Selection on stability across ecological scales

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# Abstract

Much of the focus in evolution has been on the adaptive differentiation among organisms. It is equally important to understand the processes that result in similarities of structure among systems. We discuss examples of such similarities occurring at different ecological scales, from predator-prey relationships (attack rates and handling times) through communities (food web structures) to ecosystem properties. Selection among systemic configurations or patterns that differ in their intrinsic stability should lead generally to increased representation of relatively stable structures. Such non-adaptive, but selective, processes that shape ecological communities offer an enticing mechanism for generating widely observed similarities, and have sparked new interest in stability properties. This non-adaptive systemic selection operates not in opposition to, but in parallel with adaptive evolution.

# Systemic selection generating patterns

In recent years, there has been a great dealof ferment focused on the integration of ecology and evolution [1]. Evolutionary and ecological processes can generate a variety of community configurations, but some are observed in nature more or less frequently than others. Not all of these configurations will be equally feasible ([see Glossary](#_Glossary)) or dynamically stable, such that all member species can persist in sufficiently high abundances that extinction of any one is unlikely. When interpreting broad patterns in ecology we must account for feasibility and stability. Feasible and stable configurations should be observed in nature with a higher frequency than configurations that are strongly intrinsically unstable.

If there is no way for the species to coexist (not feasible), then the configuration is doomed to extinction. In other cases, a feasible equilibrium does exist but it is intrinsically unstable. For example, in the well-studied intraguild predation module an intraguild predator and its prey (both of whom share a third resource) are unlikely to be able to coexist if both predators are equal competitors for the shared resource (Box 1). Instead, we should only expect to observe this module when the intraguild prey is able to outcompete its predator for the shared resource [2–4].

This example illustrates what we suggest to be a much more general principle. Namely, differences in observed frequency of alternative community structures might have little to do with the particular characteristics of the local environments in which such systems exist. Instead, intrinsic system properties determining feasibility and stability underlie the relative success or persistence of different configurations, and they can do so more or less independently of local environmental variation. We expect that such non-adaptive systemic selection can produce many of those recurrent ecological patterns that have been observed in nature over large scales of space and time.

Systemic selection can be a powerful yet simple explanation of many ecological patterns, including ecological allometries, species interaction strengths, food web metrics, and macroevolutionary patterns [e.g., 5,6]. This intellectual protocol for understanding ecological patterns has long been implicit in community ecology [7,8], but recent advances in theory suggest that it is time to make this principle more explicit and formal. We start with an overview of systemic selection, in the hopes of generating a “search image” for this type of process in ecology. Then, we dive deeper into (mostly) community ecology to explore the role of systemic selection on stability in connecting local to macro scales.

# Systemic selection is non-adaptive

Biologists are used to thinking of “selection” as a process that explains adaptation of individuals to local conditions, through a selection regime based on the interactions of phenotypes with a shared environment. However, macro-scale biological systems, such as communities, can also change by the preferential loss of internal configurations that are relatively unstable. This elimination of the unstable is a type of long-term selective process, but one where the selection regime, instead of being based on an interaction of the biological system or its components with local conditions, derives primarily from fundamental properties of the system itself; the process doesn’t “see” the local conditions nor is it activated by environmental change (see Box 2).

A theory that is based on such a systemic selection process can be thought of as “non-adaptive”, in the sense that the selection process it describes does not adapt the system to *local* conditions, even though (if working alone) it changes the system in a given direction. We use *adaptation* as is commonly understood, illustrated by Vermeij’s definition (p. 364 in [9]), “… as a heritable attribute of an entity that confers advantages in survival and reproduction of that entity *in a given environment*.” Vermeij goes on to say that these “advantages” are “*context dependent”* (italics added). A trait that is an adaptation in one environment might not be in a different one. By contrast, what we refer to as systemic selection involves “advantages” that are context *independent*.

 Of course, the properties of these systems likely change in significant ways by ordinary adaptive selection and/or coevolution among their members, and by processes such as migration and drift, tending to generate diversity within and among systems (e.g., communities). However, because systemic selection operates on intrinsic system properties that do not vary with the environment, it can, if not significantly overwritten by other processes, have a major role in explaining the evolution and stable properties of macro-scale biological systems. Such non-adaptive processes tend to promote similarity rather than diversity among ecological systems. A book-length treatment that explores non-adaptive selection more generally is in preparation by Damuth and Ginzburg.

