves"): (a) spatial and environmental variables all included (i.e. latitude included in the analysis); (b) environmental variables only included (i.e. latitude omitted).

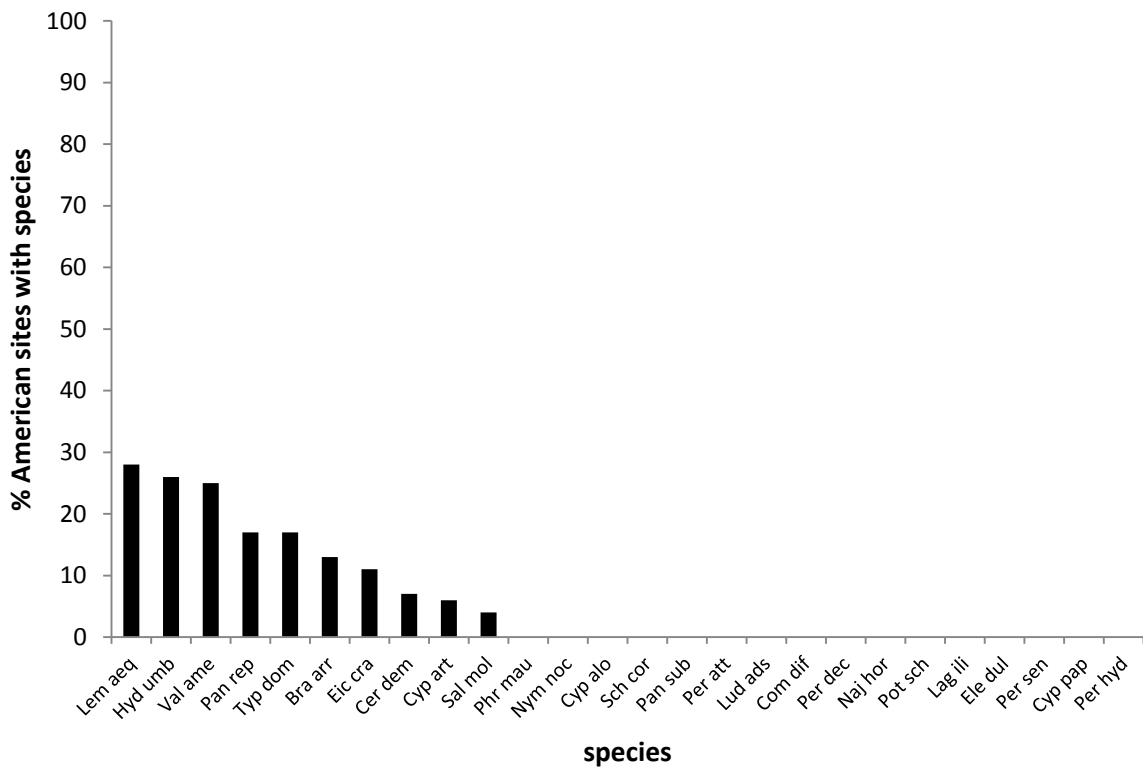
Figure 5. Boosted Regression Tree partial dependence plots of fitted function vs. observed values (primary values shown as tick marks on x-axis) for each of 5 spatial/ environmental variables significantly predicting macrophyte  $\alpha$ -diversity (species richness: S) for all-sites combined. Continuous line: fitted values; dashed line: smoothed fitted. Abbreviations: Lat: latitude (absolute  $^{\circ}$ ); Alt: altitude (m above sea level); Alk: alkalinity ( $\mu\text{Eq L}^{-1}$ ); EC: electrical conductivity ( $\mu\text{S cm}^{-1}$ ). Values given in brackets are proportion of the explained variance that each variable accounts for in the data.

Figure 6. Boosted Regression Tree partial dependence plots of fitted function vs. observed values for each of 5 spatial/ environmental variables significantly predicting macrophyte  $\alpha$ -diversity (species richness: S) for American sites. See caption to Fig. 5 for further details.

Figure 7. Boosted Regression Tree partial dependence plots of fitted function vs. observed values for each of 5 spatial/ environmental variables significantly predicting macrophyte  $\alpha$ -diversity (species richness: S) for African sites. See caption to Fig. 5 for further details.

