Abstract. Ecological structures and processes occur at specific spatiotemporal scales, and interactions that occur across multiple scales mediate scale-specific (e.g., individual, community, local, or regional) responses to disturbance. Despite the importance of scale, explicitly incorporating a multi-scale perspective into research and management actions remains a challenge. The discontinuity hypothesis provides a fertile avenue for addressing this problem by linking measureable proxies to inherent scales of structure within ecosystems. Here we outline the conceptual framework underlying discontinuities and review the evidence supporting the discontinuity hypothesis in ecological systems. Next we explore the utility of this approach for understanding cross-scale patterns and the organization of ecosystems by describing recent advances for examining nonlinear responses to disturbance and phenomena such as extinctions, invasions, and resilience. To stimulate new research, we present methods for performing discontinuity analysis, detail outstanding knowledge gaps, and discuss potential approaches for addressing these gaps.

Key words: body mass; competition; discontinuity hypothesis; extinction; function; hierarchy theory; invasion; multiple-scale analysis; nonlinear responses; regime shift; resilience.

INTRODUCTION

Toward the end of the 20th century, ecology underwent a conceptual shift from a linear, continuous view of ecosystem processes and structures to one that emphasized nonlinearity and the discontinuous nature of many variables and processes (Wiens 1989, Solé and Bascompte 2006). Ecosystems are strongly influenced by biotic and abiotic processes that operate over different spatial and temporal scales (Levin 1992, Peterson et al. 1998, Peters et al. 2007). Therefore, although small-scale observations provide an important route to explore ecosystem dynamics, it is critical to understand how patterns and processes observed at finer scales represent those operating over broader spatiotemporal scales, and similarly, how large-scale processes correspond to small-scale phenomena (Levin 1992, Cooper et al. 1998, Scheffer and van Nes 2007). These multi-scale patterns will affect the manner in which ecosystems respond to disturbance operating over different scales (Peters et al.
and organismal traits as they relate to that pattern, as cross-scale patterns in abundance, function, diversity, to address a range of ecological questions regarding analysis. We illustrate how this approach may be used the emerging literature incorporating discontinuity nuisances across a range of data types. Third, we explore we provide and describe tools for evaluating disconti-

nuities within biological fields such as hierarchy theory. Second, ecosystems and draw links to work in other ecological framework that accounts for discontinuities. Here, we outline and evaluate evidence for a conceptual approach allows the identification of scale-specific relationships among ecosystem drivers and processes, habitat structure, resource availability, and organisms. Here, we outline and evaluate evidence for a conceptual framework that accounts for discontinuities within ecosystems and draw links to work in other ecological and biological fields such as hierarchy theory. Second, we provide and describe tools for evaluating discontinuities across a range of data types. Third, we explore the emerging literature incorporating discontinuity analysis. We illustrate how this approach may be used to address a range of ecological questions regarding cross-scale patterns in abundance, function, diversity, and organismal traits as they relate to that pattern, as well as to emergent phenomena such as resilience. Finally, we highlight the potential limits of applying discontinuity theory and analyses to specific systems and the current gaps in knowledge, providing stimulus for new research.

Discontinuities: Framework, Evidence, and Extensions

Conceptual framework

The discontinuity approach is derived from hierarchy theory (Appendix A). Growing evidence from nature and ecological modeling suggests that ecosystem structure and dynamics are dominated by the influence of a small set of plant, animal, and abiotic processes operating at specific temporal periodicities and spatial scales, forming a hierarchy (O’Neill et al. 1986, Holling 1992). Each level in this nested hierarchy of variables is controlled by processes sufficiently different in speed and size to introduce discontinuities in the distribution and pattern of ecosystem attributes such as habitat structure and resource availability (Fig. 1; see Allen and Starr 1982, O’Neill et al. 1986, Kolasa 1989). Thus ecological structure varies with scale and reflects the actions of the particular processes operating at a given scale. Such discontinuous hierarchical patterns of processes, structure, and resources were first proposed in systems theory over 50 years ago (Simon 1962). Some 30 years later, ecologists began applying these concepts to describe and understand a
range of aquatic and terrestrial ecosystems (Levin 1992, Gillson 2004, Gunderson 2008). For example, the scales of food and shelter afforded by the physical structures of a coral reef ecosystem vary from individual coral branches up to multi-reef complexes, with implications for the abundance of associated organisms (Fig. 2a; see Nash et al. 2013). Discontinuous, hierarchical structure is being used to assess hydro-geomorphic processes in fluvial systems (Poole 2002), and the concept underpins considerable work in landscape ecology (e.g., Kolasa 1989, Pavlacky and Anderson 2007, Johnson 2009).

