

A comprehensive formula for decomposing change in community similarity into introduction and extinction events

Xianping Li, Céline Bellard, Feng Hu, Huixin Li

▶ To cite this version:

Xianping Li, Céline Bellard, Feng Hu, Huixin Li. A comprehensive formula for decomposing change in community similarity into introduction and extinction events. Ecography, 2019, 42 (10), pp.1714-1716. 10.1111/ecog.04613 . hal-03252885

HAL Id: hal-03252885 https://hal.science/hal-03252885

Submitted on 8 Jun 2021

HAL is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers. L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.

1	doi: 10.1111/ecog.04613
2	
3	
4	A comprehensive formula for decomposing change in community similarity into introduction and
5	extinction events
6	
7	
8	Xianping Li, Céline Bellard, Feng Hu and Huixin Li
9	
10	
11	X. Li (https://orcid.org/0000-0003-4892-5953), F. Hu and H. Li
12	(https://orcid.org/0000-0003-1058-7610) (<u>huixinli@njau.edu.cn</u>), College of Resources and
13	Environmental Sciences, Nanjing Agricultural Univ., CN-210095 Nanjing, PR China. – C. Bellard
14	(https://orcid.org/0000-0003-2012-1160), Laboratoire Ecologie, Systématique et Evolution, Univ.
15	Paris-Sud, CNRS, AgroParisTech, Univ. Paris-Saclay, Orsay Cedex, France.
16	
17	
18	Abstract
19	
20	
21	

22 (Abstract)

The global phenomenon of biotic homogenization is threatening biodiversity worldwide. To explain 23 24 homogenization, multiple frameworks based on different classifications of introduction and 25 extinction events have been proposed. However, comprehensive and generalized approaches are 26 still lacking. Using mathematical derivation, we present a single comprehensive formula, which 27 demonstrates that similarity changes can be precisely determined by the frequencies of six event 28 types, initial similarity, and total number of species. The formula also shows that the effect 29 directions for all events are type-dependent, while their effect magnitudes generally covary with 30 initial similarity and decrease with increasing species numbers. We argue that future empirical 31 and/or predictive studies of biotic homogenization should pay more attention to these direct factors 32 for a deep understanding of community similarity changes. 33 34

35 Keywords: biotic homogenization, biological invasion, community similarity

36

37 Introduction

38 Biological invasion, climate change, and urbanization are presently altering species compositions 39 worldwide (McKinney and Lockwood 1999, McKinney 2006, Qian and Ricklefs 2006, Magurran et al. 2015, Gossner et al. 2016, Newbold et al. 2018), while changes in community composition are 40 41 generally associated with higher community similarities and losses of community distinctiveness (McKinney and Lockwood 1999, Olden et al. 2004). The underlying processes of homogenization 42 are still unclear and especially lack theoretical underpinnings (Olden and Poff 2003, Rosenblad and 43 44 Sax 2017). It was theoretically and empirically proposed that similarity changes are determined by different events (Olden and Poff 2003, Villéger and Brosse 2012, Rosenblad and Sax 2017). For 45 example, the introduction of the same species to two sites would result in an increase in similarity, 46 47 while the extinction of same native species from two sites would lead to a decrease in similarity. In 48 complement to earlier works on the mathematical description of homogenization (Olden and Poff 49 2003, Villéger and Brosse 2012), Rosenblad and Sax (2017) provided a concise framework of six 50 types of introduction and extinction events with formula derivations to qualify the relative importance of these events on changes in community similarity. They found that the effects of these 51 52 events vary according to event type and initial similarity. However, their conclusions mainly derive from simplified assumptions of species richness and number of events, which prevent us from 53 54 understanding the precise relationships among initial similarity, different event types, and total 55 number of species. Through mathematical derivation (see below) and simulations (Fig. 1), we thus aim to: 1) combine previously identified determinants of change in similarity into a single 56 comprehensive formula; 2) illustrate how these determinants interact with each other; and 3) 57 58 highlight the important role of gradient in species richness in affecting the magnitude of similarity

59 changes.