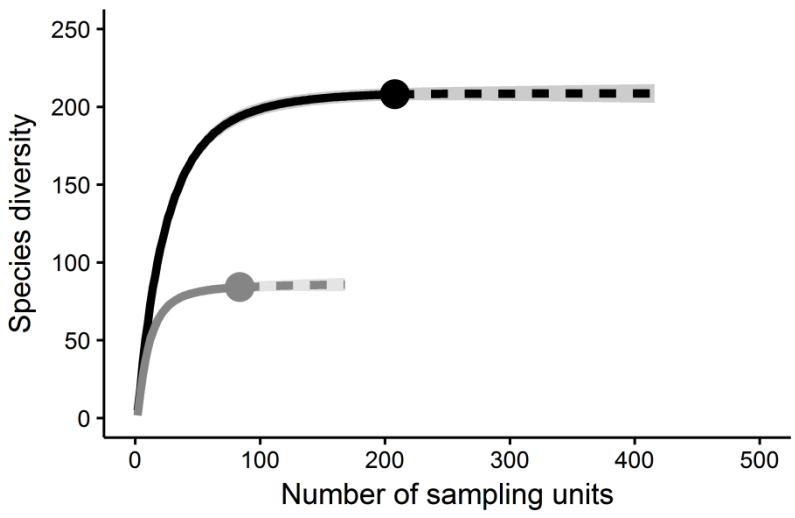


(a)

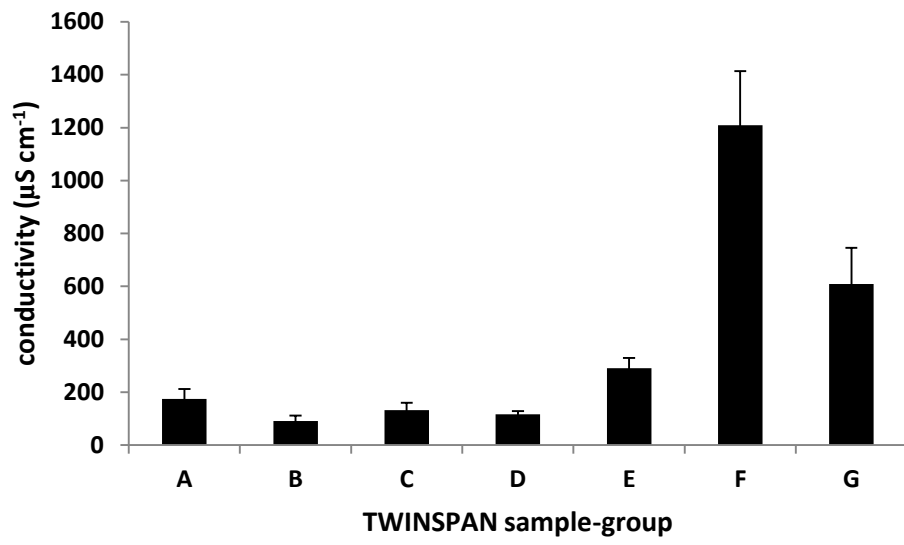


(b)

