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1 **The bioelements, the elementome and the “biogeochemical niche”.**

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21 Running Head: **Elementome and biogeochemical niche.**

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25 **Keywords:**

26 Bio-elements, elementome, biogeochemical niche, ecological niche, phylogenetic distance,
27 sympatry, plasticity, ecosystem function and organization, evolution.

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32 **Abstract**

33 Every living creature on Earth is made of atoms of the various bioelements that are harnessed in
34 the construction of molecules, tissues, organisms and communities, as we know them. Organisms
35 need these bioelements in specific quantities and proportions to survive and grow. Distinct
36 species have different functions and life strategies, and have therefore developed distinct
37 structures and adopted a certain combination of metabolic and physiological processes. Each
38 species is thus also expected to have different requirements for each bioelement. We therefore
39 propose that a “biogeochemical niche” can be associated with the classical ecological niche of
40 each species. We show from field data examples that a biogeochemical niche is characterized by
41 a particular elementome defined as the content of all (or at least most) bioelements. The
42 differences in elementome among species are a function of taxonomy and phylogenetic distance,
43 sympatry (the bioelemental compositions should differ more among coexisting than among non-
44 coexisting species to avoid competitive pressure), and homeostasis with a continuum between
45 high homeostasis/low plasticity and low homeostasis/high plasticity. This proposed
46 biogeochemical niche hypothesis has the advantage relative to other associated theoretical niche

47 hypotheses that it can be easily characterized by actual quantification of a measurable trait: the
48 elementome of a given organism or a community, being potentially applicable across taxa and
49 habitats. The changes in bioelemental availability can determine genotypic selection and
50 therefore have a feedback on ecosystem function and organization, and, at the end, become
51 another driving factor of the evolution of life and the environment.

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56 **1. The elements of life and the construction of molecules, tissues, organisms and**
57 **communities**

58 All living beings are made of atoms of various bioelements (elements used by living organisms).
59 The most common bioelements are: hydrogen (H) 59%, oxygen (O) 24%, carbon (C) 11%,
60 nitrogen (N) 4%, phosphorus (P) 1% and sulfur (S) 0.1-1% (percentages of total number of
61 atoms in organisms) (Slade 2006). C, H, and O are the basis of organismic structure, i.e. water
62 and primary structural molecules, whereas N, P and S enter in the various biomolecules allowing
63 biochemical reactions. C, H and O from CO₂ and H₂O are converted into the skeleton of all types
64 of organic molecules by photo- and chemosynthesis. The molecular structures that control life
65 processes (e.g. RUBISCO or chlorophyll), that carry information (e.g. DNA and RNA), and that
66 control the storage and release of energy when and where necessary (e.g. ATP and NAD) all
67 incorporate N, P and in lower amounts S.

68 These six bioelements, however, are not the only bioelements that have important roles in
69 organisms (Slade, 2006). Other bioelements, normally present in low concentrations such as
70 potassium (K), magnesium (Mg), iron (Fe), calcium (Ca), molybdenum (Mo), manganese (Mn)
71 and zinc (Zn) have functions that are more specific. They are cofactors in catalytic reactions in
72 several biochemical and physiological processes. For example, in plants, K, Mg, Fe, S, Ca, and
73 Zn are involved in photosynthesis, respiration, regulation of ion balance in chloroplasts and
74 vacuoles, transport of sugar into the phloem, secondary metabolism, maintenance of internal
75 cellular osmotic or pH equilibria and balances, and strategies for stomatal control and avoidance
76 of water stress (Knight et al. 1991, Bourgis et al. 1999, Shaul 2002, Tripler et al. 2006, Peñuelas
77 and Sardans 2009, Baribault et al. 2012, Naples and Fisk 2010, Rivas-Ubach et al. 2012,
78 Peñuelas et al. 2013, Sardans and Peñuelas 2015). All these bioelements thus contribute
79 essentially to organismic functions, albeit to different degrees (Figure 1).

80 The concentrations and ratios of C, N and P have been the basis of ecological
81 stoichiometry (Elser et al. 2000a,b, 2001, Sardans et al. 2015, 2016). However, to link
82 bioelemental composition and stoichiometry with ecological traits in all types of organisms and
83 communities will thus require information about bioelemental composition beyond the C:N:P
84 concentrations and ratios (Peñuelas and Sardans 2009). The other bioelements such as S, K, Ca
85 or Mg are also crucial and can be limiting in some ecosystems (Naples and Fisk 2010, Baribault
86 et al. 2012, Lapenis et al. 2013). These other bioelements have been introduced in ecological
87 stoichiometric studies to link the elemental composition of individuals and species with their
88 ecological traits, such as drought resistance (related to K), light environment (related to Mg) or
89 the levels of N and S deposition (related to K, Ca, Mg and S) (Sardans et al., 2011;2012a;

90 Sardans et al., 2015). All these bioelements should thus be considered in ecological
91 stoichiometric studies.

92

93 **2. Elementome and biogeochemical niche.**

94 Species use these bioelements to different degrees based on their ecological strategy (example
95 for plants in Figure 2). Classic Hutchinson's (1957) description of species niche has been largely
96 applied to the main taxa from a point-of-view of distinct species in a site position in a "n-
97 dimensional hypervolume" formed by the values of determined environmental variables and/or
98 organisms traits. For instance, plant species have been classified in a hypervolume formed by
99 three axes, i.e. corresponding to Grime's 3 strategies: competitor, stress-tolerant and ruderal
100 (Grime 1977). In animals, the species-specific niche has been determined and located in
101 function of hypervolumes for n-axis of different values of energy expenditure (Wilson et al.,
102 2011), distinct diet sources (Araujo et al., 2007; Bearhop et al., 2004) or different elemental
103 composition (González et al., 2017, 2018). In microbial communities, genotype-niches have also
104 been described in function of the different availabilities of different nutrients (Pereira and Berry,
105 2017) or as a function of the values of diverse functional traits (Lennon et al., 2012) thus
106 describing multivariant spaces with a determined combination of nutrients availability or
107 functional traits where a determined genotype can exist. Any living species should thus tend to
108 optimize its overall function within a determined and space within this volume generated by a
109 combination of these axes describing strategies or biotic and/or abiotic variables while avoiding
110 the overlap with other coexisting species. Following these niche approach, we can hypothesize
111 that distinct species have different functions and life strategies and have therefore developed
112 distinct structures and adopted a certain combination of metabolic and physiological processes.

113 Therefore, each species (of all taxa types) also has different requirements for each bioelement,
114 and should thus tend to have its own elemental composition (Yu et al. 2011; Kattge et al. 2011;
115 Sardans and Peñuelas 2013, 2014; Sardans et al. 2015; Pierce et al. 2016; Bartrons et al., 2018).

116 We postulate that the more functionally separated two species are, the more different the
117 content of bioelements should be. This led us to propose a “biogeochemical niche” (BN) defined
118 as the multidimensional space defined by the concentrations of bioelements in individuals of a
119 given species (Peñuelas et al., 2008). This BN has been used in the recent years by other authors
120 such as Gonzalez et al. (2017) who proposed the “multidimensional stoichiometric niche” as the
121 specific position of different taxonomic/trophic groups in the three-dimensional space
122 determined by C, N, and P, and other elements as additional axes in their application of the BN
123 to animals (Gonzalez et al., 2018), as previously proposed for plants by Peñuelas et al. (2008),
124 and for plants and animals together by Bartrons et al (2018). BN is thus based on the species-
125 specific needs and use of bioelements in different amounts and proportions resulting from the
126 organismic structure and function in the given abiotic and biotic environmental space where the
127 species has evolved (Peñuelas et al. 2008, 2010). Each species should thus have an optimal (at
128 maximum fitness) equilibrium of bioelemental composition, resulting from its functional and
129 structural adaptations, equivalent to the optimum “elemental phenotype” (Jeyasingh et al., 2014).
130 The BN represents thus the species-specific strategies of growth and resource uptake and the
131 differences in soil-space occupation.

