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Estimates of local biodiversity change over time stand up to scrutiny

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1	Running head: LOCAL BIODIVERSITY CHANGE
2	Estimates of local biodiversity change over time stand up to scrutiny
3	Mark Vellend ^{1,11} , Maria Dornelas ² , Lander Baeten ³ , Robin Beauséjour ¹ , Carissa D. Brown ⁴ ,
4	Pieter De Frenne ^{3,5} , Sarah C. Elmendorf ⁶ , Nicholas J. Gotelli ⁷ , Faye Moyes ² , Isla H. Myers-
5	Smith ⁸ , Anne E. Magurran ² , Brian J. McGill ⁹ , Hideyasu Shimadzu ¹⁰ , Caya Sievers ²
6	¹ Département de biologie, Université de Sherbrooke, 2500 boulevard de l'Université,
7	Sherbrooke, QC, J1K 2R1, Canada
8	² Centre for Biological Diversity and Scottish Oceans Institute, School of Biology, University of
9	St. Andrews, St. Andrews, Fife, KY16 9TH, UK
10	³ Department of Forest and Water Management, Forest & Nature Lab, Ghent University, BE-
11	9090 Melle-Gontrode, Belgium
12	⁴ Department of Geography, Memorial University of Newfoundland, St. John's, NL, Canada,
13	A1B 3X9
14	⁵ Department of Plant Production, Ghent University, Proefhoevestraat 22, 9090 Melle, Belgium
15	⁶ National Ecological Observatory Network, Boulder, CO 80301, USA
16	⁷ Department of Biology, University of Vermont, Burlington, VT 05405, USA
17	⁸ School of GeoSciences, University of Edinburgh, Edinburgh, EH9 3FF, UK
18	⁹ School of Biology and Ecology, Sustainability Solutions Initiative, University of Maine, Orono,
19	ME 04469, USA
20	¹⁰ Department of Mathematical Sciences, Loughborough University, Loughborough,
21	Leicestershire, LE11 3TU, UK
22	¹¹ E-mail: mark.vellend@usherbrooke.ca

23 *Abstract.* We present new data and analyses revealing fundamental flaws in a critique of two 24 recent meta-analyses of local-scale temporal biodiversity change. First, the conclusion that short-term time series lead to biased estimates of long-term change was based on two errors in 25 26 the simulations used to support it. Second, the conclusion of negative relationships between temporal biodiversity change and study duration was entirely dependent on unrealistic model 27 assumptions, the use of a subset of data, and inclusion of one outlier data point in one study. 28 29 Third, the finding of a decline in local biodiversity, after eliminating post-disturbance studies, is not robust to alternative analyses on the original dataset, and is absent in a larger, updated 30 31 dataset. Finally, the undebatable point – noted in both original papers – that studies in the ecological literature are geographically biased, was used to cast doubt on the conclusion that, 32 outside of areas converted to croplands or asphalt, the distribution of biodiversity trends is 33 34 centered approximately on zero. Future studies may modify conclusions, but at present, alternative conclusions based on the geographic-bias argument rely on speculation. In sum, the 35 critique raises points of uncertainty typical of all ecological studies, but does not provide an 36 evidence-based alternative interpretation. 37

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Key words: biodiversity, meta-analysis, species richness, temporal change.

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INTRODUCTION

Patterns of biodiversity over space and time represent the foundation of many ecological theories and conservation priortization schemes. Concerns have been raised (Gonzalez et al. 2016) about two papers that collectively analyzed >250 individual datasets on biodiversity change through time from many parts of the world (Vellend et al. 2013, Dornelas et al. 2014). Both of these studies found that the average magnitude of temporal change in alpha diversity across studies was indistinguishable from zero. Dornelas et al. (2014) additionally showed significant and