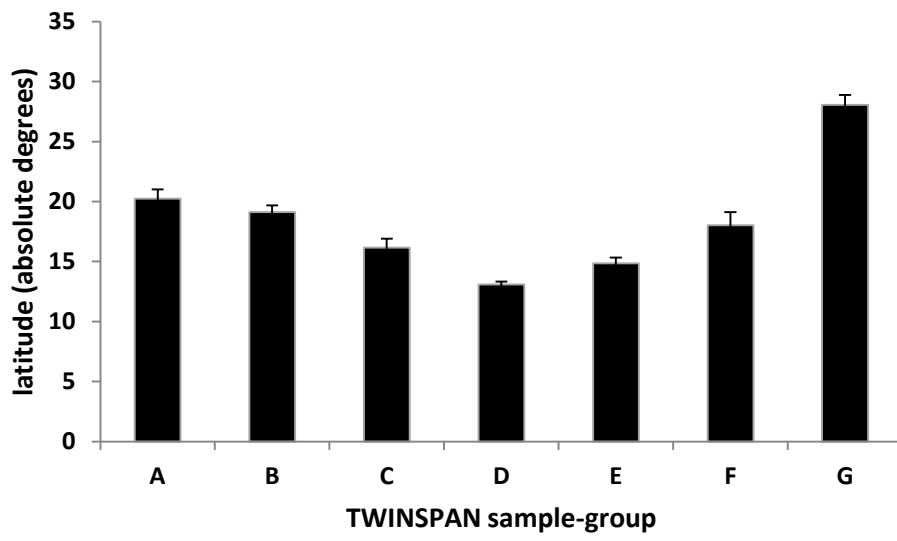
FIGURE 1



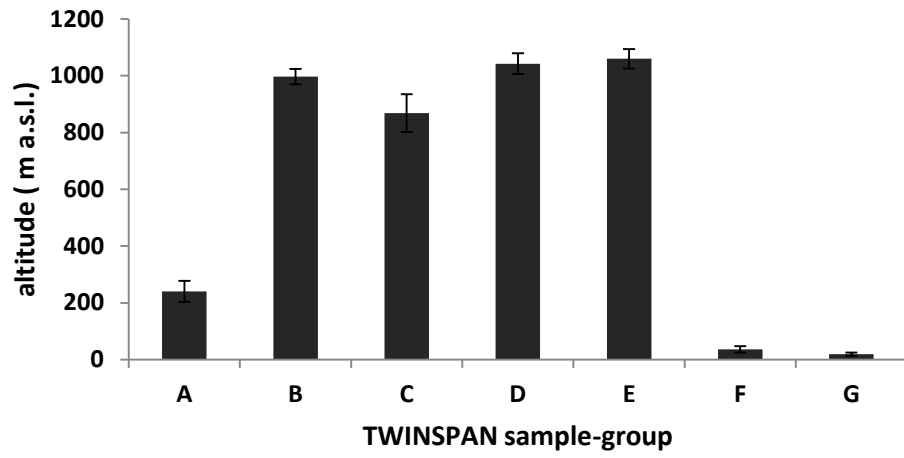
**FIGURE 2**



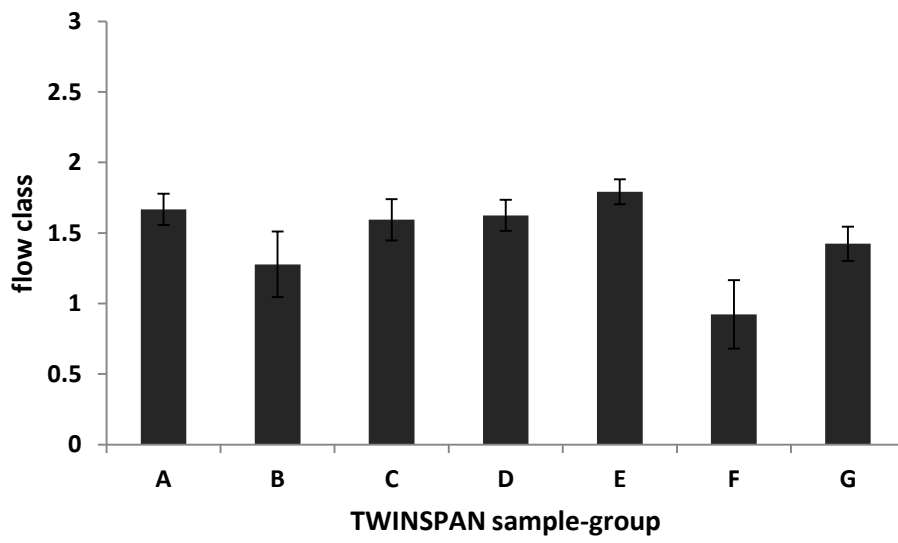
(a)



(b)

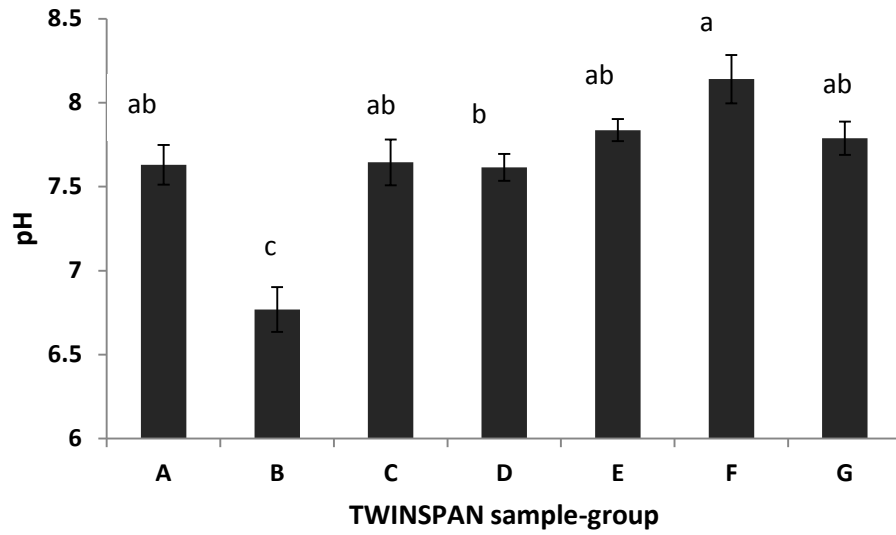


(c)

