

1 **General destabilizing effects of eutrophication on grassland productivity at multiple**  
2 **spatial scales**

3 Yann Hautier<sup>1\*</sup>, Pengfei Zhang<sup>1-4</sup>, Michel Loreau<sup>5</sup>, Kevin R. Wilcox<sup>6</sup>, Eric W. Seabloom<sup>7</sup>,  
4 Elizabeth T. Borer<sup>7</sup>, Jarrett E.K. Byrnes<sup>8</sup>, Sally E. Koerner<sup>9</sup>, Kimberly J. Komatsu<sup>10</sup>, Jonathan  
5 S. Lefcheck<sup>11</sup>, Andy Hector<sup>12</sup>, Peter B. Adler<sup>13</sup>, Juan Alberti<sup>14</sup>, Carlos A. Arnillas<sup>15</sup>, Jonathan  
6 D. Bakker<sup>16</sup>, Lars A. Brudvig<sup>17</sup>, Miguel N. Bugalho<sup>18</sup>, Marc Cadotte<sup>19</sup>, Maria C. Caldeira<sup>20</sup>,  
7 Oliver Carroll<sup>21</sup>, Mick Crawley<sup>22</sup>, Scott L. Collins<sup>23</sup>, Pedro Daleo<sup>14</sup>, Laura E. Dee<sup>24</sup>, Nico  
8 Eisenhauer<sup>25,26</sup>, Anu Eskelinen<sup>25,27,28</sup>, Philip A. Fay<sup>29</sup>, Benjamin Gilbert<sup>30</sup>, Amandine  
9 Hansar<sup>31</sup>, Forest Isbell<sup>7</sup>, Johannes M. H. Knops<sup>32</sup>, Andrew S. MacDougall<sup>21</sup>, Rebecca L.  
10 McCulley<sup>33</sup>, Joslin L. Moore<sup>34</sup>, John W. Morgan<sup>35</sup>, Akira S. Mori<sup>36</sup>, Pablo L. Peri<sup>37</sup>, Edwin T.  
11 Pos<sup>1</sup>, Sally A. Power<sup>38</sup>, Jodi N. Price<sup>39</sup>, Peter B. Reich<sup>38,40</sup>, Anita C. Risch<sup>41</sup>, Christiane  
12 Roscher<sup>25,42</sup>, Mahesh Sankaran<sup>43</sup>, Martin Schütz<sup>41</sup>, Melinda Smith<sup>44,45</sup>, Carly Stevens<sup>46</sup>, Pedro  
13 M. Tognetti<sup>47</sup>, Risto Virtanen<sup>28</sup>, Glenda M. Wardle<sup>48</sup>, Peter A. Wilfahrt<sup>7</sup>, and Shaopeng  
14 Wang<sup>49\*</sup>

15  
16 \*Corresponding authors: y.hautier@uu.nl, shaopeng.wang@pku.edu.cn

17  
18 <sup>1</sup>Ecology and Biodiversity Group, Department of Biology, Utrecht University, Padualaan 8,  
19 3584 CH Utrecht, The Netherlands. <sup>2</sup>State Key Laboratory of Grassland and Agro-  
20 Ecosystems, School of Life Science, Lanzhou University, Gansu Province, 730000, P. R.  
21 China. <sup>3</sup>Institute of Eco-Environmental Forensics of Shandong University, Shandong  
22 Province, 266237, P. R. China. <sup>4</sup>Ministry of Justice Hub for Research & Practice in Eco-  
23 Environmental Forensics, Shandong Province, 266237, P. R. China. <sup>5</sup>Centre for Biodiversity  
24 Theory and Modelling, Theoretical and Experimental Ecology Station, CNRS, 2 route du  
25 CNRS, 09200 Moulis, France. <sup>6</sup>Department of Ecosystem Science and Management,  
26 University of Wyoming, Laramie, WY, USA. <sup>7</sup>Department of Ecology, Evolution, and  
27 Behavior, University of MN, St. Paul, MN, 55108 USA. <sup>8</sup>Department of Biology, University  
28 of Massachusetts Boston, Boston, MA 02125. <sup>9</sup>Department of Biology, University of North  
29 Carolina Greensboro, Greensboro, NC, USA. <sup>10</sup>Smithsonian Environmental Research Center,  
30 Edgewater, MD 21037 USA. <sup>11</sup>Tennenbaum Marine Observatories Network, MarineGEO,  
31 Smithsonian Environmental Research Center, Edgewater, MD 21037, USA. <sup>12</sup>University of  
32 Oxford Department of Plant Sciences, OX1 3RB, UK. <sup>13</sup>Department of Wildland Resources  
33 and the Ecology Center, Utah State University, Logan, UT, 84322, USA. <sup>14</sup>Instituto de  
34 Investigaciones Marinas y Costeras (IIMyC), FCEyN, UNMdP-CONICET, CC 1260 Correo