132 We propose that the processes underlying this species-specific elementome and the BN
133 are based on three basic complementary rules (Figures 3). First, distinct taxonomic groups
134 exhibit different elementomes, with larger differences as taxonomic distance and evolutionary
135 time increase. Second, at equilibrium, coexisting species tend to have distinct elementomes to

136 minimize competitive pressure. Third, trade-offs between adaptation to being competitive in a
 137 stable environment versus being successful in more fluctuating environments cause homeostasis
 138 and plasticity differences between species in a continuum of strategies. The BN distances among
 139 species should thus be a function of taxonomic difference, sympatry (coexistence), and
 140 homeostasis/plasticity and sympatry (Figure 3). The BN of taxa i is assumed to be the result of
 141 its taxonomical evolutionary determination and its capacity to respond to changes in external
 142 conditions, which is also partly taxonomically-determined. External changes include nutrient
 143 availability (including the level of atmospheric CO₂), changes in growing season duration,
 144 disturbance regimes, and, on longer time scales, changes in soil properties. This capacity to
 145 respond to environmental changes is, in turn, a function of taxonomy, sympatry, and
 146 homeostasis:

$$147 \quad BN_i = BN \text{ of taxa}_i = \text{Elementome}_{i1-n} = (C: N: P: K: S: Mg: Ca: \dots)_{i1-n} = BN_{\text{tax}_i} +$$

$$148 \quad \frac{\partial BN_{\text{tax}_i}}{\partial Env} = BN_{\text{tax}_i} + f(\text{Tax}_i, \text{Symp}_i, \text{Hom}_i) \quad (\text{Eq. 1})$$

149 where:

150 BN_{tax_i} = Biogeochemical niche explained by taxonomy of taxa i (measured in n
 151 individuals of taxa i). A n-dimensional hypervolume describes the fundamental BN of a species.
 152 The domain of this hypervolume function is restricted to a finite region corresponding to the
 153 possible values of bioelemental combinations.

$$154 \quad \frac{\partial BN_{\text{tax}_i}}{\partial Env} = \text{Changes in } BN_{\text{tax}_i} \text{ of taxa i explained by environmental changes}$$

155 tax_i = taxonomy of taxa i; Symp_i sympatry of taxa i with other competing taxa (j), and
 156 Hom_i = elemental homeostasis-plasticity of the taxa i

157 $BN_j = BN_{taxa_j} = \text{Elementome}_{j_{1-n}} = (C: N: P: K: S: Mg: Ca \dots)_j$

158 $BN_j = BN_{tax_j} + \frac{\partial BN_{tax_j}}{\partial Env} = BN_{tax_{j_{1-n}}} + f(\text{Tax}_j, \text{Symp}_j, \text{Hom}_j)$ (Eq. 2)

159 The BN distance between taxa i and j, ΔBN_{i-j} , will be:

160 $\Delta BN_{i-j} = (BN_{tax_i} - BN_{tax_j}) + \frac{\partial BN_{tax_i} - \partial BN_{tax_j}}{\partial Env} = (BN_{tax_i} - BN_{tax_j}) +$
161 $[f(\text{Tax}_i, \text{Symp}_i, \text{Hom}_i) - f(\text{Tax}_j, \text{Symp}_j, \text{Hom}_j)]$ (Eq. 3)

162 The segregation of the BNs of the dominant species of a holm-oak evergreen
163 Mediterranean forest (Figure 4a) and their displacement when exposed to drought in a climatic
164 field experiment (Figure 4b) constitute a consistent confirmation of these two first BN rules.
165 Very similar results for segregation and displacement were found using different calculation
166 approaches, from the additional use of ratios as explanatory variables to the use of discriminant
167 and other multivariate analyses (Appendix S1: Fig. S1). Pairs of species with more overlap in
168 their distributions had greater differences in their BNs as estimated from PCA scores (PCA based
169 on N,P, K, S, Ca and Mg concentrations) in a set of the 12 most representative Mediterranean
170 tree species of South Europe (Figure 4c), confirming the third rule. The BN hypothesis is an
171 extension of the ecological niche concept (Schoener 1986) to the bioelemental components of
172 life. The BN aims to establish the morphological and functional “adaptation” to the abiotic and
173 biotic traits that define the niche of each species-genotype from simple determination of the
174 elemental composition. It is based on the different contribution of each bioelement to the
175 different organismal functions and structures. Its main advantage over other theoretical classical
176 niche concepts is that it can be characterized by the measurable bioelemental composition, the
177 elementome, of the organism, species, community or even biome (Figure 5). This BN

178 measurement is not biased by the most abundant bioelements since the analysis is conducted for
179 a multivariate space defined by the standardized elemental concentrations (Figures 4 and 6).

180 The quantitative BN framework can then be used to improve projections of how species
181 will respond to environmental changes, provided elementome homeostasis/plasticity (Hom) on a
182 given time horizon is known. This Hom can be defined for example as the coefficient of variance
183 of the scores for the different individuals of a species in the PCA. It results from both
184 intraspecific variation and phenotypic individual responses. These two responses can be
185 differentiated when the hypervolume occupied by the individuals of one species is compared
186 with the hypervolume generated after changing the environmental conditions, for instance in
187 experiments in field conditions where environmental conditions are changed.

188 BN aims to refocus the use of the hypervolume in ecological niche studies. A n-
189 dimensional hypervolume was originally proposed by Hutchinson (1957) and defined as a subset
190 of the space, i.e. an n-dimensional geometrical shape, to describe the fundamental niche of a
191 species. Hutchinson suggested that the axes correspond to requirements of the species (e.g. light,
192 temperature, food size) and that the boundaries of the shape indicate the conditions that permit
193 the growth and reproduction of the individuals of a species. This description of the species niche
194 has been widely used and discussed (Begon et al, 2006 Chase and Leibold 2003, Colwell and
195 Rangel 2009, Holt 2009, Peterson et al 2011). There is now a growing set of statistical methods
196 to operate this hypothesis of species niche with data (Blonder 2017). BN should be a tool to
197 detect the niche shifts at short-, medium- and long-term scales in response to all type of
198 environmental changes (drought, species invasion, pests, new biotic relationships, etc.). The
199 mathematical description of each specific BN position in a multi-space in function of the
200 concentrations of several bioelements can be done by using essential multivariate statistical

201 analyses such as principal components analyses (PCA) or functional/general discriminant
202 analysis (F/GDA) (Sardans et al. 2015; Urbina et al. 2017), or other analyses such as kernel
203 density estimation, support vector machine, Bayesian ellipse, etc. (Blonder 2017). BN distances
204 and the level of overlap/separation among different groups can be computed with the score
205 means (centroids) of the main axes (in the PCAs; Figure 4, Appendix S1, Fig. S1) or with
206 squared Mahalanobis distances (in the discriminant analyses; Appendix S1: Fig. S2). These
207 metrics provide numerical evidences of the “distances” between the BN of different species and
208 of the BN displacement (if any) when comparing the BN of different populations of the same
209 species along space or time. Correlations between matrices provide a way of comparing the
210 distances between hypervolumes, for instance the correlation matrix of the genome differences
211 or of the differences in some ecological or ecophysiological variables in a particular set of
212 species with the corresponding correlation matrix of the BN distances of the same set of species.
213 Statistical tests such as Mantel tests (Mantel, 1967) can be used with this aim. All these
214 mathematical tools to estimate the BN allow all levels of comparisons, from species (Sardans et
215 al. 2015) to different taxa and trophic groups (Bartrons et al. 2018), also including the
216 comparison of different populations of the same species growing under distinct environmental
217 conditions (Urbina et al., 2015, 2017).