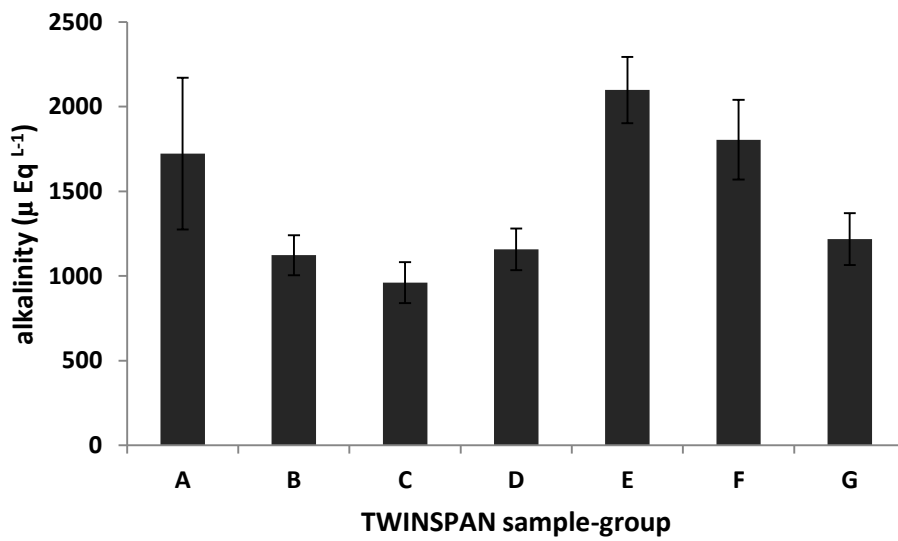


(d)

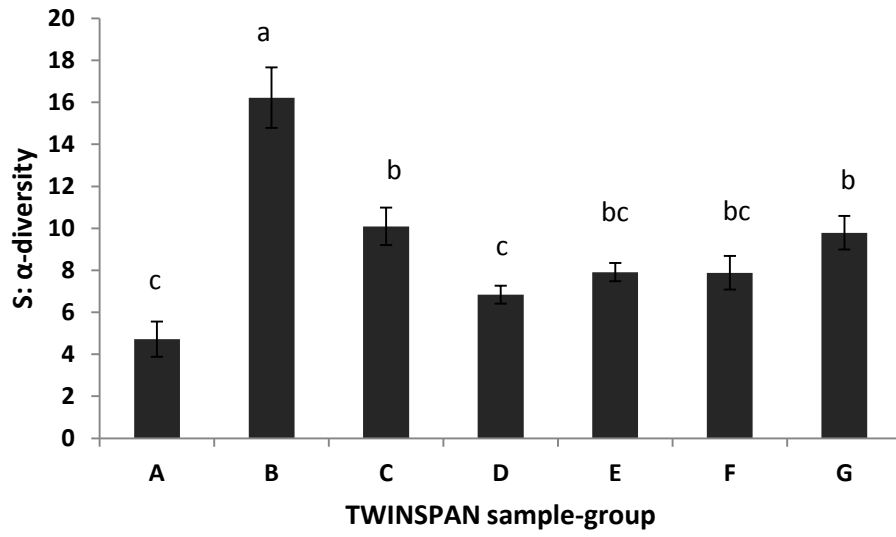




(e)



