

This is a repository copy of *Preface*.

White Rose Research Online URL for this paper: <u>https://eprints.whiterose.ac.uk/183292/</u>

Version: Published Version

### Article:

Rafajlović, M., Alexander, J.M., Butlin, R.K. orcid.org/0000-0003-4736-0954 et al. (1 more author) (2022) Preface. Philosophical Transactions of the Royal Society B: Biological Sciences, 377 (1846). ISSN 0962-8436

https://doi.org/10.1098/rstb.2021.0491

### Reuse

This article is distributed under the terms of the Creative Commons Attribution (CC BY) licence. This licence allows you to distribute, remix, tweak, and build upon the work, even commercially, as long as you credit the authors for the original work. More information and the full terms of the licence here: https://creativecommons.org/licenses/

### Takedown

If you consider content in White Rose Research Online to be in breach of UK law, please notify us by emailing eprints@whiterose.ac.uk including the URL of the record and the reason for the withdrawal request.



eprints@whiterose.ac.uk https://eprints.whiterose.ac.uk/

## PHILOSOPHICAL TRANSACTIONS B

### royalsocietypublishing.org/journal/rstb

# Preface



**Cite this article:** Rafajlović M, Alexander JM, Butlin RK, Johannesson K. 2022 Preface. *Phil. Trans. R. Soc. B* **377**: 20210491. https://doi.org/10.1098/rstb.2021.0491

One contribution of 11 to a theme issue 'Species' ranges in the face of changing environments (part I)'.

#### Subject Areas:

evolution, ecology

#### Author for correspondence:

Marina Rafajlović e-mail: marina.rafajlovic@marine.gu.se

# Preface

Marina Rafajlović<sup>1,2</sup>, Jake M. Alexander<sup>3</sup>, Roger K. Butlin<sup>2,4,5</sup> and Kerstin Johannesson<sup>2,5</sup>

<sup>1</sup>Department of Marine Sciences, and <sup>2</sup>Centre for Marine Evolutionary Biology, University of Gothenburg, 405 30 Gothenburg, Sweden

<sup>3</sup>Department of Environmental Systems Science, ETH Zürich, 8092 Zürich, Switzerland

<sup>4</sup>Ecology and Evolutionary Biology, School of Biosciences, University of Sheffield, Sheffield S10 2TN, UK <sup>5</sup>Department of Marine Sciences, University of Gothenburg, Tjärnö Marine Laboratory, 452 96 Strömstad, Sweden

MR, 0000-0003-2177-4622; JMA, 0000-0003-2226-7913; RKB, 0000-0003-4736-0954; KJ, 0000-0003-0176-7986

Species' distributions change over time. Although in a given time span a species' range may seem static, it may have undergone major changes in the past and it is likely to change in the future. For example, ranges can change in size by expanding or contracting, or shift in space while either preserving or changing their size. Furthermore, ranges can exhibit changes in the extent of (local) fragmentation, and this may or may not occur jointly with changes in range size and/or shifts. These changes can be driven by modifications to environmental factors that limit ranges, which may be biotic or abiotic, or by evolutionary changes within species, or both. Importantly, a change in a species' range may impact on populations, species, communities or ecosystem functioning in the newly occupied or abandoned habitats, with the invasion of non-native species often expected to induce negative effects. Understanding the dynamics of species' ranges thus provides the core for understanding the dynamics of biodiversity. This is a central task of modern biology, of great relevance to society, not least owing to ongoing global climate change, which has already caused range alterations of many species and loss of biodiversity, with more changes expected.

This theme issue synthesizes existing knowledge and hypotheses on the modification of species' ranges, and extends this knowledge with new hypotheses, empirical and theoretical results. Topics addressed include the role of the environmental conditions, including biotic and abiotic factors, that the species has encountered in the past and is encountering in the present, and the role of species-specific intrinsic properties, including dispersal ability of individuals, intrinsic growth rate, niche requirements, plasticity and adaptive potential for niche evolution, among others.

We hope that the theme issue will contribute to answering questions of the type: How will a range change (if at all) given a scenario of interest, at what rate, and with what consequences? Answering this sort of question is essential to design successful conservation and management actions for conserving biodiversity or mitigating its loss.

This theme issue consists of two parts. Part I begins with a discussion of macroecological patterns in species' ranges and provides a new hypothesis for how these patterns might themselves be generated through range evolution [1]. Three contributions then consider the role of plasticity in influencing range limits [2–4], followed by consideration of effects at range margins of pollen limitation [5] and genetic load [6]. Range expansion can cause surfing of underdominant alleles [7]. Finally, three contributions move towards management considerations, starting with biophysical modelling of dispersal [8] and reconstruction of invasion history [9] and then considering management of human-mediated range change [10].