218 These studies have shown the facility and consistency in the use of different types of
219 multivariate analyses to detect the BN in spatial and temporal gradients/scales. Moreover, the
220 different levels of homeostasis/plasticity of different studied groups can be calculated by the
221 area/volume in that the individuals of each group are spread in the multidimensional space
222 generated. The temporal shifts of the BN of a species or populations of one species can also be
223 calculated and thus their evolution can be followed. BN also allows the description and

224 quantification of niche appearance (i.e. when new species colonize the ecosystem), expansion,
225 contraction, and extinction. Therefore, BN constitutes a quantifiable tool to detect, quantify and
226 thus better comprehend the mechanisms and processes underlying community evolution and
227 species turn-over (Figure 7). Niche construction vs niche expansion or niche contraction vs niche
228 extinction are apparently difficult to differentiate because of their similar outcomes, and niche
229 contraction (and extinction) could be the result of environmental filtering, whereas niche
230 expansion could be due to addition of species to a community. Phenomena within the niche
231 theory such as niche convergence (Pianka et al., 2017) can be followed by monitoring the species
232 BN hyper-dimension along spatial and temporal gradients and focusing on the changes in the
233 level of overlap/distance in their BN.

234

235 **3. Evidence from observational data**

236 3.1- Taxonomy

237 The BN hypothesis is supported by strong evidence in recent studies. Observational data
238 (Sardans and Peñuelas 2014; Bartrons et al., 2018) provide evidence that plant and animal
239 species have their own elementome, their own BN. For example, different groups of species in
240 distinct climatic biomes have different foliar bioelemental compositions and stoichiometries
241 (Sardans et al. 2014, 2015, 2016). Foliar bioelemental composition of forest tree species, for
242 example, has been positively correlated with phylogenetic distance between taxa at different
243 geographic extents, from local to continental (Sardans et al. 2014, 2015, 2016). Bioelemental
244 composition and stoichiometry thus tend to differ more between phylogenetically distant than
245 between phylogenetically close tree species (Sardans and Peñuelas 2014, Sardans et al. 2015,
246 2016). These positive correlations between elemental composition differences and phylogenetic

247 distances have been also observed in animals (Gonzalez et al. 2018) in a further demonstration of
248 the first process underlying the BN hypothesis. The stoichiometry of an individual under a given
249 climatic and edaphic condition is determined in part by the uptake and the efficiency with which
250 the individual uses each bioelement, all of which depend on the trade-offs among the various
251 individual functions that maximize individual fitness under that specific climatic situation. Each
252 species and taxon have its own space in the two-dimensional plot formed by the two first
253 components of the PCA of individual composition (N, P, S, K, Ca and Mg concentrations and
254 their ratios) (See example in Figure 4A and see Appendix S1: Table S1 for the scores of
255 additional PCs).

256

257 3.2- Sympatry

258 Larger differences in the stoichiometry of foliar composition among sympatric than allopatric
259 species have recently been reported both in plant communities (Sardans et al. 2015, 2016) and in
260 animal communities (González et al., 2018) as implied in the third process involved in the BN.
261 The overlapping frequency of all pairwise comparisons among many tree species is correlated
262 with the elementome distances of the corresponding tree pairs (Figure 4c) (Sardans et al. 2015),
263 thus supporting this third process. Different grass and shrub species growing under different
264 climatic conditions in communities with distinct species compositions maintain distinct species-
265 specific foliar compositions, although there may be displacement, expansion or concentration of
266 the biogeochemical space occupied (Figure 6) depending on the coexisting species (Urbina et al.
267 2015, 2017). The species bioelemental composition changes in parallel with the shifts of
268 environmental conditions, in accordance with the new optimal functions that require new
269 proportional use of different bioelements. Contraction, expansion, and displacement at the short-

270 term would thus be complemented by construction (allopatric speciation), extinction,
271 segregation, and equalization at the long-term both for estenoic (narrow range of ecological and
272 environmental conditions) and euroic (wide range) niches (Figure 7).

273 In-depth investigation of these processes constitutes a challenge for ecologists. Some
274 studies have already demonstrated different biogeochemical niche space (euroic-estenoic) among
275 different coexisting species-genotypes in plants (Urbina et al., 2015, 2017, Sardans et al., 2015),
276 animals (González et al., 2017, 2018; Bartrons et al., 2018) and microbes (Pereira and Berry,
277 2017). Other studies have observed how the same species displaced their biogeochemical niche
278 depending on the intraspecific and intraspecific competition (Urbina et al., 2015, 2017; Sardans
279 et al., 2016). However, observational field studies along successional stages, field manipulation
280 studies of biotic (introducing new species, changes in density,...), and abiotic (changing climate
281 conditions or nutrient availability) followed along time with similar studies in greenhouse,
282 micro- and mesocosm conditions would be necessary to further test the consistency of the new
283 holistic view of ecosystem processes provided by the BN.

284

285

286 3.3- Homeostasis/Plasticity

287 Changes in environmental supply (for example soil or water bioelemental composition) can
288 affect the bioelemental composition of organisms, as observed in microbes (Frenken et al 2017)
289 and plants (He et al. 2016, Salvaggiotti et al. 2017). A trade-off between adaptations to be
290 competitive in stable environments versus being successful in less stable environments should
291 underpin the differences in the continuum of homeostasis-plasticity strategies in bioelemental
292 composition and stoichiometry among individuals, species and communities (Figure 3, Figure

293 4B). Higher N and P concentrations and lower N:P ratios coincide with higher stoichiometric
294 plasticity in opportunistic (r-strategy) than in competitor (K-strategy) plant species (Yu et al.
295 2011). Climate can also account for some of the variability in the bioelemental composition of
296 plant tissues at intra- and interspecific levels (Sardans et al. 2015, 2016, Zhang et al. 2012)
297 indirectly though climate controls on bioelemental availability and directly as a forcing of traits
298 linked to the elementome. Environments with fluctuating nutrient availabilities relative to the
299 species' lifetime would have to favor organisms with greater stoichiometric plasticity.
300 Stoichiometric plasticity tends to decrease in response to fertilization as environmental nutrient
301 richness increases, in both terrestrial and aquatic systems (Sistla et al. 2015), thus suggesting a
302 different position in the homeostatic-plasticity continuum between species in nutrient-rich versus
303 nutrient-poor environments. The BN hypothesis allows to detect the plasticity at two levels: at
304 the individual level (phenotypic plasticity) by showing how its individual elemental composition
305 can vary when environmental conditions shift, and at the intraspecific level as a result of
306 intraspecific variability (individual genotypic differences, different ontogenic stage, distinct
307 sex,..) in elemental composition of a set of individuals of the same species living under the same
308 environmental conditions. This dual source of variability due to natural intra-population
309 variability and to individual capacity to shift BN in response to environmental changes can be at
310 least partially detected experimentally with multivariate analyses working with populations of
311 the same genotype growing in different environmental conditions across space or time.