35 Central, B7600WAG, Mar del Plata, Argentina. <sup>15</sup>Department of Physical and Environmental  
36 Sciences, University of Toronto at Scarborough, Scarborough, Canada. <sup>16</sup>School of  
37 Environmental and Forest Sciences, University of Washington, Seattle, WA 98195-4115  
38 USA. <sup>17</sup>Department of Plant Biology and Program in Ecology, Evolutionary Biology, and  
39 Behavior, Michigan State University, East Lansing, MI, USA. <sup>18</sup>Centre for Applied Ecology  
40 "Prof. Baeta Neves" (CEABN-InBIO), School of Agriculture, University of Lisbon, Portugal.  
41 <sup>19</sup>Department of Biological Sciences, University of Toronto at Scarborough, Scarborough,  
42 Canada. <sup>20</sup>Forest Research Centre, School of Agriculture, University of Lisbon, Portugal.  
43 <sup>21</sup>Department of Integrative Biology, University of Guelph, Guelph, Ontario, Canada  
44 N1G2W1. <sup>22</sup>Life Sciences, Imperial College London, Silwood Park, Ascot, SL5 7PY, UK.  
45 <sup>23</sup>University of New Mexico, Department of Biology, Albuquerque, NM 87131 USA.  
46 <sup>24</sup>Department of Ecology and Evolutionary Biology, University of Colorado at Boulder, 1560  
47 30th Street, Boulder, Colorado 80309-0450 USA. <sup>25</sup>German Centre for Integrative  
48 Biodiversity Research (iDiv) Halle-Jena-Leipzig, Deutscher Platz 5e, 04103 Leipzig,  
49 Germany. <sup>26</sup>Leipzig University, Institute of Biology, Deutscher Platz 5e, 04103 Leipzig,  
50 Germany. <sup>27</sup>Department of Physiological Diversity, Helmholtz Centre for Environmental  
51 Research - UFZ, Leipzig, Germany. <sup>28</sup>Department of Ecology and Genetics, University of  
52 Oulu, Oulu, Finland. <sup>29</sup>USDA-ARS Grassland, Soil, and Water Research Laboratory,  
53 Temple, Texas 76502 USA. <sup>30</sup>Department of Ecology and Evolutionary Biology, University  
54 of Toronto, Canada M5S3B2. <sup>31</sup>Centre de recherche en écologie expérimentale et prédictive  
55 (CEREEP-Ecotron IleDeFrance), Département de biologie, Ecole normale supérieure, CNRS,  
56 PSL University, 77140, St-Pierre-les-Nemours, France. <sup>32</sup>Department of Health and  
57 Environmental Sciences, Xi'an Jiaotong Liverpool University, Suzhou, Jiangsu, 214123,  
58 China. <sup>33</sup>University of Kentucky, Plant & Soil Science, 1405 Veterans Drive, Lexington KY  
59 40546-0312. <sup>34</sup>School of Biological Sciences, Monash University, Clayton Campus, Victoria  
60 3800, Australia. <sup>35</sup>Department of Ecology, Environment & Evolution, La Trobe University,  
61 Bundoora 3086, Australia. <sup>36</sup>Graduate School of Environment and Information Sciences,  
62 Yokohama National University, 79-7 Tokiwadai, Hodogaya, Yokohama, Kanagawa, 240-  
63 8501, Japan. <sup>37</sup>INTA (National Institute of Agricultural Research)- UNPA (Southern  
64 Patagonia National University)-CONICET. Santa Cruz, Argentina. <sup>38</sup>Hawkesbury Institute  
65 for the Environment, Western Sydney University, Locked Bag 1797, Penrith, New South  
66 Wales 2751, Australia. <sup>39</sup>Institute of Land, Water and Society, Charles Sturt University,  
67 Albury, NSW, Australia, 2640. <sup>40</sup>Department of Forest Resources, University of Minnesota,  
68 Saint Paul, MN, USA. <sup>41</sup>Swiss Federal Institute for Forest, Snow and Landscape Research

69 WSL, Zuercherstrasse 111, 8903 Birmensdorf, Switzerland. <sup>42</sup>UFZ, Helmholtz Centre for  
70 Environmental Research, Physiological Diversity, Permoserstrasse 15, 04318 Leipzig,  
71 Germany. <sup>43</sup>Ecology & Evolution Group, National Centre for Biological Sciences, TIFR,  
72 Bangalore, Karnataka 560065, India (and School of Biology, University of Leeds, Leeds  
73 LS2 9JT, UK. <sup>44</sup>Department of Biology, Colorado State University, Fort Collins, CO, 80523.  
74 <sup>45</sup>Graduate Degree Program in Ecology, Colorado State University, Fort Collins, CO, 80523.  
75 <sup>46</sup>Lancaster Environment Centre, Lancaster University, Lancaster, LA1 4YQ. <sup>47</sup>IFEVA-  
76 Facultad de Agronomia, Universidad de Buenos Aires - CONICET, Av San Martin 4453  
77 C1417DSE, Ciudad Autonoma de Buenos Aires, Argentina. <sup>48</sup>School of Life and  
78 Environmental Sciences, University of Sydney, Sydney, New South Wales 2006, Australia.  
79 <sup>49</sup>Institute of Ecology, College of Urban and Environmental Science, and Key Laboratory for  
80 Earth Surface Processes of the Ministry of Education, Peking University, 100871, Beijing,  
81 China.

82

83 **Eutrophication is a widespread environmental change that usually reduces the stabilizing**  
84 **effect of plant diversity on productivity in local communities. Whether this effect is scale**  
85 **dependent remains to be elucidated. Here, we determined the relationship between plant**  
86 **diversity and temporal stability of productivity for 243 plant communities from 42**  
87 **grasslands across the globe and quantified the effect of chronic fertilization on these**  
88 **relationships. Unfertilized local communities with more plant species exhibited greater**  
89 **asynchronous dynamics among species in response to natural environmental fluctuations,**  
90 **resulting in greater local stability (alpha stability). Moreover, neighbourhood**  
91 **communities that had greater spatial variation in plant species composition within sites**  
92 **(higher beta diversity) had greater spatial asynchrony of productivity among**  
93 **communities, resulting in greater stability at the larger scale (gamma stability).**  
94 **Importantly, fertilization consistently weakened the contribution of plant diversity to**  
95 **both of these stabilizing mechanisms, thus diminishing the positive effect of biodiversity**  
96 **on stability at differing spatial scales. Our findings suggest that preserving grassland**  
97 **functional stability requires conservation of plant diversity within and among ecological**  
98 **communities. Furthermore, our results demonstrate the threat of increased ecosystem**  
99 **eutrophication to the stable provisioning of grassland services across spatial scales under**  
100 **current and future environmental conditions.**

101

102 Humans are altering global nutrient cycles via combustion of fossil fuels and fertilizer  
103 application<sup>1</sup>. We have more than doubled pre-industrial rates of nitrogen (N) and phosphorus  
104 (P) supply to terrestrial ecosystems<sup>2</sup>. Terrestrial N and P inputs are predicted to reach levels  
105 that are three to four times pre-industrial rates by 2050<sup>3</sup>. This pervasive global eutrophication  
106 will have dramatic consequences on the structure and functioning of terrestrial and aquatic  
107 ecosystems<sup>3</sup>. In grasslands, nutrient enrichment usually increases primary productivity, but  
108 reduces plant diversity and alters the ability of ecosystems to reliably provide functions and  
109 services for humanity<sup>4-7</sup>.

110

111 Concerns that eutrophication compromises both the diversity and stability of ecosystems have  
112 led to a growing number of theoretical and empirical studies investigating how these ecosystem  
113 responses may be mechanistically linked<sup>4,6,8-11</sup>. These studies, have repeatedly shown that the  
114 positive effect of plant species richness on the temporal stability of community productivity in  
115 ambient (unfertilized) conditions is usually reduced with fertilization<sup>4-6</sup>. However, these studies  
116 have primarily focused on plant responses at relatively small scales (i.e., within single local  
117 communities). Whether fertilization reduces the positive effect of diversity on temporal  
118 stability at larger scales (i.e. among neighbouring local communities) remains unclear. Filling  
119 this knowledge gap is important because the stable provision of ecosystem services is critical  
120 for society<sup>12</sup>. This is especially true, given an increasing concern for large variability of  
121 environmental conditions due to multiple anthropogenic influences including eutrophication  
122 and climate change<sup>13</sup>.