46 consistent temporal species turnover, thus highlighting marked shifts in community composition. The concerns of Gonzalez et al., which focus on species richness and diversity, are for the most 47 part typical of those that could be directed at any ecological meta-analysis: different results 48 49 might obtain in different places (underrepresented regions) or times (before people collected data of this nature), and it is possible to find data subsets that deviate from the overall pattern. These 50 concerns were used by Gonzalez et al. to call into question our conclusions. Some aspects of the 51 Gonzalez et al. critique relied on their selective use of data and methods of analysis, while others 52 focused on the nature of the data themselves and accompanying interpretations. 53 54 Here we present analyses, as well as new data, to support the following conclusions: (1) Wellreplicated short-term time series do not provide biased estimates of long-term biodiversity 55 trends. The opposite conclusion presented by Gonzalez et al. was based on two errors in their 56 57 simulation model and calculations. (2) There is no compelling evidence that studies of longer temporal duration show greater biodiversity decline. On this point, the analyses presented by 58 Gonzalez et al. were contingent on a single outlier data point and to unrealistic assumptions 59 60 about model structure, and the results were not robust to the inclusion of additional data. (3) There is no evidence in our original analyses or using a larger, updated dataset that the results 61 were biased due to combining the effects of disturbance with post-disturbance dynamics. In fact, 62 selectively excluding post-disturbance dynamics is itself a source of bias. (4) The ecological 63 literature is indeed geographically biased, a fact discussed explicitly in both Vellend et al. (2013) 64 and Dornelas et al. (2014). The analysis of Gonzalez et al. on this issue supports the undebatable 65 conclusion that new data (in this case from underrepresented regions) might modify conclusions 66 from these meta-analyses, or from any other global-scale ecological meta-analysis. 67

68

SHORT-TERM TIME SERIES DO NOT PROVIDE BIASED ESTIMATES OF LONG-TERM TRENDS

69 As we show below, a key component of the Gonzalez et al. critique is incorrect (i.e., not a matter of selective interpretation). Simulations of species richness (S) over 50-year periods and 70 subsequent calculations of log ratios (log(S_{after}/S_{before})) or slopes of richness on time during 71 72 shorter time intervals (5, 10, 20 years) were used to argue that "Estimates of biodiversity change are systematically biased when syntheses are based on datasets composed primarily of short time 73 series". Gonzalez et al. made two different errors, the first of which applies only to log ratios, 74 the second of which applies to both log ratios and slopes: 75 76 (i) When calculating a mean effect size for "short" windows of time, Gonzalez et al. did not take 77 into account the fact that a log ratio across, for example, a 10-year period is only expected (mathematically) to capture one fifth of the amount of change that occurs over 50 years. In other 78 words, they did not multiply the average of 10-year windows by 5 before comparing with the 50-79 80 year effect size. This is equivalent to the argument that, hypothetically, temperature only went 81 up by 0.5C per decade, so the estimate of the "real" increase of 2.5C over 50 years is biased. 82 (ii) The second problem is less obvious, but no less important, and it accounts for apparent diversity increases in medium-sized time windows (e.g., 20 years) when a 50-year period shows 83 a richness decline initially, followed by an increase, and then a leveling off (see Fig. 1a-c). The 84 85 problem is that with a bounded range of 50 years, "randomly" chosen segments of 20 years severely over-represent the middle portion of the time series. In another well-known ecological 86 context, this is called the mid-domain effect to explain peak species richness at central latitudes 87 or altitudes (Colwell and Lees 2000). However, whereas the boundaries in space are real, the 88 temporal boundaries are not, as time may be considered infinite in both directions. The first 89 point in the time series, for example, is only part of one 20-year segment in the "population" 90 from which the Gonzalez et al. simulations sample, 0:20. The second time point is part of two 91

segments, 0:20 and 1:21, and so on. Time points 20-30, on the other hand, are each part of 20
different segments. So, with the decline in richness happening early during the 50-year time
span, seemingly random samples of 20 years mostly miss the decline, while "detecting" a
transient increase only because it happens to occur in the middle portion of the time series. The
apparent bias detected by Gonzalez et al. is an artefact of their simulation analysis focusing on an
arbitrary bounded time interval (Fig. 1).