312

313 3.4 Evolutionary change

314 Species have stoichiometric strategies to adapt to heterogeneous environments (Sun et al. 2017)
315 and are able to adapt their elemental compositions to those of the media after several generations

316 (Turner et al. 2017). The stoichiometry of phytoplankton appears to reflect oceanic stoichiometry
317 over geological timescales, providing even stronger support to this link between organisms'
318 composition and media nutrients availability (Quigg et al. 2003; Reinhard et al. 2017). Other
319 studies of terrestrial ecosystems found that microbial community N:P ratios were correlated with
320 plant-community and soil N:P ratios (Kerkhoff and Enquist 2006, Cleveland and Liptzin 2007).
321 Close relationships between microbial and soil C:N ratios across a soil C:N gradient have been
322 also observed, e.g. in Scandinavian boreal forests (Högberg et al. 2006). Evidence suggests that
323 variation in the environmental supply of P affects the expression of highly conserved genes (e.g.
324 those of the phosphate-transporter system) and that the acquisition, assimilation and allocation of
325 P is genetically variable (Jeyasingh and Weider 2007). Transcriptomic studies of algae
326 (Grossman 2000), microbes (Baek and Lee 2007), terrestrial plants (Morcuende et al. 2007) and
327 more recently herbivores (Jeyasingh et al. 2011) have also reported that the availability of P
328 influences genetic expression. Such effects of the supply of P should impinge on fitness and
329 drive evolutionary change. Jeyasingh et al. (2009) observed that genetic variation in
330 phosphoglucose isomerase (Pg) indicated that Pg heterozygotes of *Daphnia pulicaria*
331 outcompeted Pg homozygotes under conditions of low C:P ratios, whereas the opposite was
332 observed under conditions of high C:P ratios.

333 Availability and stoichiometry of bioelements can determine transcriptional expression
334 and metabolomic structure (Rivas-Ubach et al. 2012) and phenotypic selection (Chowdhury et al.
335 2015), and can affect genomes and thereby become evolutionary drivers for organisms and
336 ecosystems on the longest timescales (Acquisti et al. 2009). For example, the mechanisms
337 controlling P acquisition are genetically variable, and the expression of the genes involved is
338 strongly affected by its environmental supply (Grossman 2000, Jeyasingh and Weider 2007,

339 Jeyasingh et al. 2011, 2014). Selection in environments with very low N availability can even
340 favor new genes richer in N-poor nucleotides (Acquist et al. 2009). Evolution could also be
341 controlled by directly affecting the quality of matter, e.g. different N:P ratios available to
342 genomes for the optimal allocation of N and P to DNA/RNA for maximizing fitness (Sardans et
343 al., 2012b).

344

345 3.5 BN of communities

346 Communities should be more capable to adjust their stoichiometry to additional nutrient inputs
347 and varying stoichiometrical conditions than single species (Danger et al. 2008). A simple and
348 comprehensive example, only considering N and P concentrations and stoichiometry, can be
349 observed when P is added to a freshwater community. In such a case, a shift in community
350 composition typically occurs in favor of N-fixing organisms with a posterior cascade of
351 composition changes towards a new equilibrium in community composition (Davidson et al.
352 2007). Environmental changes (anthropogenic or natural) such as perturbations or shifts in
353 climate, N deposition, species invasion or the increase in atmospheric CO₂ can probably generate
354 changes in the availability of various nutrients. Species of the target community can then adapt,
355 depending on their phenotypic flexibility. For example, species with a BN better adapted to new
356 environmental conditions can be favored under large long-term loadings of N by N deposition,
357 but species less well adapted can disappear. Species can also turnover, when possible, with the
358 entrance of new species with traits better adapted to life in the newly generated niches (with
359 different optimal elemental compositions). This sequence of events has been observed in several
360 field studies where shifts in the elemental composition of the media (water or soil) due to
361 changing conditions (anthropogenic or natural) have been translated into changes in the species

362 composition of species of the community (Sterner and Elser, 2002). The species with more
363 favorable elemental compositions gain then importance and/or there is species turnover
364 (Novotny et al. 2007; Yu et al. 2011, Poxleitner et al. 2016, Du 2017). Changes in N:P have ben
365 related to substantial changes in species composition of plant communities (Peñuelas et al 2013)
366 and changes in P loads to adjusted C:P ratios in zooplankton communities (Teurlink et al., 2017).

367

368 **4- Feedbacks on environment**

369 The relationships between evolution and the bioelemental composition of ecosystems are
370 bidirectional, because nutrient supply can affect evolutionary processes, while the changes
371 imposed by evolutionary processes can also affect nutrient supply (Durston and El-Sabaawi
372 2017). As a result, ocean and soil stoichiometries may have been also partly controlled by
373 organisms over long time scales (Tyrrell 1999, Lenton and Klausmeier 2007, Hatton et al. 2015,
374 Zederer et al. 2017).

375 The strongest support of Redfield ratio comes from the fact that the N:P ratio in the open
376 sea areas of all major oceans is remarkably similar to the N:P ratio of plankton (16:1). This is
377 due to the residence time of N and P in the ocean (104 years) relative to the ocean's circulation
378 time (103 years) (Falkowski and Davis 2004). However, in addition to the geochemical causes,
379 biological processes are also underlying Redfield ratios. N₂ fixation can play an important role in
380 ocean N:P ratio homeostasis (Tyrrell 1999, Lenton and Watson 2000, Ganeshram et al. 2002,
381 Kenesi et al. 2009). When ocean N:P ratios fall too low, nitrogen fixation increases by
382 augmenting the biomass of N₂-fixers and when these N₂-fixers decompose or are eaten, their N
383 rich organic matter turns to dissolved ammonium and nitrate in the water increasing water N:P

384 ratios. Conversely, when the ratio rises too high, N₂ fixation inputs decrease by P limitation of
385 N₂-fixation, thus restoring lower water N:P ratio. This process would confer to P primary control
386 of ocean production (Tyrrell 1999, Lenton and Klausmeier 2007).

387 Soil elemental composition is also controlled by the species of plants (Zederer et al.
388 2017), the soil microbiota (Hartman et al. 2017), and the interactions between plants and soil
389 fungal communities. For example, *Quercus* species control nutrient dynamics by determining the
390 composition and activity of the forest floor fungal community (Chávez-Vergara et al. 2016) and
391 the litter type (Hatton et al. 2015). Conifers acidifying soil to reduce soil nutrient availabilities
392 (base cations and P) to outcompete species with a different BN provide another consistent
393 example.

394 We should thus expect a constant feedback between organisms and their environment.
395 Understanding the processes underlying species shifts in bioelemental composition and their
396 genetic basis will help us understand the responses of organisms to environmental changes
397 (Yamamichi et al. 2015, Leal et al. 2017) and in turn the effects of organisms on ecosystem
398 functioning and services (Frisch et al. 2014, Leal et al. 2017). We are currently still moving
399 towards understanding and identifying the feedbacks between species evolution and shifts in the
400 bioelemental composition of ecosystems and how these feedbacks interact and determine
401 ecosystem and evolutionary processes. Available data seems to indicate that the long-term
402 availability and use of bioelements can determine individual selection and therefore ecosystem
403 function and organization and evolution of life.

404

405 **5- The biogeochemical niche hypothesis and related approaches**

406 Functional traits such as leaf mass area, seed size or plant height have been also used to build
407 hypervolumes to identify species-specific niche (Violle and Jiang, 2009; Violle et al., 2014;
408 Lamanna et al., 2014). However, taxonomical groups (e.g. animals, plants, fungi,..) have
409 different functional traits, preventing the use of these approaches to different taxonomy clades.
410 Instead, all the organisms, independently of their taxonomical groups, are constituted by the
411 same basic bio-elements, thus providing a general and easy to measure variable, the elementome,
412 with which to establish the species-specific niche in plants (Sardans et al., 2015), in animals
413 (Gonzalez et al., 2017) and in sets of plants and animals analyzed altogether (Bartrons et al.,
414 2018). The use of organismal elemental stoichiometry has been proven to be related to the
415 species functional traits in aquatic and terrestrial ecosystems (Meunier et al., 2017, Sardans et al.
416 2012c, Peñuelas et al. 2013). This also allows us to establish relationships between observed
417 shifts in ecosystem N:P medium (water or soil) with the shift in species dominance or
418 community and/or in trophic web composition and structure.