60 With this aim, and as documented by Rosenblad and Sax (2017) and Longman et al. (2018), the types of introductions and extinctions among assemblage pairs can be characterized into seven 61 classes, six of which (I01, I02, I12, E10, E20, and E21) influence change in similarity (Fig. 1). In 62 63 keeping with Rosenblad and Sax (2017), we use Jaccard's index to quantify community similarity, J = S / T, where J is Jaccard's coefficient, S the number of species shared by communities, and T the 64 total number of species in the two communities. We use the subscripts "ini" and "cur" to indicate 65 the metrics for the initial and current assemblage pair, respectively: for example, J_{ini} is the initial 66 67 similarity, and J_{cur} the current similarity. n_i is the number of species affected by event type *i*. Based 68 on the definitions of event types, we know that $T_{cur} = T_{ini} + n_{I02} + n_{I01} - n_{E20} - n_{E10}$, and $S_{cur} = S_{ini} + n_{I02} + n_{I01} - n_{E20} - n_{E10}$, and $S_{cur} = S_{ini} + n_{I02} + n_{I01} - n_{E20} - n_{E10}$. 69 $n_{\text{I12}} + n_{\text{I02}} - n_{\text{E21}} - n_{\text{E20}}$. Thus, the overall change in similarity is as follows:

$$70 \qquad \Delta J = J_{\rm cur} - J_{\rm ini}$$

$$71 = S_{\rm cur}/T_{\rm cur} - J_{\rm ini}$$

72 =
$$(S_{\text{ini}} + n_{\text{I12}} + n_{\text{I02}} - n_{\text{E21}} - n_{\text{E20}}) / T_{\text{cur}} - J_{\text{ini}}$$

73 =
$$(T_{\text{ini}} \times J_{\text{ini}} + n_{112} + n_{102} - n_{\text{E}21} - n_{\text{E}20}) / T_{\text{cur}} - J_{\text{ini}}$$

74 =
$$[(T_{\text{ini}} \times J_{\text{ini}} + n_{\text{I12}} + n_{\text{I02}} - n_{\text{E21}} - n_{\text{E20}}) - J_{\text{ini}} \times T_{\text{cur}}] / T_{\text{cur}}$$

75 =
$$[(T_{\text{ini}} \times J_{\text{ini}} + n_{\text{I12}} + n_{\text{I02}} - n_{\text{E21}} - n_{\text{E20}}) - J_{\text{ini}} \times (T_{\text{ini}} + n_{\text{I02}} + n_{\text{I01}} - n_{\text{E20}} - n_{\text{E10}})] / T_{\text{curr}}$$

76 =
$$[n_{I12} + (1 - J_{ini}) \times n_{I02} - J_{ini} \times n_{I01} - n_{E21} - (1 - J_{ini}) \times n_{E20} + J_{ini} \times n_{E10}] / (T_{ini} + n_{I02} + n_{I01} - n_{E20} - n_{E20} - n_{E20})$$

77 $n_{\rm E10}$)

Our formula and simulation results rigorously confirm that: 1) more I12, I02, and E10 events tend to lead to homogenization while more I01, E21, and E20 events tend to lead to differentiation (Rosenblad and Sax 2017) (Fig. 1); and 2) for I02 and E20, the effect magnitude of similarity 81 negatively covaries with initial similarity, while the effect magnitudes of IO1 and E10 positively 82 covary with initial similarity. Importantly, our results also reveal the crucial role of the total number 83 of species in determining the magnitude of similarity changes (Olden and Poff 2003) (Fig. 1). Namely, the magnitude of change in similarity for all six events decrease with the increasing 84 85 number of initial species. We further demonstrate that the effects of I12 and E21 are mediated by the initial similarity when Sørensen's index of similarity is adopted (Supplementary material 86 Appendix 1). Because the number of species that are initially shared or unique to different sites 87 88 influences the potential number of extinctions, some extinction or introduction events (e.g. E10, E20, E21, I12) are ecologically constrained by initial similarity and total species richness (Fig. 1). 89