If one examines sequential, non-overlapping portions of any length of a given time series, the average log ratio captures precisely the rate of change over the entire time series. Simulations are not required to demonstrate this point, although we provide one corrected example from Gonzalez et al. (Fig. 1), in addition to the following explanation from first principles. Imagine we have a species-richness (S) time series of five points, $t_0:t_4$, and thus four year-to-year transitions. The log ratio from beginning to end is $log(S_4/S_0)$. The average of one-year intervals is:

105 =
$$(\log(S_1/S_0) + \log(S_2/S_1) + \log(S_3/S_2) + \log(S_4/S_3)) / 4$$

$$106 = (\log(S_1) - \log(S_0) + \log(S_2) - \log(S_1) + \log(S_3) - \log(S_2) + \log(S_4) - \log(S_3)) / 4$$

 $107 = (\log(S_4) - \log(S_0))/4$

 $108 = \log(S_4/S_0)/4$

So, as long as we account for the fact that the one-year intervals cover only one quarter of the full time series (i.e., we multiply this by four), we recover the original "target" log ratio for the full time series precisely (see also Fig. 1g). The same result will hold for two-year intervals in this time series, 10-year intervals of a 50-year time series, or any other combination. The same precise mathematical equivalence does not hold for slopes, but it is equally true that there is no

systematic bias introduced by the fact of sampling a subset of a longer time series. An
incomplete sample of the portions of the longer time series will introduce variance (as is always
the case with sampling), but not systematic bias (Fig. 1). The conclusion, based on simulations,
"that short time series can provide unreliable estimates of a known trend" (Gonzalez et al. 2016)
is simply incorrect.

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LOCAL BIODIVERSITY TRENDS IN STUDIES OF DIFFERENT DURATION

120 The argument that short-term time series bias estimates of temporal biodiversity trends was used

by Gonzalez et al. as a springboard to asking whether longer duration studies tend to show

122 biodiversity declines. In this section, we address this issue for the two original studies in turn.

123 Using the data from Vellend et al. (2013), Gonzalez et al. modeled the log ratio of species

richness at the end and start of a study (see previous section) as a function of the duration of that

study, finding a statistically significant (p = 0.04) but weak relationship (Fig. 2a). They

emphasized the conclusion that longer-duration studies tend to show richness declines, although

by allowing for a non-zero intercept, their results also require explaining a nonsensical positive

biodiversity trend in studies that last zero years. If one makes the ecologically realistic

assumption that the log ratio must be zero at duration = 0 (i.e., a zero intercept), not only is the

slope not significant, but its raw value is actually positive rather than negative (Fig. 2B). This
illustrates the potentially major influence of assumptions about model structure on the spurious

132 detection of weak statistical relationships.

Given the controversy sparked by Vellend et al. (2013), we have since expanded the data set by

134 37% to include studies published through the end of 2014 (the original paper had studies

published up to July 2012; see Metadata S1 for data and computer code). The methods were

136 identical to those in Vellend et al. (2013), except that we did not additionally read through the 137 references of all new papers to find additional data sets. With the larger data set of 212 studies (the 2013 paper had 155), there is no significant relationship between local richness change and 138 study duration, regardless of whether one allows for a non-zero intercept (Fig. 2c,d). 139 140 The data in Dornelas et al. (2014) includes studies with diversity estimates for at least three time 141 points, thus allowing the estimation of slopes of diversity vs. time, rather than only before-after 142 log ratios. There is no significant relationship between the diversity-time slope and study duration (Fig. 3a,b). Gonzalez et al. chose instead to calculate log ratios using the data in 143 144 Dornelas et al. (2014; see Dataset S1 in that paper), thereby excluding most of the data used by Dornelas et al, and reported a significant negative relationship between log ratios and study 145 duration (Fig. 3c). Again their analysis allowed for a non-zero intercept; if the intercept is fixed 146 147 at zero – as expected after no time has elapsed – the relationship is not significant (Fig. 3d). In addition, the Gonzalez et al. result is highly sensitive to one outlier, depending not just on a 148 single study (reference 90 in Dornelas et al. 2014), but on a single data point in that study 149 150 (species richness = 43 in 1911, and <20 for the next 90 years). In the absence of that one data point, the relationship is not statistically significant, regardless of whether one assumes a zero or 151 non-zero intercept (Fig. 3e,f). 152