419 We first defined the BN in 2008 for plant communities (Peñuelas et al. 2008) and we
420 have since then developed it in several papers (Peñuelas et al. 2008, Sardans and Peñuelas 2014,
421 Frisch et al. 2014, Sardans et al. 2015, 2016, Urbina et al. 2017). Other groups have also used
422 this approach (De and Bhattacharyya 2009, Menge and Hedin 2009, Chimphango et al. 2015,
423 Zamora et al. 2017), obtaining results that have provided solid experimental consistency to BN
424 hypothesis. The BN that we define as species elemental composition is wide in scope and simple
425 in determination, and reflects the structure and function of each species-genotype resulting from
426 its unique evolutionary history (Sardans et al. 2008, 2015, Urbina et al. 2017).

427 BN aimed to progress beyond the classical ecological stoichiometric studies (Elser et al.
428 2000a,b, Sterner and Elser 2002, Elser and Kyle 2009, Elser et al. 2009, 2010, Rivas-Ubach et al.

429 2012, Sardans et al. 2012b). These ecological stoichiometric studies have linked several
430 important ecosystem processes with ecosystem and organismal C:N:P ratios. Several organismal
431 functions and responses to environmental shifts, though, cannot be explained only by the
432 changes in these three main bioelements (Sardans et al. 2012b,c). As stated above, other
433 bioelements (such as K, S, Ca, Mg and Fe) are needed in ecological stoichiometric studies to
434 provide more complete and global information correlating shifts in organismal elemental
435 composition with changes in organismal morphology and function in response to environmental
436 conditions (Peñuelas et al. 2008, Peñuelas and Sardans 2009, Sardans and Peñuelas 2014,
437 Sardans et al. 2015). The BN hypothesis is thus based on the entire elemental composition or at
438 least on the composition of as many elements as possible. The BN hypothesis has afterwards also
439 been used to refer to the concentrations and availabilities of nutrients in soil or water that favor
440 or determine the corresponding spatial occupation by a species or taxon (De and Bhattacharyya
441 2009, Menge and Hedin 2009, Chimphango et al. 2015, Zamora et al. 2017).

442 A similar hypothesis of a “multidimensional stoichiometric niche” was recently
443 introduced by Gonzalez et al. (2017). These authors defined a “multidimensional stoichiometric
444 niche” for taxonomic/trophic groups as their position in the three-dimensional space determined
445 by C, N, and P following Peñuelas et al. (2008). Gonzalez et al (2017) also interestingly opened
446 the possibility to incorporate more elements, and also element fluxes and transformation rates as
447 additional axes. In their studies, Gonzalez et al. (2017, 2018) proved the suitability of this
448 approach for animals. This “multidimensional stoichiometric niche” also tried to incorporate
449 rates of nutrient cycling, but these are much more difficult to quantify than elemental
450 concentrations and ratios. Thus, according with the observations in plants, animals, microbes,
451 and also in combination of different taxa altogether (animals and plants) (Peñuelas et al., 2008;

452 Sardans et al., 2015; González et al., 2017; Bartrons et al., 2008), the BN multidimensional space
453 proposed here should allow the establishment and quantification of the links between the species
454 BN distances and their taxonomic/phylogenetic distances and between homeostatic capacities
455 and sympatries, and thus to provide information on species/taxa evolutionary processes. The BN
456 thus also aims to refocus the use of the hypervolume in ecological and evolutionary niche studies
457 as discussed above in section 2.

458 Another similar approach to BN, ionomics, i.e. the profiles of mineral elements in plants,
459 has been proposed mostly focused on the genetics of plant mineral nutrition. Ionomics analyzes
460 the elemental composition of organisms in relation to soil concentrations of nutrients and trace
461 elements and also plant functional traits (Salt et al., 2008; Buescher et al., 2010; Baxter et al.,
462 2012; Huang and Salt, 2016). Ionomics is mostly used to study the content of bioelements with
463 electrical charge, mostly absorbed from soils, and mostly within a context of the impact on plant
464 health, food quality and/or identification of hyper-accumulation species. It comes from the
465 confluence of metabolomics and plant nutrition (Salt et al., 2008) that, with the incorporation of
466 modern analytical platforms such as ICP techniques, allows to determine most elements
467 altogether. The fusion of this information with bioinformatics and genetic tools, such as genomic
468 sequencing, allows the identification of the genes that control uptake, storage, and use of soil
469 nutrients and trace elements in plants (Baxter et al., 2012; Lowry et al., 2012). This further
470 allows to analyze the effects of genes and environment on plant nutrition (Buescher, et al., 2010;
471 Lowry et al., 2012; Neugebauer et al., 2018) and plant physiological status (Baxter et al., 2008).
472 BN instead considers all, or as many as possible, bioelements, i.e. also including those such as C
473 or N that are initially from atmospheric, not ionic, forms (e.g. N comes mainly from atmospheric
474 N₂; nitrates are rare in bedrock). BN focuses on organisms and on the total elemental

475 composition and stoichiometry as a proxy of their optimal function and morphology, whereas
476 ionomics focuses on the composition of the medium (soil) to describe different ionic niches in
477 the abiotic dimension of an ecosystem. Ionomics has aimed to explain shifts in ionic composition
478 by identifying specific genes and their control in changing environmental conditions, thus
479 providing knowledge and information to improve the nutrition of crop species. BN is instead
480 based on the idea that the entire genome of each genotype/species determines species-specific
481 functionality and morphology (more distinct as genomic differences increase) and thus a distinct
482 use of different bioelements, which in turn contribute asymmetrically to different functions and
483 morphological structures. BN can thus be used without knowing about the control of gene
484 expression or about gene function. Moreover, whereas ionic approach has been developed for
485 plants, BN is developed to be equally applicable to all type of taxa.

486 Even though these ionic studies are mostly agriculturally oriented, and thus the link
487 with the classical theory of ecological niche is not specifically considered, they are consistent
488 with the BN hypothesis, at least at the level to correlate elemental composition with phylogenetic
489 distance. They constitute strong evidence in favor of BN consistency and constitute a very
490 interesting contribution to the BN.

491 **7. Concluding summary and perspectives**

492 Here we propose a BN of genotypes/species/communities that should provide an improved and
493 quantifiable proxy of the general ecological niche. The BN is characterized by a particular
494 hypervolume of their individual elementomes based on their specific needs and use of
495 bioelements in different amounts and proportions for their structure and functioning. Several
496 organismal structures and functions and the responses to environmental shifts cannot be
497 explained only by the changes in the three main bioelements, C, N and P. We propose that other

498 bioelements such as K, S, Ca, Mg, Fe, Mn and Mo, among others, must be incorporated to fully
499 describe the BN. The BN extends the classical ecological niche concept but has the advantage
500 that it can be easily, directly and quantitatively analyzed.

501 The biogeochemical distances, i.e. the differences in elementome, among species, are a function
502 of taxonomy (phylogenetic distance), sympatry (the bioelemental compositions should differ
503 more among coexisting than among non-coexisting species to avoid competitive pressure), and
504 homeostasis (within the continuum between high homeostasis/low plasticity and low
505 homeostasis/high plasticity). Further studies are warranted to discern the ecological and
506 evolutionary processes involved in the BN of all types of individuals, taxa and ecosystems. The
507 changes in bioelemental availability and use at long timescales should determine phenotypic
508 selection and therefore also ecosystem function and organization, and, at the end, the evolution
509 of life and the environment.