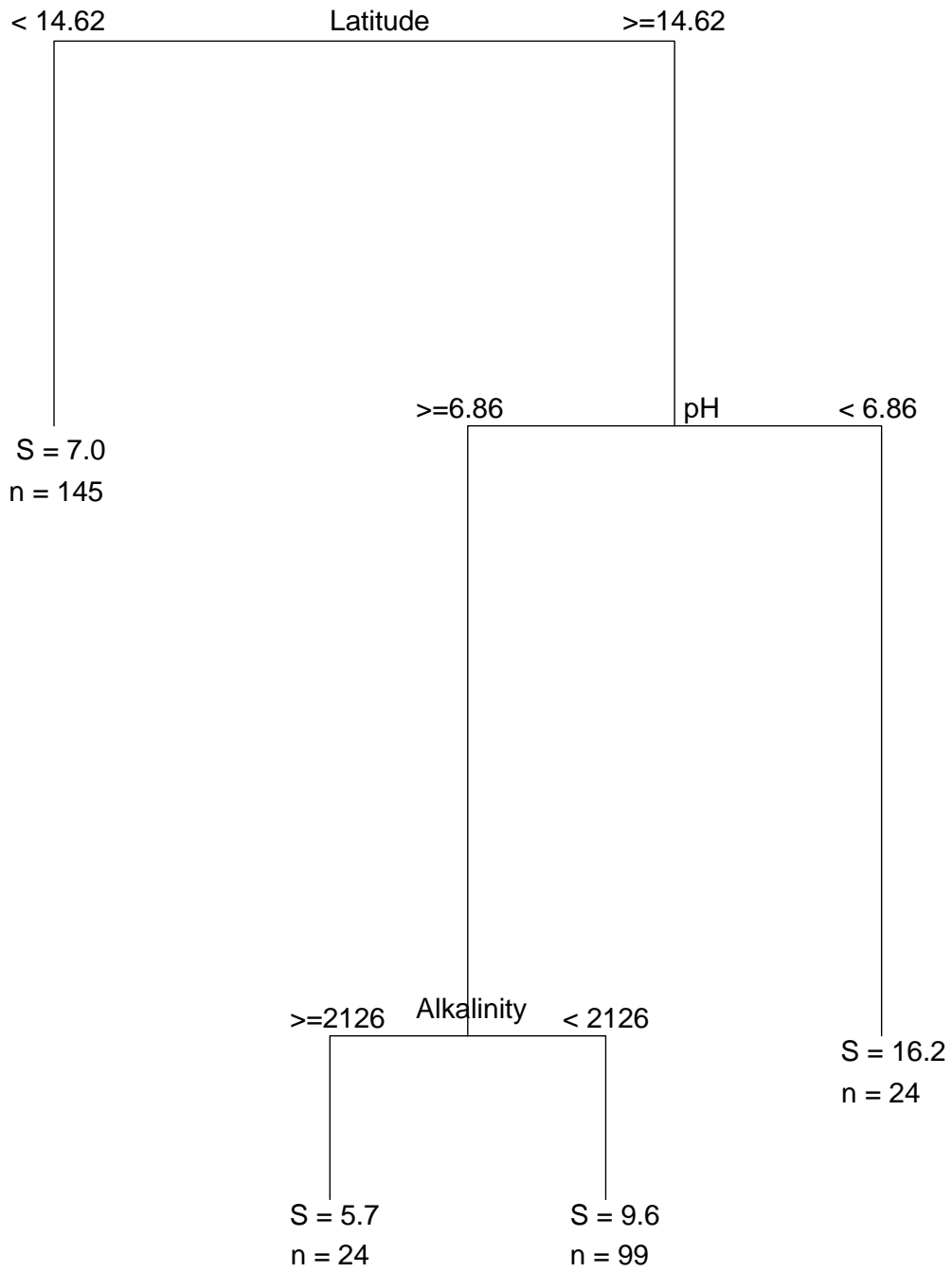
(f)