A number of theoretical frameworks have linked patterns in habitat structure to attributes of associated communities (e.g., MacArthur and Wilson 1967, Milne et al. 1992, Brown 1995, Ritchie 1998). The discontinuity hypothesis, as originally framed by Holling (1992), proposed that where ecosystem patterns are persistent over ecological time scales, biological processes unrelated to the original structuring processes will become entrained by and adapted to the pattern across scales. For example, life history, behavioral, and morphological attributes of animals may adapt to the discontinuous landscape pattern as this pattern reflects opportunities for shelter, food, and resources (Fauchald and Tveraa 2006). However, these opportunities are mediated by the scales at which individuals interact with the landscape and exploit resources (Holling 1992, Haskell et al. 2002), and the scales of these interactions are positively correlated with body size (Peters 1983).

Holling (1992) found a correlation between breaks in distributions of animal body masses and discontinuities in structures and processes in the boreal forest of Canada. At about the same time, similar hypotheses were presented in paleontology (Legendre 1986, Travoillon and Legendre 2009). Aggregations of species (or modes; Table 1) along body mass distributions indicate scales at which resources and structure are available to organisms and persist within a given landscape over ecological time scales (Fig. 2). In contrast, gaps (discontinuities or troughs) in the distribution reflect the transition to a new set of structuring processes, and therefore few and highly variable resources (Wardwell and Allen 2009). Because animals themselves often strongly modify their environment, such interactions facilitate and reinforce the resources and structure at specific scales (e.g., Bozec et al. 2012).

**Modeling and empirical evidence**

Discontinuous and multi-modal body size distributions (Table 1) have been observed in numerous ecological systems, including both terrestrial and aquatic ecosystems. Studied taxa include birds (Fischer et al. 2008, Skillen and Maurer 2008, Thibault et al. 2011), reptiles and amphibians (Allen et al. 1999), fish and plankton (Havlicek and Carpenter 2001), and mammals (Lambert 2006, Rodriguez et al. 2008, Wardwell et al. 2008). Separate work looking at other species and community attributes, such as species’ abundances and biomass (Angeler et al. 2011), richness (Warwick et al. 2006), range size (Restrepo and Arango 2008), and occupancy patterns (Hartley et al. 2004) across spatial and temporal scales, show similar discontinuous distributions (Table 1).

Similarities in body size distributions among different taxa within a single ecosystem (that are thus exposed to the same habitat structure), and body size distributions of a single taxa among structurally similar systems, have been presented as evidence of the influence of habitat on body size distributions (Holling 1992, Sendzimir 1998). However, the specific mechanisms driving the link between body size and hierarchical habitat structure need explicit exploration (Robson et al. 2005). Szabó and Meszéna (2006) modeled competitive interactions among species of different sizes and showed that the positive relationship between body size and the scale at which species perceive and use resources (Peters 1983, Laca et al. 2010) will produce discontinuous body size distributions where resources are heterogeneously distributed across scales. Empirical studies exploring the link between habitat structure and body size distributions provide further indications of the importance of scaling of the perception of resources with body size.
For example, the distributions of food and habitat resources at different spatial and temporal scales have been shown to influence body size distributions in deserts (Borthagaray et al. 2012), forests (Fisher et al. 2011), and transition zones between rain forest and savannah habitats (Smith et al. 1997), and availability of shelter to different-sized fish has been linked to body size distributions in reef ecosystems (Nash et al. 2013). Similarly, thinning of tree stands, reducing the fine-scale complexity of forest habitats, influences bird body size distributions, resulting in smaller mean body size (de la Montañá et al. 2006). In contrast, research investigating the influence of habitat structure on invertebrate body size distributions has produced mixed results (Gunnarsson 1992). Marine intertidal communities show evidence of distributions driven by sediment structure (Schwinghamer 1981), whereas seasonal and spatial changes in body size distributions are significant in freshwater sediment infaunal communities, suggesting that habitat structure is less important in this context (Stead et al. 2005).