123

124 A recent theoretical framework allows the quantification of the processes that determine the  
125 stability of ecosystem functioning at scales beyond the single local community (Fig. 1)<sup>14-16</sup>.  
126 Stability at any given scale is defined as the temporal mean of primary productivity divided by  
127 its standard deviation<sup>17</sup>. Higher local scale community stability (*alpha stability*), can result  
128 from two main processes. First, a higher average temporal stability of all species in the  
129 community (*species stability*) can stabilize community productivity due to lower variation in  
130 individual species abundances from year to year (Fig. 1b). Second, more asynchronous  
131 temporal dynamics among species in response to environmental fluctuations (*species*  
132 *asynchrony*) can stabilize community productivity because declines in the abundance of some  
133 species through time are compensated for by increases in other species (Fig. 1c). Higher  
134 stability at the larger scale (*gamma stability*) can result from higher alpha stability and more  
135 asynchronous dynamics across local communities (*spatial asynchrony*) (Fig. 1d). Thus, the

136 stabilizing effect of spatial asynchrony on productivity at the larger scale (spatial insurance  
137 hypothesis)<sup>14,18</sup> mirrors the stabilizing effect of species asynchrony on productivity at the local  
138 scale (species or local insurance hypothesis)<sup>8,16,19,20</sup>. Higher species asynchrony and species  
139 stability can result from higher local species diversity through higher *species richness*<sup>9,21,22</sup>,  
140 higher *species evenness*<sup>8</sup>, or both (e.g. higher values of diversity indices - such as the *Shannon*  
141 index - that combines the two<sup>23</sup>) (Fig. 1e). Higher spatial asynchrony can result from greater  
142 local species diversity or higher variation in species composition among communities (*beta*  
143 *diversity*)<sup>16</sup>.

144

145 According to this framework, fertilization can affect the links between diversity, asynchrony  
146 and stability across spatial scales (Fig. 1e, Table 1). At the local scale, fertilization can decrease  
147 **niche dimensionality**, and favour a few dominant plant species by affecting the competitive  
148 balance among species, potentially reducing the insurance effects of local diversity<sup>7,22</sup>. At the  
149 larger scale, fertilization can reduce spatial heterogeneity in community composition, and  
150 decrease variations among local plant community structure, potentially reducing the spatial  
151 insurance effect of beta diversity<sup>16</sup>. Moreover, fertilization often reduces plant diversity which  
152 could in turn reduce asynchrony and stability at multiple scales<sup>4,9,17,24</sup>. However, the role of  
153 fertilization in mediating the functional consequences of biodiversity changes (variations in the  
154 number, abundance and identities of species) and compensatory mechanisms (variation and  
155 compensation in species responses) that can affect the stable provisioning of ecosystem  
156 functions at larger spatial scales remains to be elucidated<sup>25</sup>.

157

158 To our knowledge, only one recent study has assessed the effect of nutrient enrichment on  
159 stability within and among interconnected communities in a temperate grassland<sup>26</sup>. By adding  
160 different nitrogen treatments to communities in ten blocks spread out within a single site, that  
161 study found that five years of chronic nitrogen addition reduced alpha stability through a  
162 decline in species asynchrony, but had no effect on spatial asynchrony. However, these  
163 conclusions were based on a single grassland site manipulating a single nutrient, with the  
164 implicit assumption that the relationship between diversity and stability was unaffected by  
165 eutrophication. This argues for multisite comparative studies assessing the generality of the  
166 mechanistic links between these ecosystem responses to eutrophication.

167

168 Here, we use a coordinated, multi-site and multi-year nutrient enrichment experiment (+/-  
169 chronic nitrogen, phosphorus and potassium addition, Nutrient Network<sup>27</sup>) to assess the scale

170 dependence of fertilization impacts on plant diversity and stability. Treatments were randomly  
171 assigned to 25 m<sup>2</sup> plots and were replicated in three blocks at most sites (Extended Data Table  
172 1). Samples were collected in 1 m<sup>2</sup> subplots across 243 communities from 42 grassland sites  
173 on six continents and followed a standardized protocol at all sites<sup>27</sup>. We selected these sites as  
174 they contained between four to nine years of experimental duration (hereafter ‘period of  
175 experimental duration’) and three blocks per site, excluding additional blocks from sites that  
176 had more than three (Extended Data Table 1). Sites spanned a broad range of seasonal variation  
177 in precipitation and temperature (Extended Data Fig.1) and a wide range of grassland types  
178 (Extended Data Table 1). In our analysis, we treated each 1 m<sup>2</sup> subplot as a ‘community’ and  
179 the replicated subplots within a site as the ‘larger scale’ *sensu* Whittaker<sup>28</sup>. We computed  
180 diversity, asynchrony, and stability within a community (local ‘alpha’ scale) and across the  
181 three replicated communities within a site (larger ‘gamma’ scale) (see Methods). We then used  
182 bivariate analysis and structural equation modelling (SEM)<sup>29</sup> to assess fertilizer impacts and  
183 disentangle the relative contributions of diversity and asynchrony to stability (Fig. 1e).

184

## 185 **Results and Discussion**

186 **Analyses of variance** revealed the negative effects of nutrient inputs on **biodiversity and**  
187 **stability** at the two scales investigated, consistent with recent findings from a single site<sup>26</sup>.  
188 **Fertilization consistently reduced species richness, alpha and gamma stability but had no effect**  
189 **on beta diversity (Extended Data Fig.2)**. Bivariate analyses **further** revealed the negative effects  
190 of nutrient inputs on biodiversity-stability relationships at the two scales investigated (Fig. 2).  
191 Relationships were generally consistent across the different periods of experimental duration  
192 considered (Extended Data Table 2). Under ambient (unfertilized) conditions, species richness  
193 was positively associated with alpha and gamma stability (Fig. 2a, b), but fertilization  
194 weakened the positive effect of species richness on stability at the two scales (Fig. 2c, d).  
195 Fertilization reduced local stability of grassland functioning by increasing temporal variability  
196 in species-rich communities (**Extended Data Fig.3**). Similarly, high beta diversity (variation in  
197 species composition among communities) was positively associated with spatial asynchrony  
198 and gamma stability under ambient conditions (Fig. 2e, f), but again fertilization weakened the  
199 positive effect of beta diversity on spatial asynchrony and gamma stability (Fig. 2g, h). These  
200 results remained when accounting for variation in climate using residual regression (**Extended**  
201 **Data Fig.4**), when using local diversity indices accounting for species abundance (**Extended**  
202 **Data Fig.5**), and when data were divided into overlapping intervals of four years (**Extended**  
203 **Data Fig.6**). Our results extend previous evidence of the negative impact of fertilization on the

204 diversity-stability relationship obtained within local plots and over shorter experimental  
205 periods<sup>4,6,26</sup>. Importantly, they show that these negative effects propagate from within to among  
206 communities. To our knowledge, our study is the first to report the negative impacts of  
207 fertilization on the relationships of beta diversity with spatial asynchrony and gamma stability.