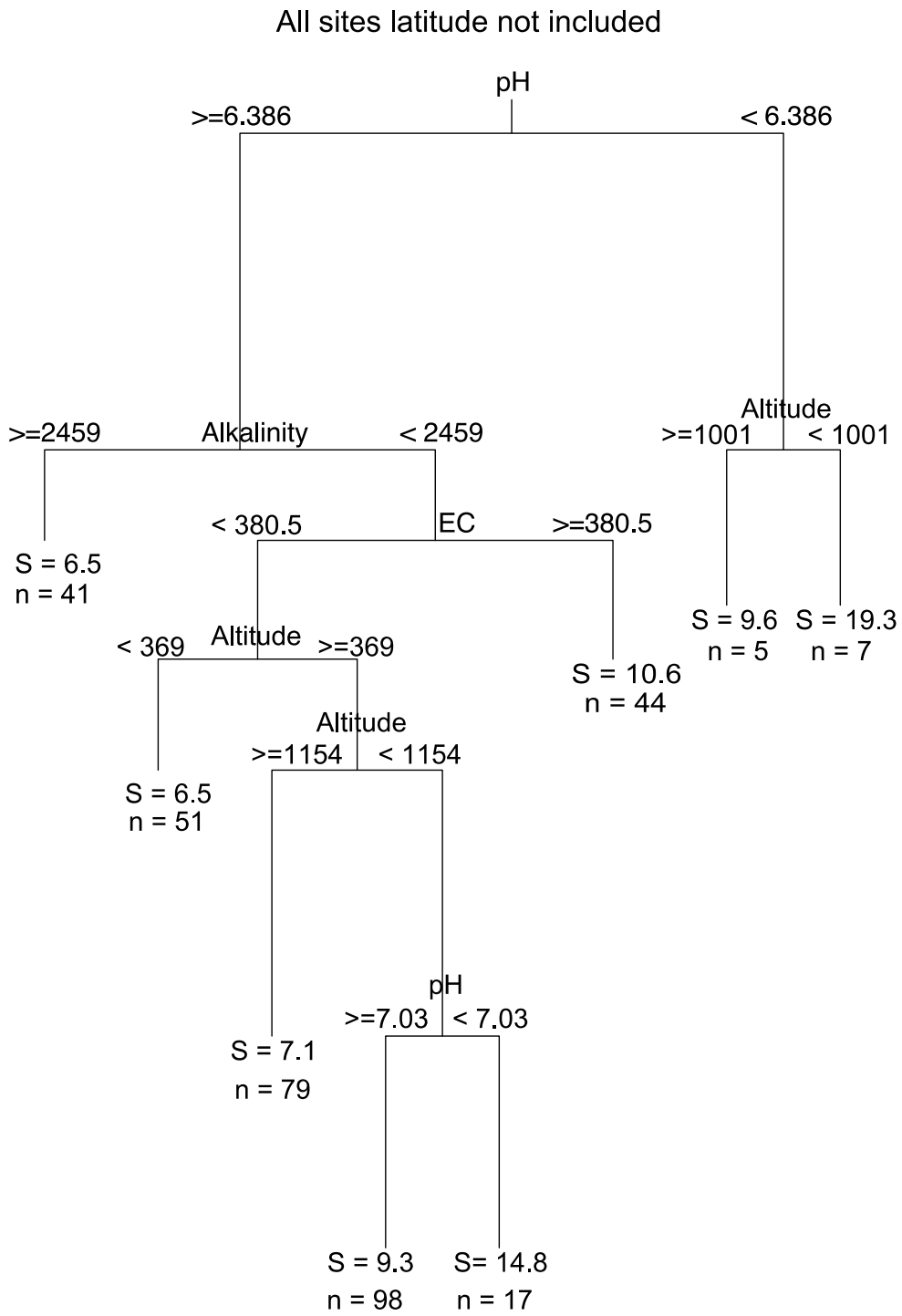
(g)