510

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516

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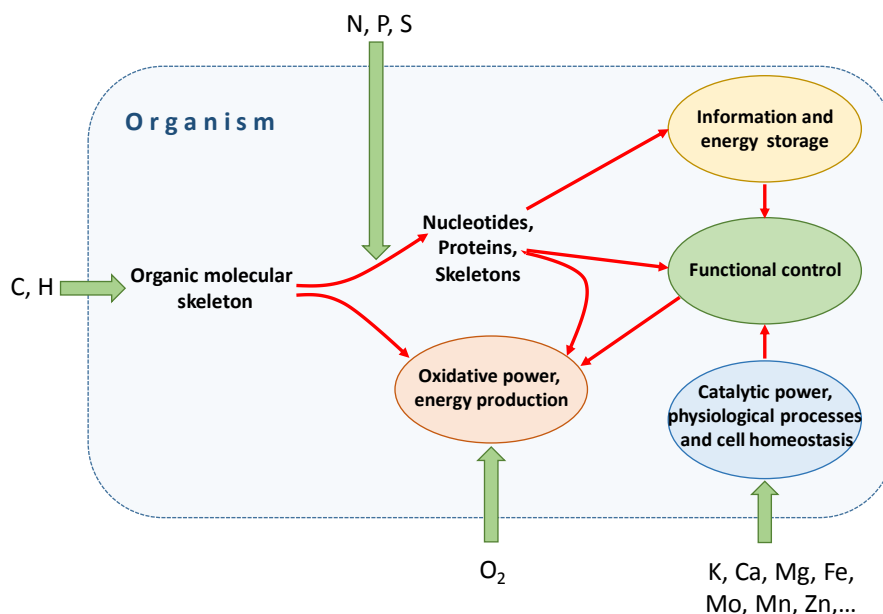
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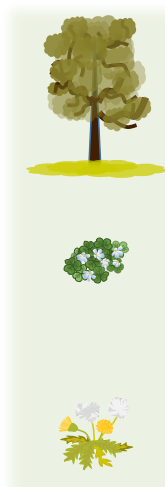
804 **Figure captions**



805

806 Figure 1. Schematic of the use of bioelements by organism in different functions. C, H, and O
 807 are the basis of organismic structure, whereas N, P and S enter in the various biomolecules
 808 allowing biochemical reactions. The molecules that control life processes (e.g. RUBISCO or
 809 chlorophyll), that carry information (e.g. DNA and RNA), and that control the storage and
 810 release of energy when and where necessary (e.g. ATP and NAD) all incorporate N, P and in
 811 lower amounts S. Other bioelements, normally present in low concentrations such as potassium
 812 (K), magnesium (Mg), iron (Fe), calcium (Ca), molybdenum (Mo), manganese (Mn) and zinc
 813 (Zn) are cofactors in catalytic reactions in several biochemical and physiological processes
 814 including for example photosynthesis, respiration, or cellular ion balance. The arrows indicate
 815 the processes in which each bioelement is involved. Green arrows indicate the main organismal
 816 functions to which each bioelement contributes. The red arrows indicate the relationships of the
 817 different bio-molecules with basic organismal structures and functions.

The example of plants



Ecological strategy	Ecological variables					Stoichiometry traits				
	Live-span	Growth rate	Body size	Reproductive strategy	Other	Stoichiometry flexibility	[C]	[N]	[P]	[K]
K-strategy (competitor)	Long	Medium	Large	Few seeds but with highly caloric reserves	Strong investment in structure	Medium	—	↑	↑	—
A-strategy (stress-tolerator)	Medium	Slow	Medium-small	Medium	Strong investment in C-rich secondary compounds	Low	↑	↓	↓	↑
r-strategy (ruderal)	Short	Fast	Small	Many seeds but with low caloric reserves	General trend to develop belowground vegetative & reproductive structures	High	—	↑	↑↑	—

818

819 Figure 2. The example of plants. Hypothesized asymmetric uses of bioelements in evolution,
 820 depending on the ecological strategy of each species. Vertical arrows indicate low or high
 821 concentration values (double arrows indicate especially high values). Dashes indicate not clear
 822 patterns. There is a different use of all these elements depending on the ecological strategy of
 823 each species (Grime 1977).

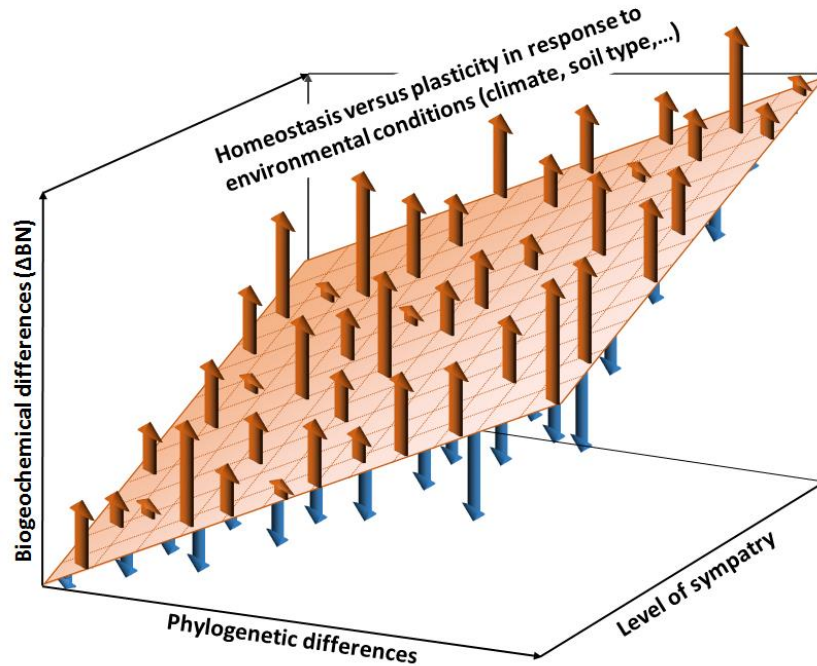
824

$$BN_i = BN \text{ of } \text{taxa}_i = \text{Elementome}_{e_{i1-n}} = (C: N: P: K: S: Mg: Ca: \dots)_{i1-n} = BN_{\text{tax}_i} + \frac{\partial BN_{\text{tax}_i}}{\partial \text{Env}} = BN_{\text{tax}_i} + f(\text{Tax}_i, \text{Symp}_i, \text{Hom}_i)$$

$$BN_j = BN \text{ taxa}_j = \text{Elementome}_{e_{j1-n}} = (C: N: P: K: S: Mg: Ca \dots)_j$$

$$BN_j = BN_{\text{tax}_j} + \frac{\partial BN_{\text{tax}_j}}{\partial \text{Env}} = BN_{\text{tax}_{j1-n}} + f(\text{Tax}_j, \text{Symp}_j, \text{Hom}_j)$$

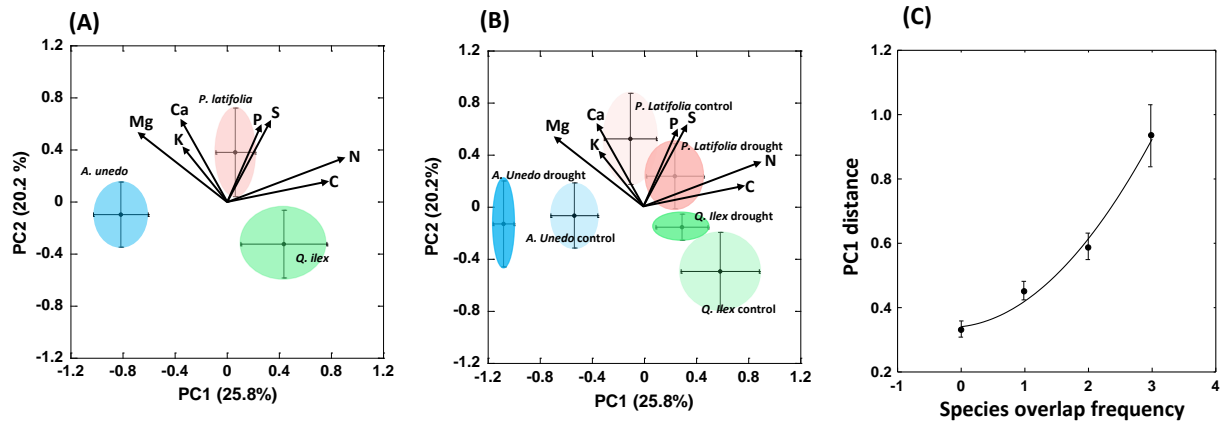
$$\Delta BN_{i-j} = (BN_{\text{tax}_i} - BN_{\text{tax}_j}) + \frac{\partial BN_{\text{tax}_i} - \partial BN_{\text{tax}_j}}{\partial \text{Env}} = (BN_{\text{tax}_i} - BN_{\text{tax}_j}) + [f(\text{Tax}_i, \text{Symp}_i, \text{Hom}_i) - f(\text{Tax}_j, \text{Symp}_j, \text{Hom}_j)]$$