The discontinuity hypothesis represents one of a number of proposed drivers of body size distributions (e.g., Brown et al. 1993, Hubbell 2001, Scheffer and van Nes 2006). However, these drivers are, by and large, complementary as they reflect multiple mechanisms operating at distinct scales. Hypotheses such as community interaction, biogeographical, phylogenetic, and discontinuity hypotheses explain pattern and allometry at distinctly different spatial and temporal scales (reviewed in Allen et al. 2006). The phylogenetic hypothesis, for example, is appropriate at continental scales (Cassey and Blackburn 2004), whereas the discontinuity hypothesis is relevant at regional scales (Allen et al. 2006).

There has been some debate regarding the relative importance of the discontinuity hypothesis vs. emergent neutrality in driving discontinuous distributions at similar scales. The latter hypothesis proposes that competitive interactions alone can generate a discontinuous body mass distribution, although the locations of the modes are the result of stochasticity and so differ from community to community (Scheffer and van Nes 2006).
This is in contrast to the discontinuity hypothesis, which proposes that the location of modes is driven by biota interacting with habitat structure, and therefore would be similar among sites characterized by the same habitat. Thibault et al. (2011) examined biomass size spectra of breeding and overwintering birds at multiple scales and found strong consistency in the number and location of the modes, suggesting non-stochastic structuring processes at work. In contrast, Vergnon et al. (2012) found evidence supporting emergent neutrality in plankton communities, although their treatment of migrants may not be applicable to terrestrial systems, and their representation of the discontinuity hypothesis should be modified; the discontinuity hypothesis predicts weaker interactions among species operating at different scales compared with those operating at similar scales (Fig. 3); this is contrary to the idea of no interaction as suggested by Vergnon et al. (2012). These contrasting results may be a function of the relative complexity of the different ecosystems under study. More work is needed to understand the mechanisms responsible for body size distributions at defined scales and in a wider range of ecosystems to assess scale- and system-specific factors that may influence this relationship (Sendzimir et al. 2003, Robson et al. 2005, Yvon-Durocher et al. 2011).

**Extensions to original framework**

Coincident with the growing body of evidence for discontinuities in numerous ecological systems, a number of species’ attributes have been shown to be associated with discontinuous body mass patterns. These include invasion, extinction, high population variability, migration, and nomadism (Allen et al. 1999, Allen and Holling 2002, Wardwell and Allen 2009). Additionally, the roles that species play and the distribution of the functional attributes of these species within and across scales may strengthen the resilience of ecological systems (Peterson et al. 1998, Walker et al. 1999). Peterson et al. (1998) expanded upon Holling’s (1992) discontinuity hypothesis by proposing that functional diversity within body mass aggregations and redundancy of functional groups across body mass aggregations (i.e., scales) support system resilience (see Applications section). Despite these advances, much of the potential of evaluating discontinuities and their implications for addressing a broad range of ecological questions remains unexplored.

**Applications of Discontinuity Analysis**

Evaluating and analyzing data for discontinuous patterns (Table 2) has two primary uses. First, it is an independent method for identifying “intrinsic” scales (Table 1, Fig. 1) of pattern and process in ecosystems. Second, it provides a platform from which to assess the distribution of key traits or processes within and across the scales of any given system. As a result, it may be used to explain cross-scale patterns such as abundance, functionality, diversity, and organismal traits as they relate to that pattern, as well as emergent phenomena such as resilience.

**Identification of scales**

Incorporating a multi-scaled perspective into empirical research remains a key issue, with choice of discrete scales often being arbitrary (Addicott et al. 1987, Roubicek et al. 2010), with the inherent danger that findings are an artifact of ad hoc scale choices and effects (Wiens 1989, McGeoch and Gaston 2002, Halley et al. 2004, Lechner et al. 2012).

Selecting scales for investigation and analysis that are relevant to the particular individual, population, or community is, therefore, a goal of effective ecological research. This has resulted in the development of a range of multi-scale methods for identifying “intrinsic” scales in ecosystems, largely arising out of work on hierarchy theory in landscape ecology (Wu and Li 2006). However, the information needed to make such informed decisions is often considerable (Addicott et al. 1987). Discontinuity analysis provides a method for detecting underlying scales of process and structure in a system, which is not dependent on arbitrary methodological choices and is relatively data inexpensive, using simple proxies such as animal body size (Appendix B; Holling 1992, Wardwell and Allen 2009) or how patterns change across scales (Bradbury et al. 1984, Hartley et al. 2004).