FIGURE 3

All Sites

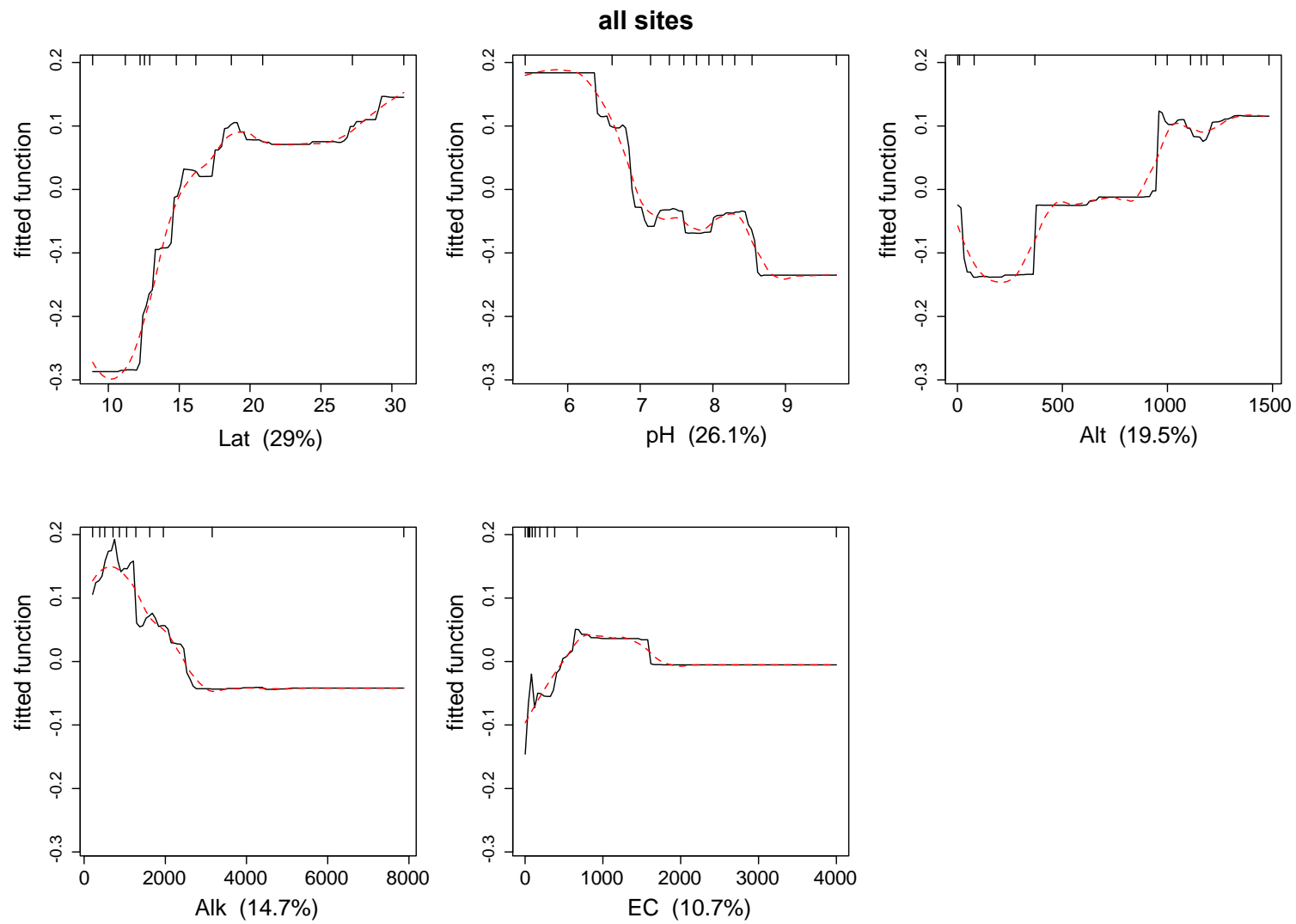


(a)

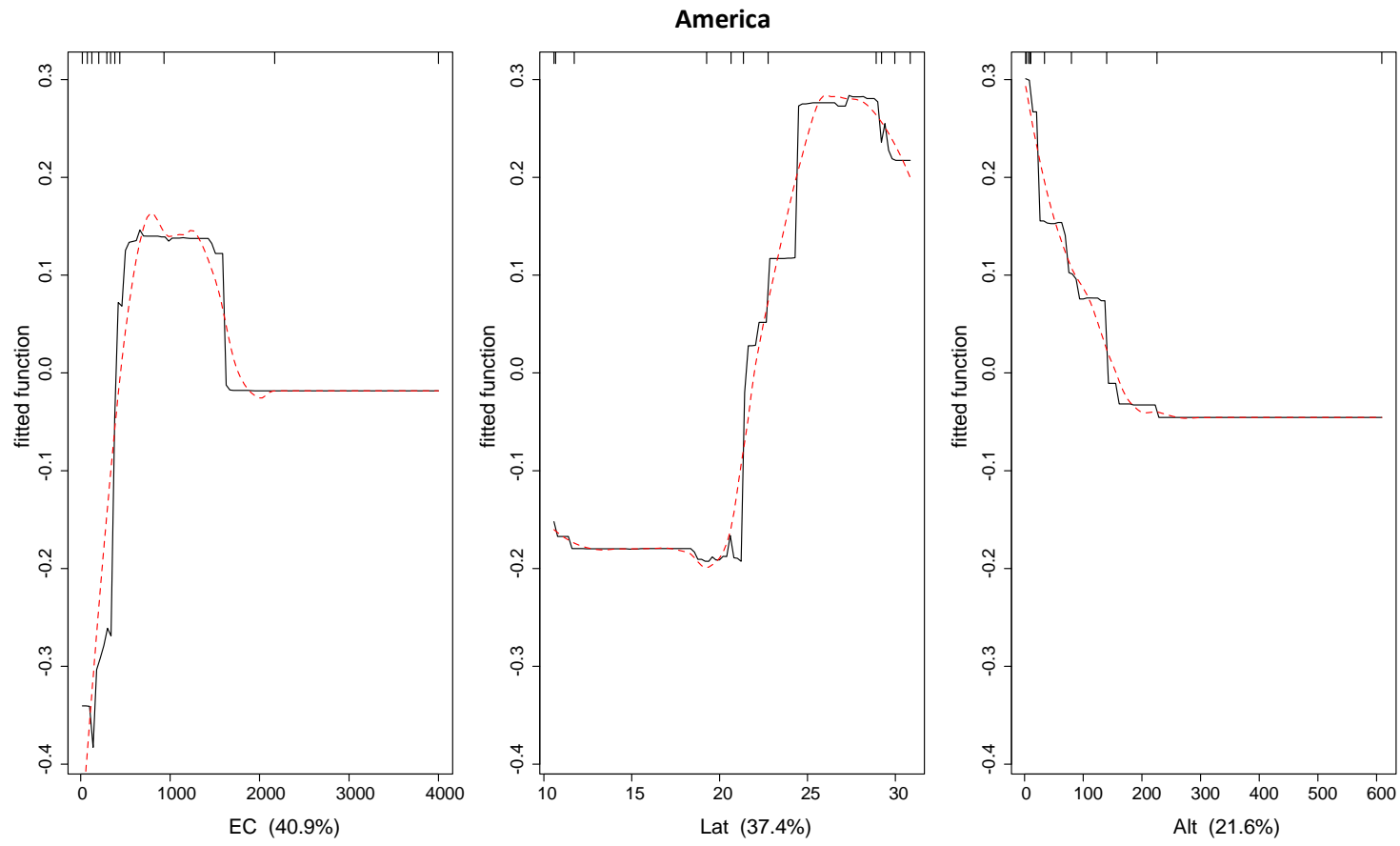


(b)