# Constraints on attack rates and handling times

Non-adaptive systemic selection can act on whole communities and ecosystems, but we begin by describing how it can also act on the pairwise interactions of species. Each species in an interacting pair brings with it attributes that may affect its interaction’s feasibility and stability. Attributes that affect large amplitude oscillatory population dynamics, for example, can be considered to affect less stable interactions than attributes affecting dynamics with low or absent oscillations.

The well-studied Rosenzweig and MacArthur [10] model provides a conceptual foundation for understanding how consumer traits such as attack rates and handling times influence oscillatory dynamics in a broader array of predator-prey models [11]. In this model, oscillations arise because the predator’s handling time induces a time delay between prey consumption and predator reproduction, which allows prey to increase in abundance; the ensuing increase in the predator population leads to prey overexploitation and a decline in the prey’s abundance followed by the predators [12]. Oscillatory dynamics are more likely to occur when prey are weakly self-limited (e.g., high carrying capacity) and predators have high attack rates and long handling times [11].

Oscillations in population abundances predispose species, and the interaction modules in which they occur, to stochastic extinctions during phases of low abundance. Disproportionate extinction of interactions exhibiting large amplitude oscillations imposes selection at the population level. The expectation that species with oscillatory dynamics should therefore be uncommon in nature [13] is borne out by analyses of time-series data, which show only a minority (~30%) of species with evidence of consumer-driven cycles [14]. Such cycles are, however, particularly common in tightly coupled interactions between specialist predators and their prey [15].

Because systemic selection tends to cull predator-prey interactions in which oscillations lead to periods of low abundances, predators’ attack rates and handling times should be inversely related, i.e., high attack rates and short handling times (greater overexploitation), low attack rates and long handling times (low exploitation potential), or low attack rates and short handling times (short time delay), which dampen oscillations.

Johnson and Amarasekare [12] establish the veracity of this prediction through analysis of the attack rates and handling times of 57 functional response experiments as well as the functional responses and self-limitation strength of two specialist parasitoids and their herbivorous insect host (Figure 1A) [16]. Likewise, estimating the attack rates and handling times of two generalist intertidal whelk predators feeding on 7 to 33 different prey species [17], Novak (Ph. D. thesis, University of Chicago, 2008) showed that 181 pairwise interactions measured across six sites exhibited combinations of attack rates and handling times consistent with the predicted inverse relationship (Figure 1B).

Empirical evidence thus supports the theoretical prediction that stability of consumer-resource interactions constrains attack rate and handling time combinations to be either both low or inversely related. An important task for future work will be to examine the simultaneous operation of systemic selection due to instability, and ongoing natural selection within each of the coevolved species that influence attack rates and handling times [4].

Similar results have been found based on models of entire food webs. Non-adaptive systemic selection, through stability and feasibility constraints, can restrict observed attack rates in a community to values where new consumers invade but do not over-exploit their resources. The possibility of such selection among otherwise unconstrained attack rates during food-web composition turnover was first demonstrated in a model by Rossberg et al. [18].

Community turnover is essential to this mechanism, as this allows species with a lower attack rate than current residents to invade, at least when residents and invader sufficiently differ from each other to not strongly compete through shared resources and consumers. This could result from the immigration of individuals from a large species pool. Variants with lower attack rates can have higher chances of establishing in a food web and persisting than those with higher attack rates, even though their invasion fitness [19] is lower, if they avoid over-exploiting and extinguishing their main resources.

A detailed analysis of this phenomenon must account for the distribution of trophic interaction strengths among consumer-resource pairs, and trophic niche widths [20]. This analysis thus identifies an “optimal” degree of dietary diversity for consumers, where the risk of resource overexploitation is minimized, matching well the dietary diversity actually found for marine fish [20,21]. Another line of evidence for systemic selection of attack rates is that aquatic size spectra (the distribution of community biomass over individuals of different sizes [22]) should theoretically have the observed power-law forms only if mean attack rates are constrained to appropriate values, at least to an order of magnitude [23].

Thus, systemic selection at the community level, over long time scales, can lead to predictable changes of attack rates and niche widths, because these traits have similar ecological implications across communities of different composition. Traits with dynamical implications that depend more sensitively on community composition, such as some specialized foraging or defense strategies [24], might be less likely to exhibit predictable patterns across space or time.

# Community modules

The structure of ecological networks strongly differs from random, particularly at the level of small-scale structures (network modules or motifs), considered to be the "building blocks" of networks. Network patterns can be altered by changing the number of species through invasion, speciation, and extinction, or by changing the way species interact (e.g., by changing trophic relations or interaction strengths).