208

209 To understand the relative role of local vs. larger scale community properties in determining  
210 asynchrony and stability at different spatial scales, we conducted SEM analyses including all  
211 measures in a single causal model (Fig. 3, [Extended Data Fig.7](#), Extended Data Table 3). Under  
212 ambient conditions, SEM revealed that higher plant species richness contributed to greater  
213 alpha and gamma stability largely through higher asynchronous dynamics among species  
214 (species asynchrony, standardized path coefficient = **0.39**), and not necessarily through greater  
215 species stability (standardized path coefficient = **0.01**) (Fig. 3a, [Extended Data Fig.8a, b](#)). The  
216 positive association between species richness and alpha stability is consistent with existing  
217 experimental<sup>17,24</sup> and shorter-term observational evidence<sup>4,30,31</sup>. Our results confirm that the  
218 stabilizing effects of species richness in naturally-assembled grassland communities is largely  
219 driven by species asynchrony, but not species stability<sup>4,6,22,26</sup>. In addition, they show that the  
220 positive impact of species richness on the stability of community productivity via species  
221 asynchrony in turn leads to greater stability of productivity at the larger spatial scale.

222

223 While correlated with species richness, higher beta diversity also contributed to greater gamma  
224 stability through an independent pathway, namely via higher asynchronous dynamics among  
225 local communities (spatial asynchrony, standardized path coefficient = **0.20**, Fig. 3a). While  
226 theoretical studies have suggested a role for beta diversity in driving spatial asynchrony<sup>15,16</sup>,  
227 previous empirical studies conducted along a nitrogen gradient at a single site<sup>26</sup> or across 62  
228 sites with non-standardized protocols<sup>21</sup> did not find an association between these two variables.  
229 Here, we show that the presence of different species among local communities is linked to  
230 higher variation in dynamics among them, demonstrating the stabilizing role of beta diversity  
231 at larger spatial scales through spatial asynchrony. This also indicates the need for multi-site  
232 replication with standardized treatments and protocols to detect such effects.

233

234 Importantly, fertilization acted to destabilize productivity at the local and larger spatial scale  
235 through several mechanisms ([Fig. 3, Table 2](#)). At the local scale, fertilization weakened the  
236 positive effects of plant species richness on alpha and gamma stability (Fig. 2a, c and b, d) via  
237 a combination of two processes (Fig. 3b, [Extended Data Fig.8c, d](#)). First, the positive

238 relationship between species richness and species asynchrony in the control communities  
239 (standardized path coefficient = 0.39, Fig. 3a), was weaker in the fertilized communities  
240 (standardized path coefficient = 0.20, Fig. 3b). Moreover, this general positive effect of  
241 richness on asynchrony was counteracted by a second stronger negative relationship of richness  
242 with species stability (standardized path coefficient = -0.37). Such negative effect of  
243 fertilization on species stability was not observed under ambient conditions and could be due  
244 to shifts in functional composition in species-rich communities from more stable conservative  
245 species to less stable exploitative species in a temporally variable environment<sup>32,33</sup>. Together,  
246 these two effects explain the overall weaker alpha stability at higher richness with fertilization.  
247 We did not find evidence that the loss of diversity caused by fertilization (an average of  $-1.8 \pm$   
248  $0.5$  species  $m^{-2}$ , Extended Data Fig.2a, Extended Data Fig.9a) was related to the decline of  
249 alpha stability, confirming results from other studies<sup>5,6</sup> and earlier Nutrient Network results<sup>4</sup>  
250 obtained over shorter time periods. This could be because the negative feedback of the loss of  
251 richness caused by fertilization on stability requires a longer experimental duration, or greater  
252 loss of plant diversity, to manifest<sup>9,34</sup>. Another possible explanation is that fertilization may  
253 have a direct positive effect on stability, by increasing community biomass ( $t = 2.41$ , d.f. = 326,  
254  $P = 0.016$ ) and enhancing stability via overyielding effects<sup>35</sup>, a formal test that would require  
255 monocultures.

256

257 At the larger scale, fertilization reduced the strength of the relationship between beta diversity  
258 and gamma stability by reducing the strength of the relationship between beta diversity and  
259 spatial asynchrony (standardized path coefficient = 0.20 in Fig. 3a vs. standardized path  
260 coefficient = 0.03 in Fig. 3b). This result provides evidence that fertilization can reduce the  
261 stabilizing role of spatial asynchrony among initially dissimilar communities. We did not find  
262 evidence that this was due to a negative feedback of changes in beta diversity caused by  
263 fertilization on gamma stability (Extended Data Fig.2b, Extended Data Fig.9b). The positive  
264 relationship between beta diversity and spatial asynchrony, and the negative impact of  
265 fertilization on that relationship, suggests that the spatial insurance effect caused by variation  
266 in species composition among local communities may be disrupted in a eutrophic world.

267

268 Our results support the idea that asynchronous dynamics among species in species-rich  
269 communities play a stabilizing role and show that this effect propagates to larger spatial  
270 scales<sup>21,26</sup>. Furthermore, our study is the first to report the positive association between beta  
271 diversity and gamma stability through spatial asynchrony in real-world grasslands.



272 Importantly, fertilization reduced the contribution of biodiversity to these stabilizing  
273 mechanisms at both scales, diminishing the local and spatial insurance of biodiversity on  
274 stability. Such diminished insurance effects lead to a reduced ecosystem stability at larger  
275 scales. Future climate will be characterised by more variability including more frequent  
276 extreme events<sup>13</sup>. Our results indicate that preserving ecosystem stability across spatial scales  
277 in a changing world requires conserving biodiversity within and among local communities.  
278 Moreover, policies and management procedures that prevent and mitigate eutrophication are  
279 needed to safeguard the positive effects of biodiversity on stability at multiple scales.

280

## 281 **Methods**

282

### 283 *Study sites and experimental design*

284 The study sites are part of the Nutrient Network (NutNet) experiment (Extended Data Table 1;  
285 <http://nutnet.org/>)<sup>27</sup>. Plots at each site are 5 × 5 m separated by at least 1 m. All sites included  
286 in the analyses presented here included unmanipulated plots and fertilized plots with nitrogen  
287 (N), phosphorus (P) and potassium and micronutrients (K) added in combination (NPK<sub>+</sub>). N, P  
288 and K were applied annually before the beginning of the growing season at rates of 10 gm<sup>-2</sup> y<sup>-1</sup>  
289 <sup>1</sup>. N was supplied as time-release urea ((NH<sub>2</sub>)<sub>2</sub>CO) or ammonium nitrate (NH<sub>4</sub>NO<sub>3</sub>). P was  
290 supplied as triple super phosphate (Ca(H<sub>2</sub>PO<sub>4</sub>)<sub>2</sub>), and K as potassium sulphate (K<sub>2</sub>SO<sub>4</sub>). In  
291 addition, a micronutrient mix (Fe, S, Mg, Mn, Cu, Zn, B and Mo) was applied at 100 gm<sup>-2</sup> y<sup>-1</sup>  
292 to the K-addition plots, once at the start of the experiment but not in subsequent years to avoid  
293 toxicity. Treatments were randomly assigned to the 25 m<sup>2</sup> plots and were replicated in three  
294 blocks at most sites (some sites had fewer/more blocks or were fully randomised). Sampling  
295 was done in 1 m<sup>2</sup> subplots and followed a standardized protocol at all sites<sup>27</sup>.