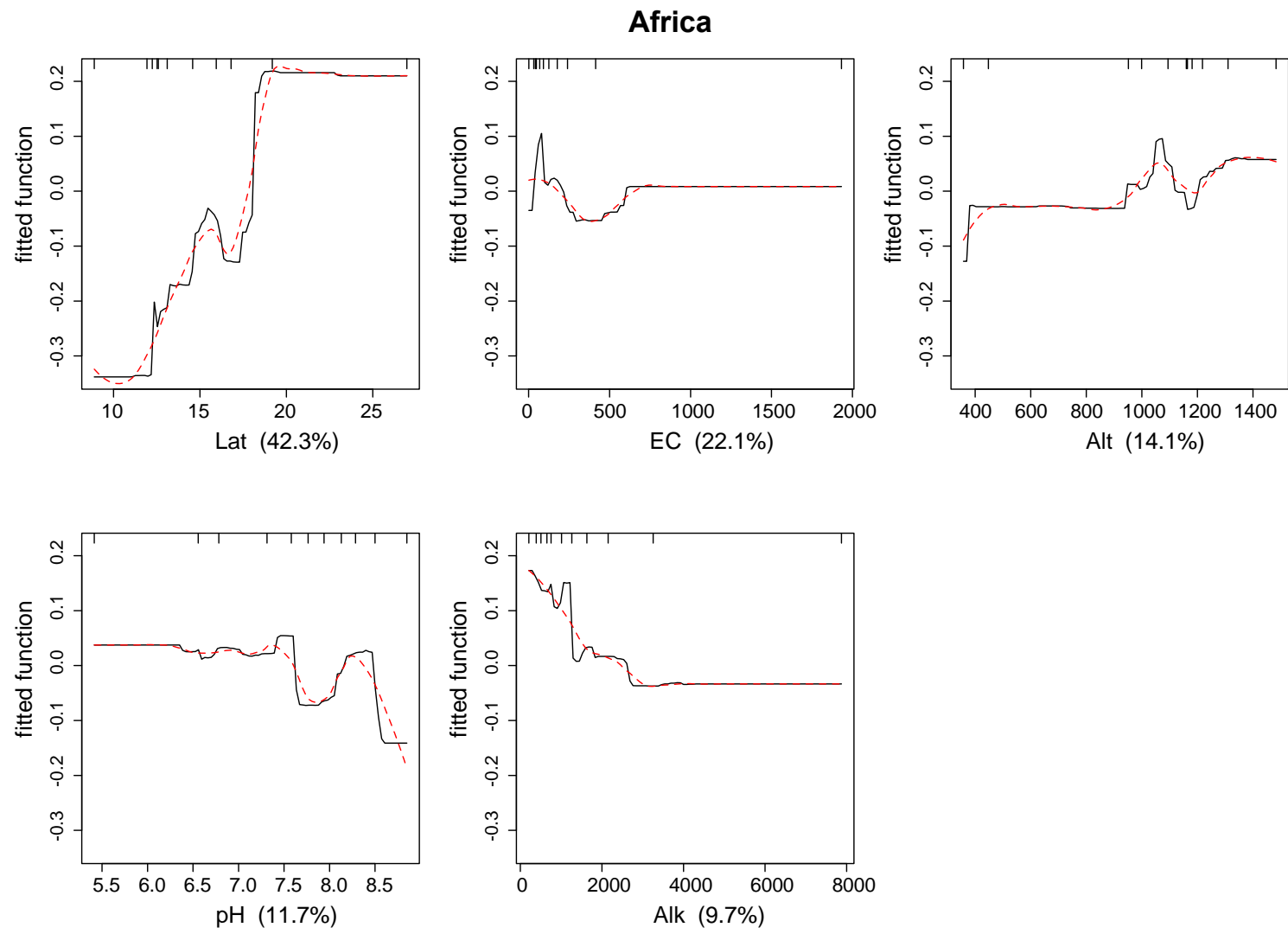
**FIGURE 4**



**FIGURE 5**



**FIGURE 6**



**FIGURE 7**