Such analyses present a number of important opportunities, including the ability to: (1) differentiate between systems exhibiting scale invariance of variables and processes (i.e., consistent patterns across scales) vs. those with discrete, “intrinsic” scales (see “power laws” and “scale domains” in Appendix B; Wiens 1989, Kerkhoff and Enquist 2007); (2) reduce the arbitrariness of scale selection and increase the likelihood of designing effective multi-scale studies (Wheatley and Johnson 2009); (3) delimit the appropriate scales for ecological surrogates (Hartley et al. 2004, Januchowski-Hartley et al. 2011); (4) predict congruence in the response of species to disturbance or environmental drivers (Peterson et al. 1998, Chen et al. 2011); and (5) partition out variance associated with scale effects prior to running other analyses. An example of such an analytical integration is seen in tests for priority effects in Hawaiian avifauna (C. R. Allen, M. P. Moulton, and C. S. Holling, unpublished data). Inhibitory priority effects describe the negative influence of species already present at a site on the colonizing ability of new species (Belyea and Lancaster...
The strength of the negative relationship between the number of species present in the Hawaiian bird community and the success of introduced species was strengthened when body size was accounted for, i.e., membership within a specific body size aggregation was used as a blocking factor in the analysis of variance. This outcome is linked to scale-specific competitive interactions; species within the same aggregation are predicted to experience greater interspecific competition than with species in other aggregations (Fig. 3; see Peterson et al. 1998). As a result, priority effects are stronger in aggregations containing greater numbers of species (C. R. Allen, M. P. Moulton, and C. S. Holling, unpublished data).

Identifying nonlinearities and regime shifts

Interactions among processes operating at different temporal and spatial scales can generate nonlinear behavior (Burkett et al. 2005, Peters et al. 2007). To model these dynamics and minimize “ecological surprises” at local and system-wide scales, development of robust methods for detecting and evaluating nonlinearities is essential (Peters et al. 2004). Discontinuity analysis may be used to explicitly identify nonlinear
patterns within social and ecological systems. For example, these methods have highlighted nonlinearities in both city size and plankton biomass distributions (Garmestani et al. 2007, Angeler et al. 2012). This approach can be extended to characterize nonlinear temporal behavior at the system level, to detect impending regime shifts (Allen et al. 2014).

The capacity for leading indicators, such as recovery rate, rising variance, skewness, or “flickering,” to reveal approaching regime shifts has generated considerable interest (e.g., Scheffer et al. 2009, Wang et al. 2012). Nonetheless, there is concern that proposed metrics may provide an inadequate warning period to allow policy changes in time to address and counteract forecasted shifts (Biggs et al. 2009). Discontinuity analysis may contribute to regime shift detection using existing indicators, by highlighting more sensitive variables that will provide earlier warning signals. For example, rising variance has been presented as a prospective leading indicator (Carpenter and Brock 2006). However, some populations, communities, and abiotic variables are likely to show greater variability than others; thus a method of selecting appropriate variables is needed to inform the design of monitoring programs developed to highlight ecosystem changes (Carpenter and Brock 2006, Wardwell and Allen 2009). Wardwell and Allen (2009) found rising variance in bird population abundance close to discontinuities in body mass distributions (Fig. 4), and proposed that this type of analysis could be used to highlight which variables are likely to show increased variance prior to a regime shift (Wardwell and Allen 2009).

T. L. Spanbauer et al. (unpublished manuscript) present a novel regime shift indicator based on analysis of discontinuities in species abundances over time. Multivariate time series analysis (Angeler et al. 2011) was used to successfully delimit regime shifts in lakes using paleo-diatom data. Where large spatial and temporal data sets are not available (Biggs et al. 2009), other indicators based on discontinuity analysis that require comparatively little data are proposed. The cross-scale pattern of habitat structure and body size distributions is driven by dominant processes and drivers operating over specific spatial and temporal scales (Holling 1992). Therefore, changes in the number or location of discontinuities within habitat or body size