296

### 297 *Site selection*

298 Data were retrieved on 1 **May 2020**. To keep a constant number of communities per site and  
299 treatment, we used three blocks per site, excluding additional blocks from sites that had more  
300 than three (Extended Data Table 1). Sites spanned a broad envelope of seasonal variation in  
301 precipitation and temperature (Extended Data Fig.1) and represent a wide range of grassland  
302 types including alpine, desert and semi-arid grasslands, prairies, old fields, pastures, savanna,  
303 tundra and shrub-steppe (Extended Data Table 1).

304 Stability and asynchrony measurements are sensitive to taxonomic inconsistencies. We  
305 adjusted the taxonomy to ensure consistent naming over time within sites. This was usually

306 done by aggregating taxa at the genus level when individuals were not identified to species in  
307 all years. Taxa are however referred to as “species”.

308 We selected sites that had a minimum of four years, and up to nine years of post-treatment data.  
309 Treatment application started at most sites in 2008, but some sites started later resulting in a  
310 lower number of sites with increasing duration of the study, from 42 sites with four years of  
311 post-treatment duration to 15 sites with nine years of duration (Extended Data Table 1). Longer  
312 time series currently exist but for a limited number of sites within our selection criteria.

313

#### 314 *Primary productivity and cover*

315 We used above-ground live biomass as a measure of primary productivity, which is an effective  
316 estimator of above-ground net primary production in herbaceous vegetation<sup>36</sup>. Primary  
317 productivity was estimated annually by clipping at ground level all aboveground live biomass  
318 from two 0.1 m<sup>2</sup> (10 x 100 cm) quadrats per subplot. For shrubs and subshrubs, leaves and  
319 current year’s woody growth were collected. Biomass was dried to constant mass at 60°C and  
320 weighed to the nearest 0.01 g. Areal percent cover of each species was measured concurrently  
321 with primary productivity in one 1 x 1m subplot in which no destructive sampling occurred.  
322 Cover was visually estimated annually to the nearest percent independently for each species,  
323 so that total summed cover can exceed 100% for multilayer canopies. Cover and primary  
324 productivity were estimated twice during the year at some sites with strongly seasonal  
325 communities. This allowed to assemble a complete list of species and to follow management  
326 procedures typical of those sites. For those sites the maximum cover of each species and total  
327 biomass were used in the analyses.

328

#### 329 *Diversity, asynchrony and stability across spatial scales*

330 We quantified local scale and larger scale diversity indices across the three replicated 1-m<sup>2</sup>  
331 subplots for each site, treatment and duration period using cover data<sup>37,38</sup>. In our analysis, we  
332 treated each subplot as a ‘community’ and the collective subplots as the ‘larger scale’ *sensu*  
333 Whittaker<sup>28</sup>. Local scale diversity indices (species richness, species evenness, Shannon and  
334 Simpson) were measured for each community and averaged across the three communities for  
335 each treatment at each site resulting in one single value per treatment and site. Species richness  
336 is the average number of plant species. Shannon is the average of Shannon–Weaver indices<sup>39</sup>.  
337 Species evenness is the average of the ratio of the Shannon–Weaver index and the natural  
338 logarithm of average species richness (i.e. Pielou’s evenness<sup>40</sup>). Simpson is the average of  
339 inverse Simpson indices<sup>41</sup>. Due to strong correlation between species richness and other

340 common local diversity indices (Shannon:  $r = 0.90$  (95% CIs =  $0.87 - 0.92$ ), Simpson:  $r = 0.88$   
341 ( $0.86 - 0.91$ ), Pielou's evenness:  $r = 0.62$  ( $0.55 - 0.68$ ), with  $df = 324$  for each), we used species  
342 richness as a single, general proxy for those variables in our models. Results using these  
343 diversity indices did not differ quantitatively from those presented in the main text using  
344 species richness (Extended Data Fig.5), suggesting that fertilization modulate diversity effects  
345 largely through species richness. Following theoretical models<sup>15,16</sup>, we quantified abundance-  
346 based gamma diversity as the inverse Simpson index over the three subplots for each treatment  
347 at each site and abundance-based beta diversity as the multiplicative partitioning of abundance-  
348 based gamma diversity: abundance-based beta equals the abundance-based gamma over  
349 Simpson<sup>28,42</sup>, resulting in one single beta diversity value per treatment and site. We used  
350 abundance-based beta diversity index because it is directly linked to ecosystem stability in  
351 theoretical models<sup>15,16</sup> and thus directly comparable to theories. We used the R functions  
352 'diversity', 'specnumber', and 'vegdist' from the vegan package<sup>43</sup> to calculate Shannon-  
353 Weaver, Simpson and species richness indices within and across replicated plots.  
354 Stability at multiple scales was determined both without detrending and after detrending data.  
355 For each species within communities, we detrended by using species-level linear models of  
356 percent cover over years. We used the residuals from each regression as detrended standard  
357 deviations to calculate detrended stability<sup>17</sup>. Results using detrended stability did not differ  
358 quantitatively from those presented in the main text without detrending. Stability was defined  
359 by the temporal invariability of biomass (for alpha and gamma stability) or cover (for species  
360 stability and species asynchrony), calculated as the ratio of temporal mean to standard  
361 deviation<sup>14,17</sup>. Gamma stability represents the temporal invariability of the total biomass of  
362 three plots with the same treatment, alpha stability represents the temporal invariability of  
363 community biomass averaged across three plots per treatment and per site, and species stability  
364 represents the temporal invariability of species cover averaged across all species and the three  
365 plots per treatment<sup>14</sup>. The mathematical formula are:

$$366 \quad \text{Species stability} = \frac{\sum_{i,k} m_{i,k}}{\sum_{i,k} \sqrt{w_{ii,kk}}}$$

$$367 \quad \text{Alpha stability} = \frac{\sum_k \mu_k}{\sum_k \sqrt{v_{kk}}}$$

$$368 \quad \text{Gamma stability} = \frac{\sum_k \mu_k}{\sqrt{\sum_{k,l} v_{kl}}}$$

369 where  $m_{i,k}$  and  $w_{ii,kk}$  denote the temporal mean and variance of the cover of species  $i$  in subplot  
370  $k$ ;  $\mu_k$  and  $v_{kk}$  denotes the temporal mean and variance of community biomass in subplot  $k$ ,  
371 and  $v_{kl}$  denotes the covariance in community biomass between subplot  $k$  and  $l$ . We then define

372 species asynchrony as the variance-weighted correlation across species, and spatial asynchrony  
373 as the variance-weighted correlation across plots:

$$374 \quad \text{Species asynchrony} = \frac{\sum_{i,k} \sqrt{w_{ii,kk}}}{\sum_k \sqrt{\sum_{i,j,kl} w_{ij,kl}}}$$

$$375 \quad \text{Spatial asynchrony} = \frac{\sum_k \sqrt{v_{kk}}}{\sqrt{\sum_{k,l} v_{kl}}}$$

376 where  $w_{ij,kl}$  denotes the covariance in species cover between species  $i$  in subplot  $k$  and  
377 species  $j$  in subplot  $l$ .