In sum, the evidence provided by Gonzalez et al. to support their claim that longer-duration studies tend to show biodiversity decline is exceedingly weak at best. Their conclusions depend on specific and unrealistic assumptions, and provide negligible predictive value. Whether using the realistic assumption of zero biodiversity change at duration = 0, using a larger data set, taking account of an outlier, or analyzing slopes instead of log ratios, we find no convincing evidence that estimates of biodiversity change depend on study duration. In any given time series, it is

clearly possible (and indeed likely) that trend detection will depend on the particular period of

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160 time analyzed. In our analyses, the observed trends were evenly spread above and below zero for the range of durations, and well-populated with data (<50 years or so). There is thus, at 161 162 present, no evidence to support Gonzalez et al.'s conclusion that longer-duration studies systematically show average local biodiversity declines. We recognize that all of these analyses 163 were carried out with respect to baselines determined by the beginning of the time series 164 involved. There will be cases where ecosystems have lost or gained biodiversity before these 165 observations began, but at present we cannot assess the frequency of these different scenarios. 166 167 EFFECTS OF DISTURBANCE Another concern of Gonzalez et al. (see also Eisenhauer et al. 2016) was the simultaneous 168 inclusion of (i) studies that characterize the effects of disturbance ("impacts") and (ii) studies that 169 characterize recovery from disturbance ("recovery"). If our goal had been to assess only the 170 direct effects of disturbance (e.g., Supp and Ernest 2014), this would indeed have been 171 172 inappropriate. But that was not the goal of either original paper. In a world with fire, grazing, logging, and other disturbances of varying intensity and frequency, to include only the effects of 173 such disturbances and to ignore locations recovering from past disturbances constitutes the 174 175 introduction of a bias in itself. Gonzalez et al. pursued this line of inquiry by selecting studies classified as "post-disturbance", "post-fire", and "cessation of grazing", for elimination from the 176 Vellend et al. (2013) dataset (i.e., leaving the impact-only data subset). 177 178 Unlike the primary analyses in Vellend et al. (2013), Gonzalez et al. analyzed raw log ratios 179 rather than those expressed as change per decade, and they weighted studies by the square root of sample size. The result was a distribution of effect sizes with an upper confidence limit that was 180 slightly negative (1% loss). However, this significant result depends on the combined influence 181

182 of the two analysis modifications introduced by Gonzalez et al. Using the original unweighted 183 analyses and only the data reported in Vellend et al. (2013), the 95% credible intervals for the impact-only data subset selected by Gonzalez et al. overlap zero both for raw log ratios, if only 184 slightly ([-0.095, 0.012] or between 9% loss and 1% gain), and for change expressed per decade 185 ([-0.061, 0.031]). More importantly, using the larger, updated dataset, the upper credible 186 intervals are well above zero (raw log ratio: [-0.053, 0.034]; per decade: [-0.038, 0.032]), 187 allowing us to reject the conclusion that eliminating "recovery" studies reveals an average 188 decline in local plant biodiversity. 189 190 More generally, we disagree with the argument that disturbances (anthropogenic or otherwise) 191 are generally expected to cause a decline in local biodiversity. Empirical studies testing the effect of disturbance on species diversity find a wide range of results, including no effects, 192 193 positive effects, negative effects, and hump-shaped relationships (Mackey and Currie 2001; Supp and Ernest 2014; Newbold et al. 2015; Vellend et al. 2017). Even logging – implicitly 194 emphasized in Gonzalez et al.'s analysis of forest cover change - is often followed by a short-195 196 term increase in local plant diversity (due to colonization of early-successional species) and a subsequent longer-term decline back to levels similar to old-growth forest (Halpern and Spies 197 1995; Duguid and Ashton 2013). In this scenario, capturing only the long-term "recovery" phase 198 would bias results against positive trends rather than negative trends. The opposite scenario (the 199 concern of Gonzalez et al.) certainly exists as well, with biodiversity potentially decreasing after 200 disturbance. Disturbance does not have a unidirectional effect with a single recovery trajectory, 201 so even if studies captured some phases of the disturbance cycle more than others, the 202