**Table 2. Practical tools for detecting discontinuities.**

<table>
<thead>
<tr>
<th>Method</th>
<th>Discontinuities or multi-modality</th>
<th>Data</th>
<th>Platform</th>
<th>Description</th>
<th>Example references</th>
</tr>
</thead>
<tbody>
<tr>
<td>Hierarchical cluster analysis</td>
<td>discontinuities mean values‡</td>
<td>R: hclust in stats library</td>
<td>identifies groups using successive partitions of the data</td>
<td>Fischer et al. (2007)</td>
<td></td>
</tr>
<tr>
<td>Multivariate time series modeling</td>
<td>discontinuities species abundance</td>
<td>R: quickPCNM in PCNM library</td>
<td>identifies groups of species exhibiting different temporal trends</td>
<td>Angeler et al. (2009, 2012)</td>
<td></td>
</tr>
<tr>
<td>Fractal analysis</td>
<td>discontinuities various</td>
<td>various</td>
<td>identifies groups based on changes in fractal dimension across scales</td>
<td>Krummel et al. (1987), Nash et al. (2013)</td>
<td></td>
</tr>
<tr>
<td>Kernel-density estimation</td>
<td>modality</td>
<td>mean values†‡</td>
<td>R: density within stats library</td>
<td>density function of a variable</td>
<td>Havlicek and Carpenter (2001), Xu et al. (2010), Wang et al. (2012)</td>
</tr>
<tr>
<td>Mixture models</td>
<td>modality</td>
<td>mean values†§</td>
<td>R: OpenBUGS</td>
<td>uses MCMC estimation to model modality</td>
<td></td>
</tr>
</tbody>
</table>

**Notes:** Several methods have been described for identifying discontinuities and multi-modality within the distributions of variables such as body size or biomass. The suitability of these methods varies with respect to the type of data available and the research question (e.g., identifying discontinuities or multi-modality). All techniques have their biases (reviewed in Stow et al. 2007); therefore a combination of methods, followed by triangulation of their respective results, has been identified as the most robust approach (Stow et al. 2007). To date, mean body mass has been primarily used as a measure of body size, although for species with indeterminate growth, other metrics may be more appropriate (Robson et al. 2005). The list of platforms specified is not exhaustive.

‡ Other descriptive statistics, such as mode, median, or maximum value may be used, depending on the research question and data.
† May incorporate a measure of dispersion.
§ May be modified to incorporate abundance.
distributions over time would indicate changes in the dominant processes driving these discontinuous patterns; i.e., would provide evidence of a regime shift (Holling 2001). Such changes have been observed spatially in bird communities, among sites experiencing differing levels of landscape modification (Fischer et al. 2007). Temporal changes in the structure of body size distributions could be used to forewarn of reorganization within a system leading to a new regime. In light of research highlighting the need for robust multi-metric early warning frameworks (Lindegren et al. 2012), discontinuities present an innovative method with which existing metrics can be compared and combined.

Functional distributions, macroecology, and resilience

Species may be grouped according to the functional role that they play in the environment. Functions performed by vertebrate and invertebrate species include pollination, grazing, nitrogen fixation, seed dispersal, decomposition, soil nutrient generation, modification of water flows, opening up patches, and modifying environmental gradients within the landscape (Folke et al. 2004). Body size is a proxy for the scale at which species operate (Peters 1983); therefore, body size distributions can be combined with functional classifications to describe and enumerate the distribution of members of functional groups within and across scales, i.e., the range of scales over which each group delivers its functional role. This information is of direct interest to two fields of ecology: macroecology and resilience (Kerkhoff and Enquist 2007).

There has been an increasing recognition that competitive and other forms of intra- and interspecific interactions need to be incorporated into macroecological studies (Araújo and Luoto 2007), but to date there has been a lack of clarity regarding the influence of competition on local assembly (Gotelli et al. 2010). Where studies have incorporated competition, the strength of interactions has been inferred from membership within species or functional groups (e.g., Gotelli et al. 2010). Peterson et al. (1998) predict that species using similar resources could minimize competitive interactions via differentiation of the scales at which they operate (Fig. 3). Therefore, members of a functional group are more likely to be distributed across scales (and thus body size aggregations) than expected by chance. This nonrandom pattern has been demonstrated in bird and mammal populations (Wardwell et al. 2008), suggesting that functionally similar species within the same body size aggregation are subject to stronger interactions than those operating at different scales (Peterson et al. 1998). As a result, macroecological studies that group species according to body size aggregation would provide a clearer picture of the likely strength of competitive interactions among species using similar resources, and may help to resolve some of the current uncertainties. The effectiveness of this approach was illustrated in the discussion on Hawaiian avifauna priority effects.