378 These two asynchrony indices quantify the incoherence in the temporal dynamics of species  
379 cover and community biomass, respectively, which serve as scaling factors to link stability  
380 metrics across scales<sup>14</sup> (Fig. 1). To improve normality, stability and asynchrony measures were  
381 logarithm transformed before analyses. We used the R function ‘var.partition’ to calculate  
382 asynchrony and stability across spatial scales<sup>14</sup>.

383

#### 384 *Climate data*

385 Precipitation and temperature seasonality were estimated for each site using the long-term  
386 coefficient of variation of precipitation (MAP\_VAR) and temperature (MAT\_VAR)  
387 respectively derived from the WorldClim Global Climate database (version 1.4;  
388 <http://www.worldclim.org/>)<sup>44</sup>.

389

#### 390 *Analyses*

391 All analyses were conducted in R 4.0.2<sup>45</sup> with N=42 for each analysis unless specified. First,  
392 we used analysis of variance (ANOVA) to determine the effect of fertilization and period of  
393 experimental duration on biodiversity and stability at the two scales investigated. Models  
394 including an autocorrelation structure with a first-order autoregressive model (AR(1)), where  
395 observations are expected to be correlated from one year to the next, gave substantial  
396 improvement in model fit when compared with models lacking autocorrelation structure.  
397 **Second, we used** bivariate analyses and linear models to test the effect of fertilization and  
398 period of experimental duration on biodiversity-stability relationships at the two scales  
399 investigated. **Again,** models including an autocorrelation structure gave substantial  
400 improvement in model fit (Extended Data Table 2)<sup>46</sup>. We ran similar models based on nutrient-  
401 induced changes in diversity, stability and asynchrony. For each site, relative changes in  
402 biodiversity, stability and asynchrony at the two scales considered were calculated as the  
403 natural logarithm of the ratio between the variable in the fertilized and unmanipulated plots

404 (Extended Data Fig.9). Because plant diversity, asynchronous dynamics and temporal stability  
405 may be jointly controlled by inter-annual climate variability<sup>22</sup>, we ran similar analyses on the  
406 residuals of models that included the coefficient of variation among years for each of  
407 temperature and precipitation. Results of our analyses controlling for inter-annual climate  
408 variability did not differ qualitatively from the results presented in the text (Extended Data  
409 Fig.4). Additionally, to test for temporal trends in stability and diversity responses to  
410 fertilization, we used data on overlapping intervals of four consecutive years. Results of our  
411 analyses using temporal trends did not differ qualitatively from the results presented in the text  
412 (Extended Data Fig.6). Inference was based on 95% confidence intervals.

413 Second, we used structural equation modelling (SEM)<sup>29</sup> with linear models, to evaluate  
414 multiple hypothesis related to key predictions from theories (Table 1). The path model shown  
415 in Fig. 1e was evaluated for each treatment (control and fertilized) and we ran separate SEMs  
416 for each period of experimental duration (from 4 to 9 years of duration). We generated a  
417 summary SEM by performing a meta-analysis of the standardized coefficients across all  
418 durations for each treatment. We then tested whether the path coefficients for each model  
419 differed by treatment by testing for a model-wide interaction with the 'treatment' factor. A  
420 positive interaction for a given path implied that effects of one variable on the other are  
421 significantly different between fertilized and unfertilized treatments. We used the R functions  
422 'psem' to fit separate piecewise SEMs<sup>49</sup> for each duration and combined the path coefficients  
423 from those models using the 'metagen' function<sup>50</sup>.

424

#### 425 **Data availability**

426 Data will be made publicly available upon acceptance via GitHub  
427 (<https://github.com/YannHautier/NutNetStabilityScaleUp>). Source data are provided with this  
428 paper.

429

#### 430 **Code availability**

431 R code of all analyses will be made publicly available upon acceptance via GitHub  
432 (<https://github.com/YannHautier/NutNetStabilityScaleUp>).

433

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541

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554

## 555 **Author Contributions**

556 YH, PZ, KRW, ML and SW developed and framed research questions. YH and SW analysed  
557 the data with help from PZ, KRW, EWS, JEKB, SEK, KJK and JSL. YH wrote the paper with  
558 contributions and input from all authors. EWS and ETB are Nutrient Network coordinators.  
559 The author contribution matrix is provided as Extended data table 4.

560

## 561 **Competing interests**

562 The authors declare no competing interests.

563

## 564 **Additional information**

565 Supplementary Information is available for this paper.

566

## 567 **Materials & Correspondence**

568 Correspondence and requests for materials should be addressed to [y.hautier@uu.nl](mailto:y.hautier@uu.nl) and  
569 [shaopeng.wang@pku.edu.cn](mailto:shaopeng.wang@pku.edu.cn).

570



571 **Table 1. Hypotheses related to key predictions from theories relating biodiversity,**  
 572 **asynchrony and stability within and among interconnected communities.**

Pathway	Hypotheses and mechanisms	References
<b>Within communities</b>		
Species richness -> species stability	Higher plant richness within a community either increases or decreases the temporal stability of species abundances within the community by either decreasing or increasing variation in individual species abundances from year to year.	8
Species richness -> species asynchrony	Higher plant richness within a community provides greater likelihood for asynchronous fluctuations among species to compensate one another when the number of species is higher.	51
Species stability -> alpha stability	Higher temporal stability of species abundances within the community increases the temporal stability of community productivity due to lower variation in individual species abundances from year to year	8,14
Species asynchrony -> alpha stability	Higher species asynchronous responses to environmental fluctuations within the community increases the temporal stability of community productivity because declines in the abundance of some species are compensated for by increases in others, thus buffering temporal fluctuation in the abundance of the whole community (species or local insurance hypothesis).	8,14,19,20
<b>Among communities</b>		
Beta diversity -> spatial asynchrony	Higher variation and dissimilarity in species composition among communities increase asynchronous community responses to environmental fluctuations.	16
Alpha stability -> gamma stability	Higher temporal stability of local communities cascades to larger scales and increase the temporal stability of total ecosystem function at the landscape level	15
Spatial asynchrony -> gamma stability	Higher asynchronous community responses to environmental fluctuations increase temporal stability of productivity at the larger scale because declines in the productivity of some communities are compensated for by increases in others, thus buffering temporal fluctuation in the productivity of interconnected local communities (spatial insurance hypothesis).	15,18