203 consequences would not be easily predictable.

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204 In sum, both short-term and longer-term changes in local biodiversity caused by disturbance are 205 relevant to understanding temporal biodiversity trends, and the empirical data indicate highly context dependent effects of disturbance on the magnitude and direction of biodiversity change. 206 THE ECOLOGICAL LITERATURE IS INDEED GEOGRAPHICALLY BIASED 207 Ecological studies of all kinds have been conducted far more often in Europe and North 208 209 America, and nearby waters, than elsewhere. In the case of our meta-analyses, we are unable at present to estimate with confidence how local biodiversity has changed in under-recorded or 210 unrecorded regions, such parts of Africa or the Indian Ocean. This is a challenge for global 211 212 analysis of biodiversity change, and we hope that highlighting this challenge will instigate more data collection in these regions. However, while any given subset of data might deviate slightly 213 from the overall pattern, there was no obvious signal that geographic bias led to bias against 214 finding biodiversity decline. For example, in Vellend et al. (2013), the estimated mean log ratios 215 of species richness change over time for South America (N = 12), Asia (N = 9), Australia (N =216 217 5), and Africa (N = 2) were all positive. One could choose to conduct an analysis giving greater weight to these understudied regions: this would shift the estimated central tendency towards 218 biodiversity increases rather than decreases. In addition, the analyses of Gonzalez et al. show 219 220 that one of the original meta-analyses had an underrepresentation of places with high recent human impacts (Vellend et al. 2013), while the other had an overrepresentation of places with 221 high human impacts (Dornelas et al. 2014), and yet both studies show a similar distribution of 222 temporal changes in local diversity. 223

Gonzalez et al. have identified some important axes along which we might improve the future representativeness of biodiversity studies (e.g., regional diversity, human impacts), but ultimately only new data from underrepresented regions can speak directly to what is happening

227 in those parts of the world, and thus prompt a potential re-assessment of conclusions. Local 228 biodiversity change is very much dependent on specific, local circumstances, and new and interesting results from poorly known regions may well emerge in the future. Improving the 229 230 spatial representation of these regions is a high priority in obtaining better estimates of local biodiversity change. 231 232 In sum, Gonzalez et al. present analyses to demonstrate a point noted in both original papers: the 233 data are geographically biased. Precisely the same limitation applies to most ecological synthesis and meta-analysis papers (e.g., Cardinale et al. 2012, Hooper et al. 2012, Elahi et al. 234 235 2015, Haddad et al. 2015), in which there was no such vigorous effort to quantify geographic 236 bias and its attendant consequences for limiting the scope of conclusions. We are working with the best data available, and continue to assemble data, to directly document temporal biodiversity 237 238 change at the local scale. Converting natural ecosystems to croplands or parking lots causes a local loss of biodiversity (Newbold et al. 2015), but otherwise there is a great deal of variation 239 but no clear tendency for the net temporal local biodiversity trend to be different from zero 240 241 across the sites in the available data (Vellend et al. 2013, Dornelas et al. 2014, Elahi et al. 2015). TO CONCLUDE 242 243 We agree with Gonzalez et al. concerning the need for better biodiversity monitoring in the future. Our knowledge of a great many places on earth is quite limited, and many drivers of 244 biodiversity change are expected to push in opposite directions (Vellend et al. 2017). For 245 example, non-native species introductions typically increase regional-scale species richness (Sax 246 and Gaines 2003, Winter et al. 2009), and in areas that are currently cold and humid (e.g., 247 temperate-zone mountain tops), species richness is also expected to increase due to climate 248