Part II will include further contributed papers, a general Introduction [11] and a Conclusion that draws together all contributions to the theme issue [12].

Data accessibility. This article has no additional data.

 $\odot$  2022 The Authors. Published by the Royal Society under the terms of the Creative Commons Attribution License http://creativecommons.org/licenses/by/4.0/, which permits unrestricted use, provided the original author and source are credited.

2

Authors' contributions. M.R.: writing—original draft, writing—review and editing; J.M.A.: writing—review and editing; R.K.B.: writing review and editing; K.J.: writing—review and editing. All authors gave final approval for publication and agreed to be held accountable for the work performed therein.

Competing interests. This theme issue was put together by the Guest Editor team under supervision from the journal's editorial staff, following the Royal Society's ethical codes and best-practice guidelines. The Guest Editor team invited contributions and handled the review process. Individual Guest Editors were not involved in assessing papers where they had a personal, professional or financial conflict of interest with the authors or the research described. Independent reviewers assessed all papers. Invitation to contribute did not guarantee inclusion.

Funding. This work was funded by the Hasselblad Foundation Grant for Female Scientists (to M.R.), and by a grant from the Swedish Research Council Formas (to K.J. and M.R.; grant number 2019-00882). R.K.B. was funded by the European Research Council.

### References

- Lancaster LT. 2022 On the macroecological significance of eco-evolutionary dynamics: the range shift–niche breadth hypothesis. *Phil. Trans. R. Soc. B* **377**, 20210013. (doi:10.1098/rstb. 2021.0013)
- Maccagni A, Willi Y. 2022 Niche breadth and elevational range size: a comparative study on Middle-European Brassicaceae species. *Phil. Trans. R. Soc. B* 377, 20210005. (doi:10.1098/rstb. 2021.0005)
- Noer NK, Ørsted M, Schiffer M, Hoffman AA, Bahrndorff S, Kristensen TN. 2022 Into the wild—a field study on the evolutionary and ecological importance of thermal plasticity in ectotherms across temperate and tropical regions. *Phil. Trans. R. Soc. B* **377**, 20210004. (doi:10.1098/rstb. 2021.0004)
- Eriksson M, Rafajlović M. 2022 The role of phenotypic plasticity in the establishment of range

margins. *Phil. Trans. R. Soc. B* **377**, 20210012. (doi:10.1098/rstb.2021.0012)

- Dawson-Glass E, Hargreaves AL. 2022 Does pollen limitation limit plant ranges? Evidence and implications. *Phil. Trans. R. Soc. B* 377, 20210014. (doi:10.1098/rstb.2021.0014)
- Sachdeva H, Olusanya O, Barton N. 2022 Genetic load and extinction in peripheral populations: the roles of migration, drift and demographic stochasticity. *Phil. Trans. R. Soc. B* 377, 20210010. (doi:10.1098/rstb.2021.0010)
- Gilbert KJ, Moinet A, Peischl S. 2022 Gene surfing of underdominant alleles promotes formation of hybrid zones. *Phil. Trans. R. Soc. B* 377, 20210006. (doi:10.1098/rstb.2021.0006)
- Jahnke M, Jonsson PR. 2022 Biophysical models of dispersal contribute to seascape genetic analyses. *Phil. Trans. R. Soc. B* 377, 20210024. (doi:10.1098/ rstb.2021.0024)

- Hudson J, Bourne SD, Seebens H, Chapman MA, Rius M. 2022 The reconstruction of invasion histories with genomic data in light of differing levels of anthropogenic transport. *Phil. Trans. R. Soc.* B 377, 20210023. (doi:10.1098/rstb.2021.0023)
- Holman LE, Parker-Nance S, de Bruyn M, Creer S, Carvalho G, Rius M. 2022 Managing humanmediated range shifts: understanding spatial, temporal and genetic variation in marine nonnative species. *Phil. Trans. R. Soc. B* 377, 20210025. (doi:10.1098/rstb.2021.0025)
- Rafajlović M, Alexander JM, Butlin RK, Johannesson K. 2022 Introduction. *Phil. Trans. R. Soc. B* 377, 20210xxx. (doi:10.1098/rstb.2021.0xxx)
- Bridle J, Hoffmann A. 2022 Understanding the biology of species' ranges: when and how does evolution change the rules of ecological engagement? *Phil. Trans. R. Soc. B* **377**, 20210xxx. (doi:10.1098/rstb.2021.0xxx)