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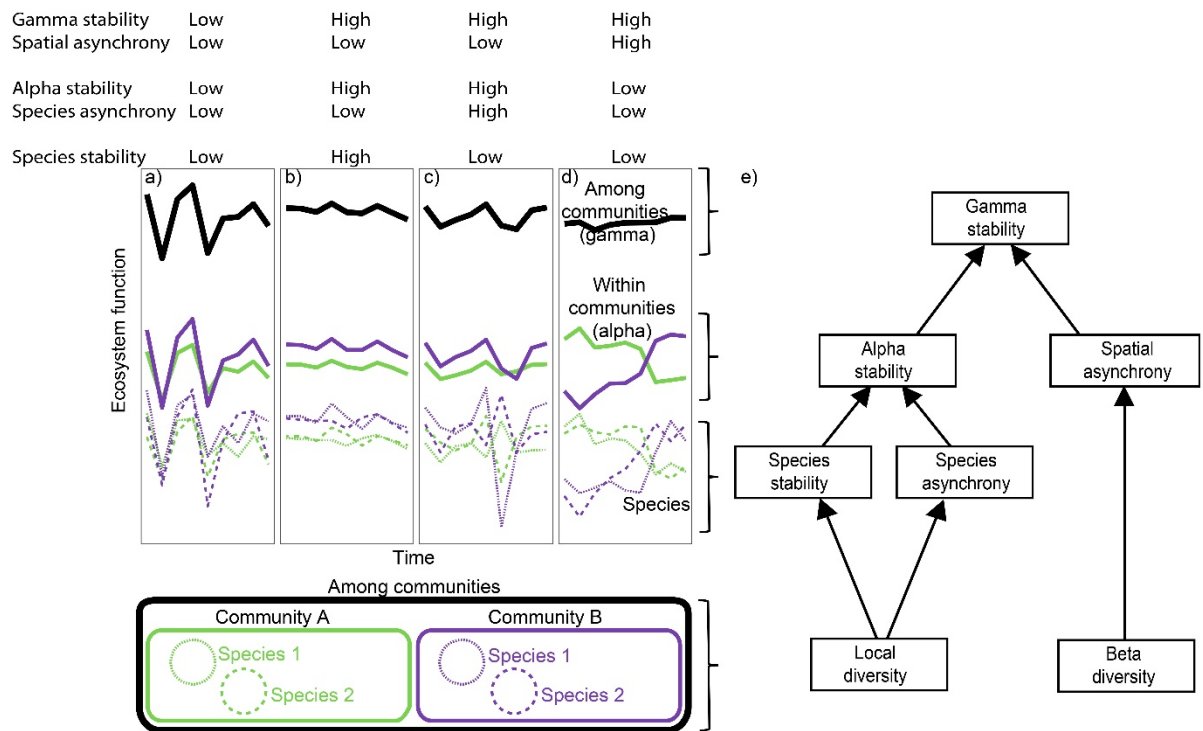
574

575 **Table 2. Summary of meta-analysis results showing tests for differences of model paths**  
 576 **between the unmanipulated control and fertilized conditions, including Cochran Q**  
 577 **statistics for the treatment effect (unmanipulated control versus fertilized condition) with**  
 578 **associated degrees of freedom and p-values.**

Pathway	Cochrane Q statistics	d.f.	P-value
<b>Within communities</b>			
Species richness -> species stability	36.52	1	<0.001
Species richness -> species asynchrony	3.44	1	0.064
Species stability -> alpha stability	0.09	1	0.77
Species asynchrony -> alpha stability	7.15	1	0.008
<b>Among communities</b>			
Beta diversity -> spatial asynchrony	4.52	1	0.034
Alpha stability -> gamma stability	5.27	1	0.022
Spatial asynchrony -> gamma stability	0.11	1	0.74