825

826 Figure 3. Theoretical distribution of the biogeochemical differences between species as a
 827 function of (1) the taxonomic and phylogenetic distances between them, (2) the level of
 828 sympatry (overlap frequency in their corresponding areas of distribution) and of (3) the
 829 homeostatic or flexible response to current environmental conditions (e.g. climate, soil traits).
 830 The surface represents the biogeochemical distance resulting from phylogenetic distance and
 831 level of sympatry, and the brown and blue arrows the positive and negative residuals of the BN
 832 distance induced by variable current environmental conditions (e.g. climate, soil traits,
 833 neighbors).

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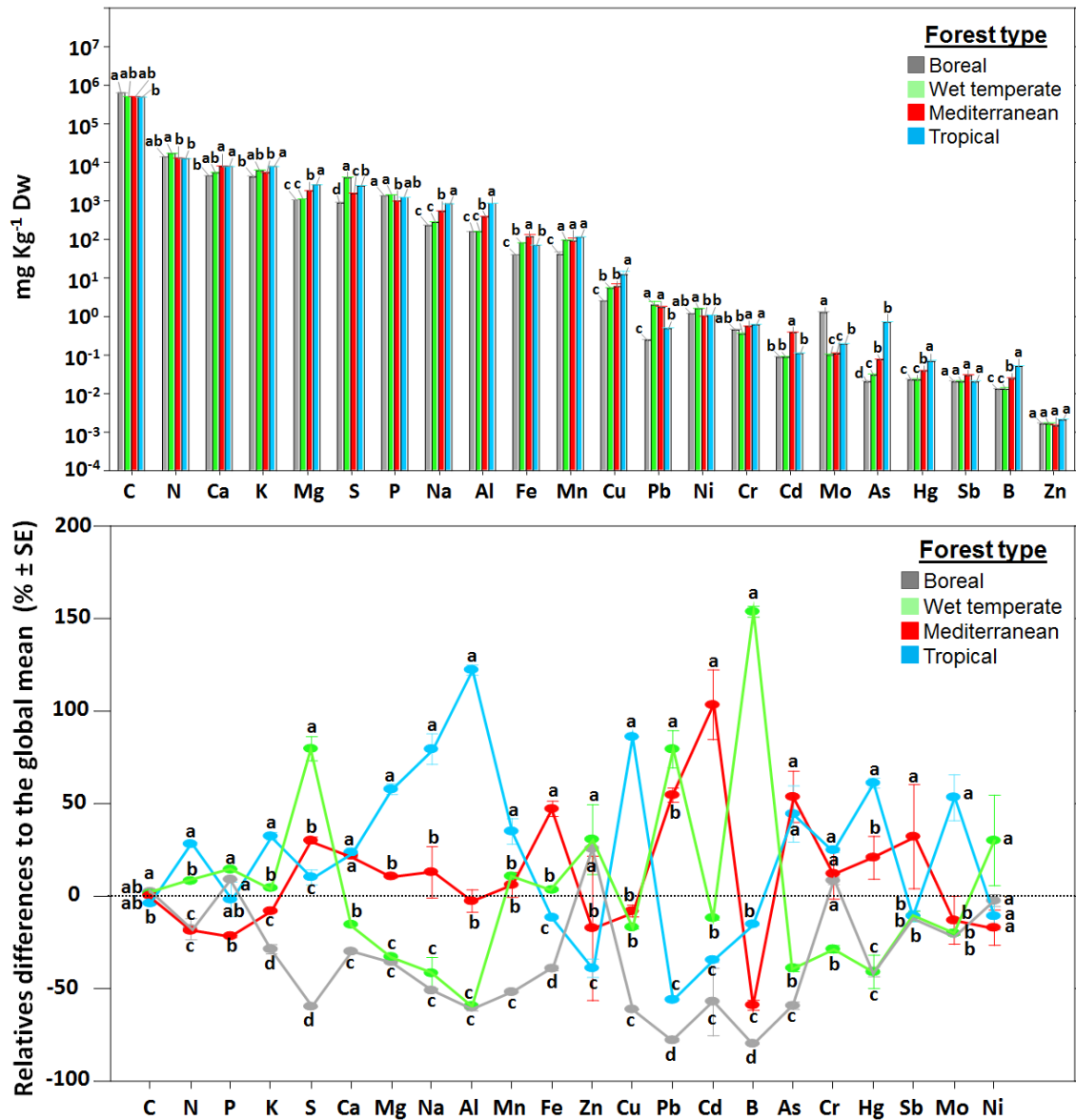


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836 Figure 4. (A) Biogeochemical niche segregation among coexisting plant species. We analyzed
 837 the foliar elementomes of the three dominant trees (*Quercus ilex*, *Arbutus unedo* and *Phillyrea*
 838 *latifolia*) in a field experiment of climatic manipulation in an evergreen Mediterranean forest in
 839 the Prades Mountains (Catalonia). We plotted the plant scores for the first two principal
 840 components of the principal component analysis (PCA) conducted with foliar N, P, K, S, Ca, Mg
 841 concentrations as variables. (B) Shifts in biogeochemical niche in a drought experiment. The
 842 mean \pm confidence intervals (95%, represented by the ellipses) of the species \times climate
 843 interaction groups are depicted in the plot thus comparing control plants to those submitted to a
 844 decrease of 20 % of water availability during six years in field conditions. The other PCAs (from
 845 PC3 to PC7) and the corresponding other axes are now shown in the Appendix S1: Table S1.
 846 (C) Score distances for PC1 of the PCA of the foliar stoichiometry (foliar N, P, K, S, Ca and Mg
 847 concentrations and their ratios) in a set of the 12 most representative Mediterranean species of
 848 South Europe as a function of the frequency of pairwise species overlap (coincidence in the same
 849 community): six broadleaf species, *Quercus ilex*, *Q. faginea*, *Q. pyrenaica*, *Q. suber*, *Castanea*
 850 *sativa* and *Eucalyptus globulus*, and six needleleaf species, *Pinus halepensis*, *P. pinaster*, *P.*
 851 *nigra*, *P. pinea*, *Juniperus oxycedrus* and *J. thurifera*). 0, no overlap; 1, occasional overlap; 2,

852 moderate overlap; 3, frequent overlap. Based on Rivas-Martinez et al. (2001) De Bolos and Vigo
 853 (2001), CREAM, Catalonia cover map, (2013) and Folk et al. (1984).

