

Meeting report



What is macroecology?

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The symposium ‘What is Macroecology?’ was held in London on 20 June 2012. The event was the inaugural meeting of the Macroecology Special Interest Group of the British Ecological Society and was attended by nearly 100 scientists from 11 countries. The meeting reviewed the recent development of the macroecological agenda. The key themes that emerged were a shift towards more explicit modelling of ecological processes, a growing synthesis across systems and scales, and new opportunities to apply macroecological concepts in other research fields.

Keywords: macroecology; spatial scale; process-based model; theory; ecosystem; disease

1. INTRODUCTION

The idea of macroecology as a distinct field of research has been around for more than two decades [1] and was conceived as a response to the realization that small-scale local processes alone were not able to fully explain the abundance and distribution of species. This led to a broader perspective that searched for generalized patterns at large spatial and temporal scales [2], characterized by the search for statistical relationships to explain the distribution of biodiversity from a historical and geographical perspective [2,3]. Ten years ago, a symposium of the British Ecological Society (BES) was convened with the aim of reconciling divergent perspectives on large-scale ecological patterns. This ‘Causes and Consequences’ symposium set the tone for a decade of research in macroecology [4].

Recently, a macroecology special interest group of the BES was formed. The inaugural meeting brought together a diverse group of researchers to review the evolution of macroecology as a research discipline, highlight recent notable developments and explore new applications. Nick Isaac described the aims of the BES macroecology group, which include providing a forum to share ideas and concepts, promoting data access and standards, showcasing methodological advances and setting the agenda for future research. This was followed by a keynote address from Ian Owens, who presented a personal perspective on the development of macroecology throughout the past decade. Owens argued that macroecology has been revolutionized by a combination of the availability of large molecular phylogenies, high-resolution datasets on geographical distribution, extensive computational power and new analytical approaches. As a result, rapid advances have been made towards answering many of the questions that originally occupied macroecologists, such as variation in body size, geographical range dynamics, and the role of neutral processes. These advances have brought with them a new set of opportunities and challenges [5], many of which were recurrent themes during the day. These themes are summarized below.

2. FROM PATTERN TO PROCESS

The strongest theme that percolated all of the talks was the increased emphasis on the processes that drive biodiversity patterns (see also [5]). This theme was introduced by Owens, who outlined a shift from describing patterns to a search for mechanistic understanding. In other words, the way we address key research questions has changed, notably by the increased use of process-based conceptual models of biodiversity [6]. This theme was further developed by Sean Connolly, who identified a mismatch between the biological reasoning that underpins hypotheses about the drivers of macroecological patterns and the statistical models that are actually fitted to data. Connolly illustrated how this has hindered progress in our understanding of large-scale species-richness gradients, and demonstrated how models based on biological processes can be used to derive testable hypotheses [7]. Although macroecology is relatively advanced in its use of statistical methods, the theoretical basis of the predictions involved is sometimes poorly developed. Connolly argued that the explicit formulation of theoretical models, and the robust derivation of statistical expectations from those models, is one of macroecology’s most significant challenges.

Katrin Böhning-Gaese provided a clear demonstration of how incorporating local processes can influence large-scale patterns of species distributions. For example, projections of the impact of climate change on bird species richness yielded very different results when biotic interactions with tree species were taken into account [8]. Similarly, Trevor Price emphasized that both biotic and abiotic factors can explain large-scale diversity gradients. He showed how niche conservatism is not enough to explain diversity gradients of Himalayan birds, unless competitive interactions are incorporated. Kate Jones and David Redding showed how the spread of a zoonotic disease (Lassa fever) can only be understood with

reference to the distribution of the host (a rat). Moreover, Nicholas Dulvy described how the thermal tolerance of individual organisms underpins the distribution of poikilothermic animals in the oceans, and their responses to recent climate change, but that this was not the case on land [9]. Dulvy speculated that gross differences between marine and terrestrial environments can be attributed to the importance of behavioural thermoregulation and interspecific competition on land, contrasting with the dominance of size-based competition in marine systems.

The increasing focus on mechanistic understanding in macroecology is not confined to this meeting [5,10], and many of the recent attempts to build unified theories in ecology have been process-based [11–14]. A key challenge now is to derive general and testable predictions via robust theoretical modelling, underpinned by biologically reasonable assumptions. Recent progress in this area has been substantial [6], although many current theories may not be testable even for data-rich taxa, such as mammals [15]. Thus, further research to bridge the gap between theory, predictions and data is a priority for the development of macroecology into the future.