249 warming (Pauli et al. 2012). On the other hand, nitrogen deposition often causes plant diversity

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250	to decline (Simkin et al. 2016), and for some taxa habitat fragmentation can do the same (Haddad
251	et al. 2015). How different forces balance out in the future can best be determined by systematic,
252	long-term monitoring – a major priority for future research in ecology and conservation.
253	Causes and trends of local biodiversity, and therefore any applied consequences, are just as
254	described: local. The global average across many local trends is thus of applied significance
255	only indirectly, via framing arguments about the consequences of biodiversity change (e.g.,
256	Hooper et al. 2012). Given the data at hand, we can reject the notion of local biodiversity loss as
257	the general rule, and whether new data reveal a ratio of positive vs. negative trends at 50:50,
258	60:40 (positive mean), or 40:60 (negative mean), context dependence and site-specificity would
259	remain the dominant pattern. The most generally applicable statement we can make at present is
260	that in most situations we expect substantial changes in species composition – that is, species
261	turnover - with important implications for biodiversity conservation efforts (Dornelas et al.
262	2014, Magurran 2016).
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329 Figure captions

FIG. 1. Mean \pm standard deviation of log ratios (log(S_{after}/S_{before})) and slopes (species richness vs. 330 time) for repeated samples of short time series sampled from a longer (50-year) duration data set. 331 (a) A reproduction of Fig. S3D from Gonzalez et al., showing one example of species richness 332 dynamics over time that appeared to lead to especially biased results. (b,c) Results of 1000 333 seemingly random samples of different duration conducted according to the methods of 334 Gonzalez et al.; these results appear to show an average positive trend among moderate-duration 335 samples, despite a long-term negative (log ratio) or flat (slope) trend over the full duration. (d,e) 336 Log ratio results when correcting separately for duration (problem (i) in main text) and overlap 337 (problem (ii) in main text); here we see that just accounting for the duration of data subsets 338 removes bias from short-duration samples, while correcting for overlap removes any tendency 339 for positive average trends. (f) Slope results after correcting the overlap problem. (g) Log ratio 340 341 results after correcting for both problems; here the averages are precisely equal to the long-term trend. Note that when correcting for overlap, we only use durations that are multiples of the 50-342 vear total time span. 343

FIG. 2. Relationships between local plant species richness change over time (y-axis) and the duration of a study, modeled assuming either a non-zero y-intercept (a,c) or a zero y-intercept (b,c), using the original data in Vellend et al. (2013) (a,b) or an expanded dataset (c,d; new data points shown in red). The effect size for temporal richness change is expressed as the log ratio of species richness in the final year of study (SR₂) and in the initial year of study (SR₁). Lines represent the estimated effect size with credible intervals. See Appendix S1 for statistical methods and Metadata S1 for all data and computer code.

FIG. 3. Relationships between species richness change over time (y-axis) and the duration of a

- study, using data from Dornelas et al. (2014). Relationships were modeled assuming either a
- 353 non-zero y-intercept (a,c,e) or a zero y-intercept (b,e,f), using either slopes (a,b) or log ratios (c-
- f) to express temporal biodiversity change, and either including one outlier (a-d, "Original
- dataset") or not (e,f, "Modified dataset"). Lines represent the estimated effect size with credible
- intervals. See Appendix S1 for statistical methods and Metadata S1 for all data and computer
- 357 code.