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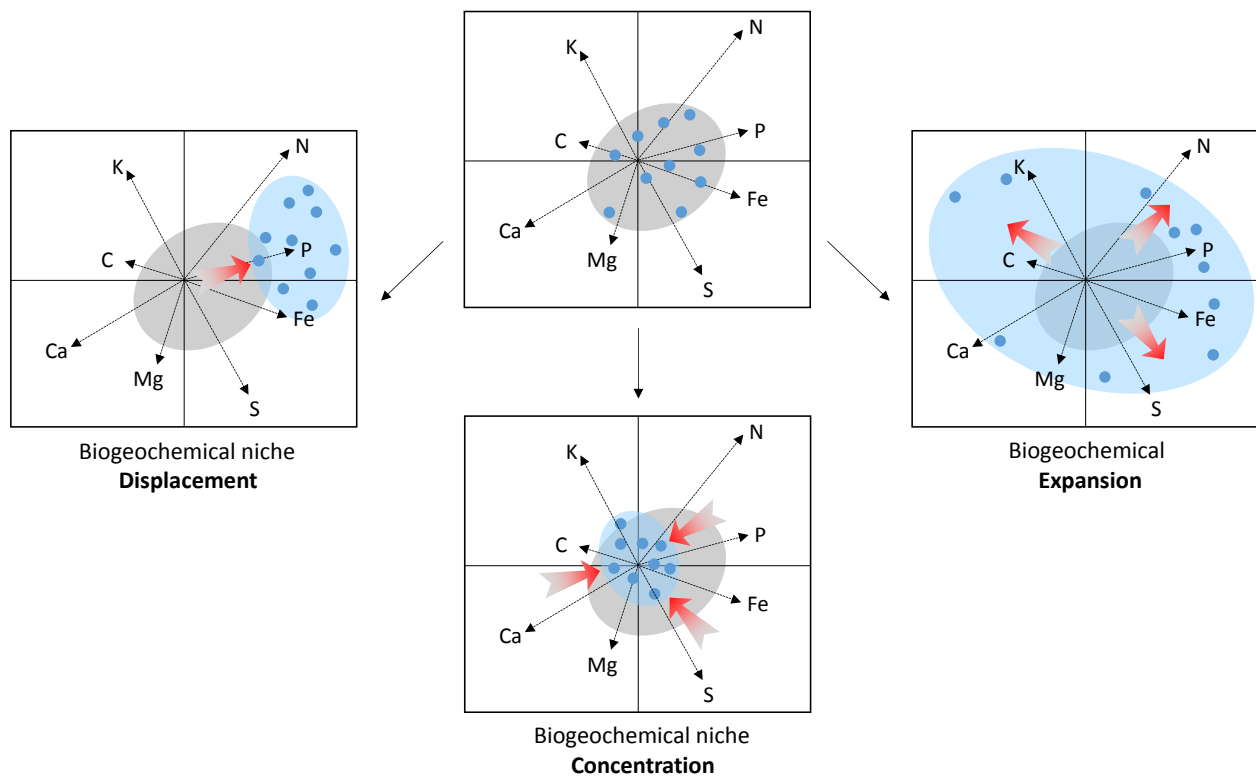


855

856 Figure 5. Elementome, full spectrum of bioelemental content, of boreal, wet temperate,
 857 Mediterranean and tropical forests. (A) Elementomes calculated from web of science data
 858 incorporated to our GEU-CREAM dataset (Boreal forest: 473 sites and 1890 trees; Wet temperate

859 forests: 2882 sites and 12665 trees; Mediterranean forests: 2001 sites and 2901 trees; Tropical
 860 wet forest: 1100 sites and 1821 trees). The values in plot B show the difference of each forest
 861 type relative to the global mean ($\% \pm SE$). We have used data from the main available data bases
 862 (e.g. TRY and ICP forest) and from an exhaustive search for woody plants from non-
 863 manipulated areas in WEB of Science

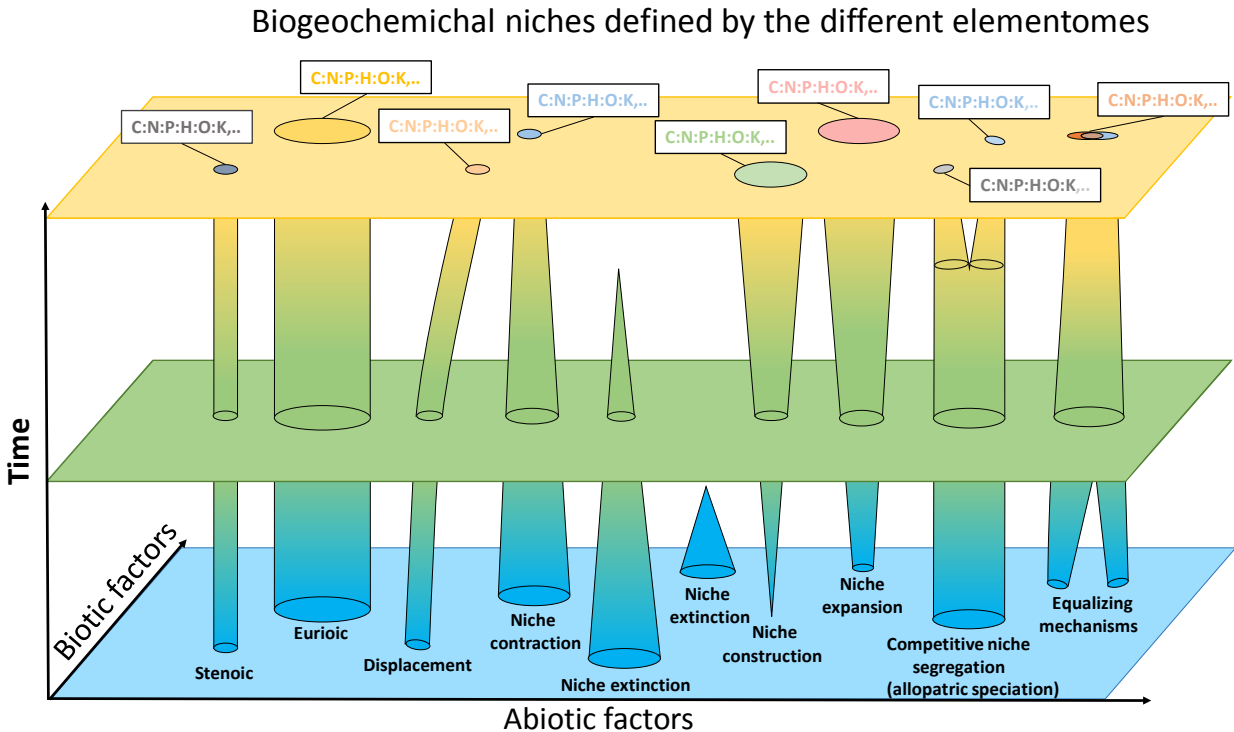
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865

866 Figure 6. Three possible biogeochemical niche responses to short-term changes in the
 867 environmental conditions: Contraction, Expansion or Displacement. Blue dots represent
 868 individuals of a species, grey and blue ellipses represent the initial and final BN space.

869



870

871 Figure 7. Possible responses of species biogeochemical niches to long-term changes in the
 872 abiotic and biotic environmental conditions (possible evolutionary changes in the elementome of
 873 a species). We hypothesize that each species has an optimal function related with its niche traits
 874 and thus an optimal content of bioelements. Species have an optimal function related with its
 875 niche traits and thus an optimal content of bioelements. The species content of bioelements
 876 changes in parallel with the species niche shifts according with the new optimal functions for the
 877 new environmental conditions. Contraction, expansion, and displacement are here complemented
 878 by construction (allopatric speciation), extinction, segregation, and equalization at the long-term
 879 both for estenoic (narrow range of ecological and environmental conditions) and euroic (wide
 880 range) niches. The communities resulting of interactions with changing abiotic and biotic
 881 conditions are even more capable to adjust their stoichiometry than single species to new nutrient
 882 supply and stoichiometrical conditions. Each particular niche constitutes a singular functional

883 and physical position in the ecosystem that can vary with time in response to changes in abiotic
884 variables (climate, perturbations such as volcanic eruptions, etc.) and to changes in the impacts
885 and feed-backs of the other organisms living in it (biotic factors). In this context, each niche
886 should determine a specific and particular content of different bioelements in each moment. This
887 should also determine a singular elementome accompanying the niche characteristics of each
888 genotype and species at both micro- and macro-evolutionary timescales.