3. BREAKING DOWN BARRIERS

Traditionally, macroecology focused on processes operating at large scales (e.g. climatic and phylogenetic), largely ignoring the potential for small-scale processes to generate a coherent signal in macroecological patterns [16]. One reason is the deficit of fine-grained (e.g. population-level) datasets that are replicated over large spatial extent [5]: national monitoring schemes have great potential in this regard [17]. A growing body of evidence, both theoretical and empirical, suggests such signals can be detected (see above). Conversely, Böhning-Gaese showed large-scale abiotic gradients can influence community assembly. One striking example is that the degree of specialization, identified using interaction networks among pollinator and frugivore species, is greater in temperate than in tropical communities, contrary to expectation [18,19]. Böhning-Gaese argued that advances in understanding how ecological patterns are generated at multiple spatial scales, and how they are interrelated, are important steps towards a multi-scale synthesis across ecology.

An additional barrier to progress within ecology in general is the lack of synthesis across taxonomic groups and biomes. Historically, macroecology was no exception, being predominantly focused on terrestrial vertebrates [5], although marine macroecology was well-represented at this meeting. A feature of the presentations by marine ecologists was that the concepts and analyses they use are not exclusive to the marine environment. Connolly's process-based models of species richness are wholly transferrable to terrestrial cases. Dulvy went further, arguing that contrasts between realms can discriminate among hypotheses. For instance, equator-ward range limits on land were previously explained as an artefact of under-sampling in the tropics, but the contrast with changing marine range limits in the tropics, where scientific capacity is also low, suggested that stagnant terrestrial ranges are real [9]. More generally, inter-realm comparative

analyses provide many novel opportunities to test mechanistic macroecological hypotheses [20].

4. NEW APPLICATIONS

The meeting demonstrated well how macroecology has influenced diverse research agendas, further reinforcing its application to public policy on biodiversity [21,22]. Owens argued that the influence of macroecology has been unusually broad and deep at the interface of science and policy, especially around land-use, climate change and biodiversity loss. Thus, a significant opportunity exists for macroecology to remain influential and adapt to changing priorities of stakeholders and funding bodies. Two talks focused specifically on the extent to which macroecological ideas are gaining traction in mapping ecosystem services (MES) and epidemiology.

MES, and the potential trade-offs among them, is ripe for the application of macroecological approaches. Like macroecology, MES examines correlations in space over large scales, for example, calculating the degree of spatial overlap of multiple services. Felix Eigenbrod argued MES should adopt macroecological tools to identify the mechanisms underpinning the distributions of ecosystem services. A further challenge for MES lies in the necessity to consider linkages between the distribution of biophysical stocks and their potential beneficiaries, which is somewhat analogous to modelling overlapping geographical ranges of interacting species. For example, Böhning-Gaese incorporated species richness of fig trees (the stock) into predictive models for frugivorous birds (the beneficiaries) [23]. Therefore, the incorporation of co-occurrence and subsequent interactions within both research agendas may be an area that would benefit from collaboration.

A further case study was presented by Jones and Redding, who argued that biodiversity may provide an ecosystem service of disease regulation, thereby contributing to human health. They contrasted traditional epidemiology, which is highly mechanistic and often treats diseases in isolation, with the emerging field of 'disease macroecology', which searches for general patterns in the emergence of novel diseases [24,25]. Jones described how this approach can address policy-relevant questions about emerging infectious diseases and provide a context for mechanistic models of epidemiology at large spatial scales.

5. CONCLUSIONS

Macroecology has clearly matured from its descriptive, pattern-based, roots and now strives for explicit mechanistic ecological understanding. Key questions about the distribution of organisms in space and time remain central to the research agenda, but the conceptual and analytical approaches have changed markedly [5]. The growth of macroecology as both an applied science and a theoretical endeavour is also remarkable. In conclusion, we identify three key ways in which macroecology could progress: (i) close the conceptual gap between data and theory; (ii) enhance integration of replicated field (i.e. fine-grained) studies across the

macroecological scale; (iii) deepen and extend collaboration across realms, biomes and taxonomic groups (including microbes [26]), in order to determine the extent to which patterns and processes are truly general across all biodiversity.

The symposium was organized by N.J.B.I., S.A.K. and T.J.W., and was funded by the BES. T.J.W. is supported by the Royal Society, S.A.K. and S.R.C. by the Australian Research Council and N.K.D. by Natural Sciences and Engineering Research Council of Canada. We are grateful to Rob Freckleton, Georgina Mace and Albert Phillimore for advice and support, and to all the participants for attending.

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