90 Using the formula, we demonstrate that the direction and magnitude of change in community 91 similarity are determined by the frequencies of six introduction and extinction event types, initial 92 similarity, and total species richness. The formula provides a unified theoretical framework for 93 understanding homogenization and differentiation processes, which are documented in numerous 94 empirical studies (Qian and Ricklefs 2006, Villéger et al. 2011, Toussaint et al. 2014, Magurran et al. 95 2015). We also provide an interactive web application for visualizing the simultaneous effects of different factors on similarity changes (<https://similarity-changes.shinyapps.io/sim_app/>). Our 96 97 formula could potentially be used to predict future patterns of similarity changes based on current 98 community compositions and the estimated introduction and extinction rates under future global 99 change (Olden and Poff 2003, Rosenblad and Sax 2017, Longman et al. 2018). The formula could 100 also provide a useful framework in which to explore how other factors (such as spatial variation in 101 human activities) determine the frequencies of different event types, which in turn drive patterns of 102 change in similarity. Our work comprehensively and rigorously combines previously identified

- 103 determinants of change in similarity such as different event types and initial similarity. Further, our
- 104 results emphasize that total richness should be considered when predicting or explaining the

105 magnitude of change in similarity among assemblage pairs.

- 106
- 107 Acknowledgements: This project was funded by National Natural Science Foundation of China
- 108 (no. 41571244).
- 109 **Conflict of interest:** The authors declare no conflict of interest.
- 110

111 **References**

- Gossner, M. M. et al. 2016. Land-use intensification causes multitrophic homogenization of
 grassland communities. Nature 540: 266-269.
- Longman, E. K. et al. 2018. Extreme homogenization: The past, present and future of mammal
 assemblages on islands. Global Ecol. Biogeogr. 27: 77-95.
- Magurran, A. E. et al. 2015. Rapid biotic homogenization of marine fish assemblages. Nature
 Communications 6: 8405.
- McKinney, M. L. 2006. Urbanization as a major cause of biotic homogenization. Biol. Conserv.
 119 127: 247-260.
- McKinney, M. L. and Lockwood, J. L. 1999. Biotic homogenization: a few winners replacing many
 losers in the next mass extinction. Trends Ecol. Evol. 14: 450-453.
- Newbold, T. et al. 2018. Widespread winners and narrow-ranged losers: Land use homogenizes
 biodiversity in local assemblages worldwide. PLoS Biol. 16: e2006841.
- Olden, J. D. et al. 2004. Ecological and evolutionary consequences of biotic homogenization. –
 Trends Ecol. Evol. 19: 18-24.
- Olden, J. D. and Poff, N. L. 2003. Toward a Mechanistic Understanding and Prediction of Biotic
 Homogenization. The American Naturalist 162: 442-460.
- Qian, H. and Ricklefs, R. E. 2006. The role of exotic species in homogenizing the North American
 flora. Ecol. Lett. 9: 1293-1298.
- Rosenblad, K. C. and Sax, D. F. 2017. A new framework for investigating biotic homogenization
 and exploring future trajectories: oceanic island plant and bird assemblages as a case study. –
 Ecography 40: 1040-1049.
- Toussaint, A. et al. 2014. Historical assemblage distinctiveness and the introduction of widespread
 non-native species explain worldwide changes in freshwater fish taxonomic dissimilarity. –
 Global Ecol. Biogeogr. 23: 574-584.
- Villéger, S. et al. 2011. Homogenization patterns of the world's freshwater fish faunas. Proc. Natl.
 Acad. Sci. 108: 18003-18008.
- Villéger, S. and Brosse, S. 2012. Measuring changes in taxonomic dissimilarity following species
 introductions and extirpations. Ecol. Indicators 18: 552-558.



143

144 Figure 1. Relationship between initial similarity and change in similarity for each 145 event type under different initial richness scenarios based on the Jaccard-derived 146 formula. The top diagrams illustrating the events are adopted from Longman et al. 147 (2018); circles indicate communities, and stars the introduced or extinct species. Low, moderate, and high richness were set at 50, 100, and 1000, respectively. For a given 148 149 event type, four instances (orange line for 5 events, blue for 10, green for 50, and 150 brown for 100) were used to demonstrate the effects of event frequency on the 151 relationship, with the numbers of other event types kept as zero.

- 152
- 153