579

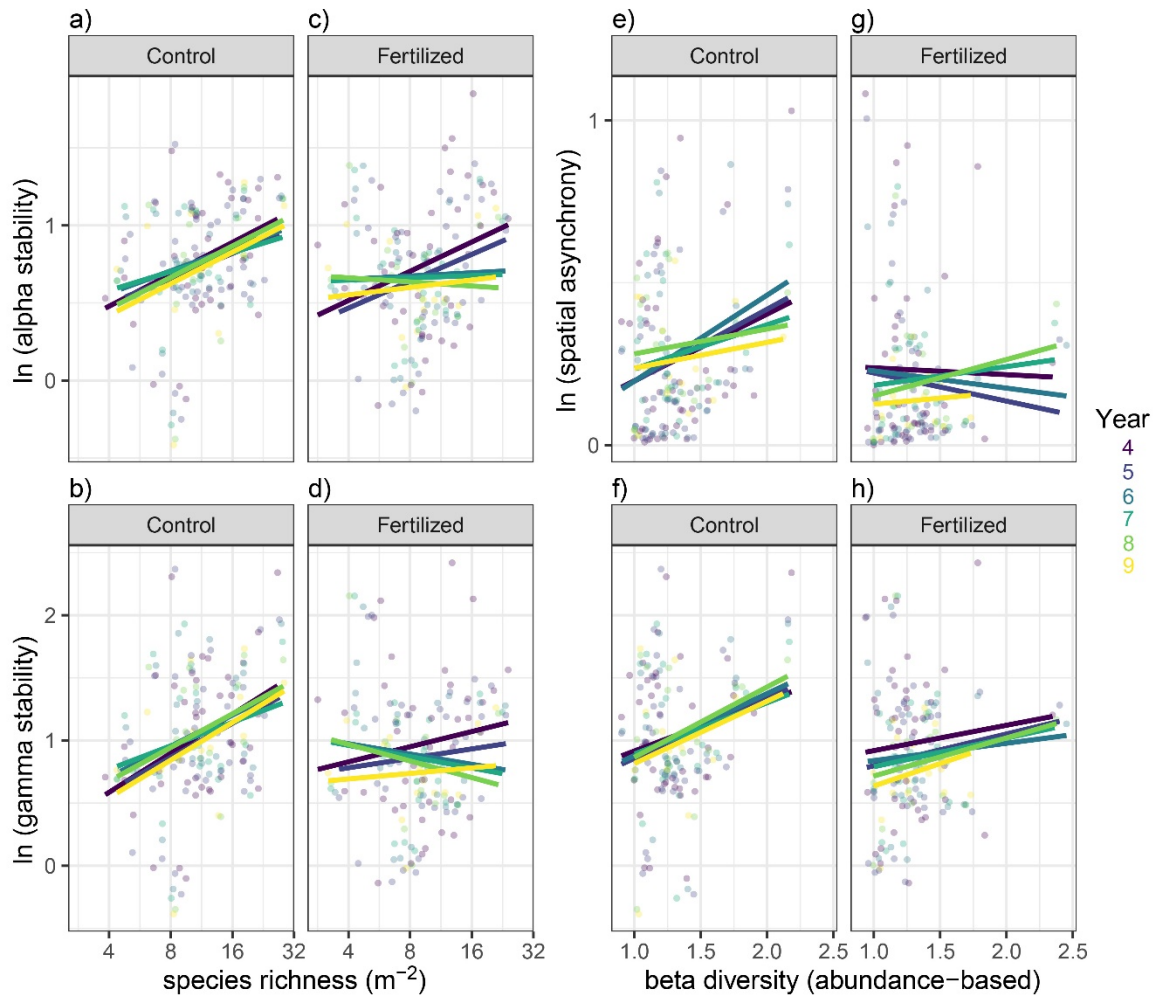
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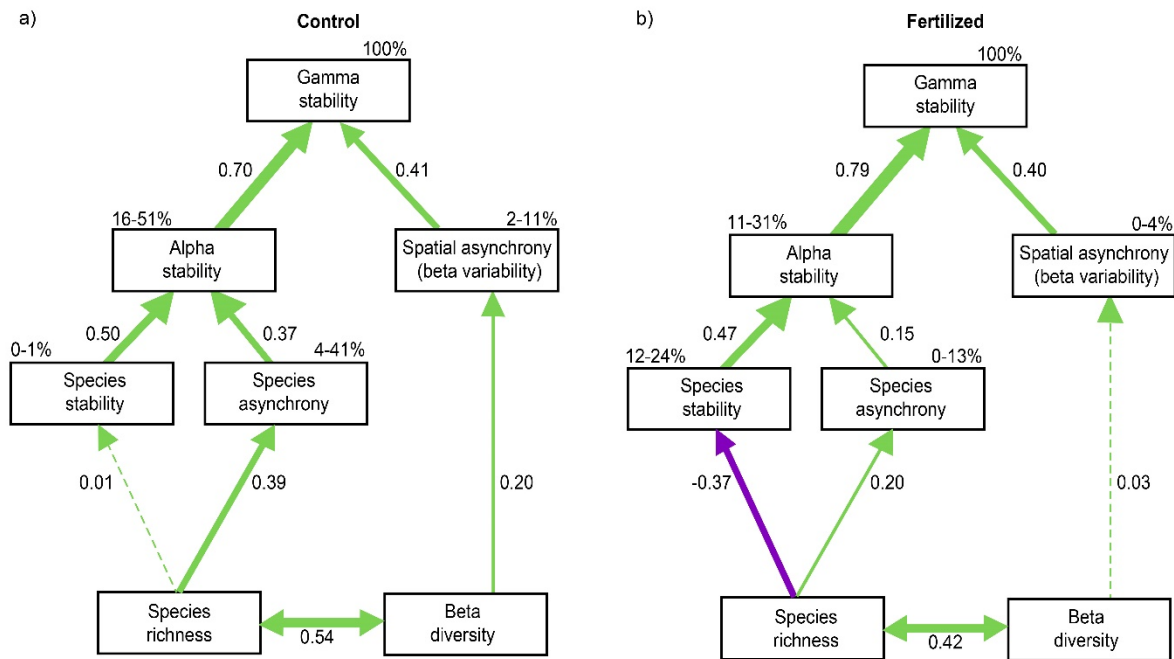
582 **Figure 1. Conceptual figure illustrating the non-exclusive processes by which species**  
 583 **stability, species asynchrony and spatial asynchrony may contribute to stabilize**  
 584 **functioning (such as productivity) within (alpha stability) and among communities**  
 585 **(gamma stability).** a) Low stability and asynchrony of species within communities result in  
 586 low alpha stability that in turn results in low gamma stability under low degree of asynchronous  
 587 dynamics among communities (spatial asynchrony). Relatively high alpha and gamma stability  
 588 may result from b) high species stability and c) high species asynchrony. d) Relatively high  
 589 gamma stability may additionally result from high spatial asynchrony. e) Path analysis used to  
 590 assess the relationship of local and beta diversity with the mechanisms promoting stability at  
 591 multiple spatial scales under unmanipulated control or fertilized condition. Note that species  
 592 names belong to a given community, they could or could not be the same species among  
 593 communities. Adapted from Wilcox et al.<sup>21</sup> and Mellin et al.<sup>52</sup>.

594



595

596 **Figure 2. Impact of fertilization on biodiversity-stability relationships across spatial**  
 597 **scales.** Stability was measured as the temporal mean of primary productivity divided by its  
 598 temporal standard deviation. Relationships were generally consistent among the periods of  
 599 experimental duration considered (Extended Data Table 2). Species richness was positively  
 600 associated with a) alpha (slope and 95% CIs across time = **0.17 (0.08 – 0.26)**) and b) gamma  
 601 stability (**0.27 (0.15 – 0.39)**) in the unmanipulated communities, but unrelated to c) alpha (**0.01**  
 602 **(-0.07 – 0.10)**) and d) gamma stability (**-0.02 (-0.09 – 0.14)**) in the fertilized communities. Beta  
 603 diversity was positively related to e) spatial asynchrony (0.18 (0.06 – 0.30)) and f) gamma  
 604 stability (0.47 (0.19 – 0.74)) in the unmanipulated communities, but unrelated to g) spatial  
 605 asynchrony (-0.01 (**-0.13 – 0.12**)) and h) gamma stability (**0.21 (-0.07 – 0.50)**) in the fertilized  
 606 communities. Note the scale of y-axis differ across panels and this needs to be considered when  
 607 visually inspecting slopes.



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**Figure 3. Summary of meta-analysis results showing the direct and indirect pathways through which biodiversity, asynchrony and stability at multiple spatial scales determines gamma stability under a) unmanipulated control or b) fertilized condition.** Boxes represent measured variables and arrows represent relationships among variables. Numbers next to the arrows are averaged effect sizes as standardised path coefficients. Solid green and pink arrows represent significant ( $P \leq 0.05$ ) positive and negative coefficients, respectively, and dashed green and pink arrows represent non-significant coefficients. Widths of paths are scaled by standardized path coefficients. Percentages next to endogenous variables indicate the range of variance explained by the model ( $R^2$ ) across period of experimental duration.