Redundancy of species within functional groups is thought to underpin ecological resilience, as it reflects the potential for each group to compensate for the loss of one or multiple species in the face of disturbance, and thus continue to drive ecological processes (Fig. 5a, b; see Walker et al. 1999, Sundstrom et al. 2012). But the value of this redundancy is misleading if each species responds to a disturbance in a similar manner. Species interacting with their environment at different scales are...
likely to respond to disturbances differently (Fig. 5b,c; see Elmqvist et al. 2003); therefore, the scale-specific distribution of functions may be used to quantify the degree of cross- and within-scale redundancy of an assemblage (Allen et al. 2005). This approach, known as the cross-scale resilience model, has been tested on bird community data from southeastern Australia, providing promising results that indicate reduced resilience of modified landscapes (Fischer et al. 2007). However, there is a need to evaluate these metrics in a wider range of ecosystems and taxa. Furthermore, opportunities remain to examine the effects of abundance on functional redundancy (Walker et al. 1999) and to incorporate trait-based functional categorization that transcends the relatively coarse nature of some functional classifications (Fischer et al. 2007).

The distribution of functional groups in time and space may indicate the scales at which species are fulfilling their role. However, the relative impact of organisms of different body sizes is inextricably linked to individual abundance (White et al. 2007). Therefore, decline in the abundance of common species, which may form habitat structure within an ecosystem and/or drive key processes (Gaston and Fuller 2008), has significant implications for functional impact at different spatial and temporal scales, which is not quantified by functional distributions alone. The role of species abundance in resilience is largely unexplored, except in the general sense that minor species can sometimes be functional substitutes for more dominant species whose populations are depressed after a disturbance (Walker et al. 1999). To date, the cross-scale resilience model and empirical evaluations of this model have not incorporated abundance and its influence on the functions that species perform (Peterson et al. 1998, Fischer et al. 2007). There is a clear need to address this gap through the addition of abundance into current models relating biological diversity to resilience.

A more detailed characterization of species roles, other than simple functional groups, is possible through the use of multidimensional functional space indices. This approach has been used to examine drivers of community assembly (Ackerly and Cornwell 2007), the delivery of ecosystem processes (Pakeman 2011), and has been proposed as a way of predicting the response of communities to specific disturbances (Mouillot et al. 2012). Mouillot et al. (2012) discuss body size as one of a number of possible traits that may vary in response to disturbance. Therefore, classifying species according to body mass aggregation in a functional trait-based analysis would explicitly group species operating at similar scales and thus incorporate scale-specific response to disturbance, increasing the sensitivity of such analyses.

The resilience of an ecosystem to specific disturbances may be related to connectivity among habitat patches. This connectivity may be passive (e.g., propagules) or due to mobile links, i.e., individuals moving between areas (Nyström and Folke 2001). Characterizing distributions of function across scales will highlight the likely spatial extent of mobile links and identify vulnerabilities due to a narrowing of the range of scales over which an assemblage is functionally effective (Nyström 2006). For example, large reef fish are subject to extreme fishing pressure in certain areas, resulting in the removal of those species that operate over large scales and thus provide critical linkages among locations across the broader seascape (Jackson et al. 2001, McAuley et al. 2012). This loss has significant implications for the connectivity and spatial resilience of coral reefs, and limits the likelihood of mobile links connecting undamaged reefs with those impacted by disturbance (Fig. 5).

**Extinctions and invasions**

The rising number of invasions by nonindigenous species and extinctions in terrestrial and aquatic environments are of serious concern (Pimentel et al. 2005, Vie et al. 2009). These changes are often associated with significant modifications to habitats and food webs, with important ramifications for the delivery of ecosystem services and the maintenance of key ecosystem processes (Simberloff et al. 2013). For example, in New Zealand, functional extinction of some bird species has significantly reduced pollination of endemic plant species, leading to reduced plant density (Anderson et al. 2011). The cost associated with the impact of invasive species in the United States alone is estimated to exceed US$100 billion per annum (Pimentel et al. 2005).

Predicting the likelihood of a species either becoming extinct or being introduced and then successfully establishing a breeding population, is critical for management and for mitigation efforts. However, such prediction is extremely difficult due to the range of species-, community-, and habitat-level factors influencing the decline of species and the success of invasions (Brook et al. 2008, Hayes and Barry 2008, Harnik et al. 2012). Nonetheless, proximity to the edge of body mass aggregations has been found to be a significant predictor of invasion success for both bird and mammal species, and of extinction risk among mammals (Allen et al. 1999, Allen 2006). Edges of body mass aggregations are associated with increased variability in abundance (Wardwell and Allen 2009), and are linked to less predictable resource availability (Fig. 4; see Wiens 1989). Consequently, these edges represent locations where species may be more susceptible to extinction or more able to exploit opportunities (Allen et al. 1999, Allen 2006). The strength of using proximity to a discontinuity in the body mass distribution as a predictor is that it incorporates both community- and habitat-level factors. Specifically it indicates the likely level of competition experienced by an invading species (Fig. 3) and the scales at which resources are available to species, because body mass distributions are thought to reflect underlying habitat structure. In ecosystems where body size aggregations are demonstrated to reflect the
underlying scales of pattern–process relationships (Borthagaray et al. 2012), the distribution of extinctions and invasions across body mass distributions can also be used to identify scales particularly vulnerable to change and impact (Cardillo and Bromham 2001, Petchey and Gaston 2002, Woodward et al. 2005). In addition, proximity to discontinuities can be used to predict extinctions or the likely success of invasions of introduced species, prior to their occurrence or establishment, respectively. Considering the global threats that extinction and invasion present (for example, 56 species of amphibians and reptiles have successfully invaded and established breeding populations in Florida alone; Krysko et al. 2011), such a predictive ability is of considerable importance. There is however, a clear need to assess the relationship between discontinuities and invasions or extinctions among taxa other than birds and mammals.

**FUTURE DIRECTIONS**

The discontinuity hypothesis provides a conceptual framework, arising from hierarchy theory, within which to examine the organization of ecosystems. However, much of the potential of this framework is unexploited and presents a fertile arena for original research in a wide range of ecological fields. To date, discontinuity research has primarily focused on adult body size as the variable of interest. Performing such studies on plants and modular organisms with no discrete body size, or in species that experience significant ontogenetic changes and indeterminate growth such as fish, remains a challenge that invites innovative approaches. For example, work by Angeler et al. (2013) used multivariate
time series modeling of invertebrate species abundance in subarctic lakes as a novel way to examine patterns of function and resilience at multiple scales, which did not rely on body size metrics. To broaden the utility of discontinuity analysis, similar efforts are needed to address this challenge across the range of potential applications. Furthermore, although there is an emerging literature on the mechanistic links between habitat and body size distributions, additional studies are needed to understand the mechanisms driving discontinuities in variables such as species biomass (Angeler et al. 2011), richness (Warwick et al. 2006), and occupancy (Hartley et al. 2004).

Our discussion of modeling and empirical evidence illustrates the range of studies providing support for the discontinuity hypothesis. However, to date there has been little exploration of those systems where discontinuities are less likely to be found. The discontinuity hypothesis links cross-scale habitat structure to community attributes, such as body size, over ecological time scales. In ecosystems where such structure is less consistent over time, e.g., pelagic systems with dynamic oceanographic conditions and boundaries, discontinuous signals may not be evident. Some work has looked at discontinuities in the body mass (Havlícek and Carpenter 2001), biomass (Angeler et al. 2012), and abundance (Angeler et al. 2011) distributions of lake system communities; however, little work has investigated discontinuities in marine pelagic systems (but see Vergnon et al. 2009). It therefore remains to be seen whether, in comparatively dynamic pelagic environments, discontinuities consistently arise and may be detected.

Discontinuity analysis offers a powerful tool for investigating cross-scale interactions, as it identifies scale-specific relationships between ecosystem drivers and processes, habitat structure, resource availability, and organisms. As a consequence, it provides a platform from which to assess the distribution of key traits or processes within and across the scales of any given ecosystem, e.g., the distribution of ecological function or invasive species. There is considerable scope to broaden the application of discontinuity analysis across ecosystems and taxa, and beyond its current focus on body size applications.

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LITERATURE CITED


SUPPLEMENTAL MATERIAL

Appendix A
Table comparing hierarchy theory, panarchy theory, scale domains, and power laws (Ecological Archives E095-055-A1).

Appendix B
Table describing different methods used to examine the shape of body size distributions (Ecological Archives E095-055-A2).