 In food webs, some modules are greatly over-represented (e.g., the tritrophic chain)[25,26]. The preponderance of certain motifs could reflect their dynamical properties [26–28], and simulations indeed show that motif profiles have strong influence on network dynamics. If different food web configurations vary in stability, systemic selection should cull out unstable configurations [2,7]. Such systemic selection could operate through two intertwined processes: network assembly and dynamical pruning [29,30]

The degree to which a community is stable can be determined through simulation [28], given assumptions about the equations governing dynamics (e.g., Box 3). The network motifs that are observed in food webs are also the structures with the greatest probability of being stable (Figure 2) [26]. Aside from motif patterns, food webs exhibit a pattern of short food chains that can be explained by stability constraints [8,31]. Food webs built with longer chains are less stable than those with shorter chains [32]. These results match what we expect from systemic selection against instability.

Non-adaptive systemic selection can act on networks via dynamical pruning, favoring configurations that are more likely to lead to system persistence [31]. Similar non-random motif profiles could also arise by network assembly, the effects of which have been found in both mutualistic and trophic networks [33–37]. Only by developing strong models of network assembly can one separate the effect of dynamical pruning from that of network growth and development. Assembly might involve systemic selection, but can also include other processes as well (e.g., priority effects, constraints on colonization).

# Limits of feasible coexistence

When modeling large numbers of interacting species with arbitrary, yet biologically plausible, combinations of traits and interactions it is very difficult to achieve coexistence of all species. During simulations at least some species will inevitably decline to zero abundance. Often this is because there is no equilibrium where all species can coexist (infeasibility) rather than any existing equilibrium point being unstable.

Feasibility constraints can be powerful modes of systemic selection. Models have shown that the number of species that can coexist in equilibrium communities (with no direct competition) cannot be greater than the number of resources mediating indirect exploitative competition [38,39]. When there are more species than resources, some will invariably outcompete the others resulting in a final community with, at most, one species per resource.

Only recently has it become clear [20] that these constraints can become tighter when the competing species target resources selectively, so that most species have just a few – but strong – competitors. In two-level food webs where a community of consumer species feeds on, and competes for, members of a community of resource species, and similarly for host-parasite communities, it has been shown that feasible coexistence of the consumer species becomes unlikely if their number exceeds half the number of resource species [20]. Beyond this point, small changes in model parameters lead to large changes in abundances. While co-existence of more consumers might then still be possible, it would require a degree of fine-tuning of parameters that is rather unlikely.

When several trophic levels are stacked upon each other, the feasibility constraint tightens even further, and only about 1/3 of the number of species at one trophic level are likely to co-exist in the next higher trophic level. The potential explanatory power of this mode of systemic selection matches the consistent observations of a 1/2 species-richness ratio in hosts-parasite communities [20,40,41] and 1/3 richness ratios at adjacent trophic levels in multitrophic food webs [20,42–44], see also Figure 3.

Although other potential mechanisms producing correlations of richness across trophic levels have been proposed [42,43,45], none to date predict specific values for the richness ratio. For size-structured food webs where the adult body sizes of consumers and resources are related by a characteristic predator prey mass ratio (PPMR), systemic selection arising from feasibility constraints implies a decline of species richness with body mass as a power law with exponent $–log⁡(3)/log⁡(PPMR) ≈–0.2$, consistent with numerical and empirical observations [46–49].

# Systemic selection at the community level and beyond

Systemic selection at the scale of entire communities should result in patterns in ecological systems that are more feasible and stable than expected at random. One way to characterize a community is the species by species community matrix, whose off-diagonal elements quantify interspecific interactions, specifically how fast the population of one species changes as a result of small deviations of another species from its equilibrium abundance. The diagonal entries describe the strengths of intraspecific density dependencies.

Even without knowing the magnitude of these interactions, one can still address the question of which kinds of structure have a *selective advantage* (i.e., are more stable), by comparing structures of interactions formalized as *signed directed graphs*. Such a graph describes the community matrix in a *magnitude-independent* way, i.e., only by the pattern of pluses, minuses, and zeros within the matrix. Patterns of interactions that are *sign-stable* (as magnitude-independent community matrices) are robust, in that environmental changes affecting strengths (but not signs) of interactions cannot flip a system from stability to instability. Mathematical results on criteria for sign stability (which are known [50–53]) thus provide qualitative predictions about patterns in community structure that should result.

Sign-stable configurations in nature are rare: simple trophic chains (without omnivory) or a few chains linked by amensalistic or commensalistic links with no cycles. *Horizontally structured* communities (i.e., those without trophic interactions) cannot be sign-stable, but can be stable depending on the strength of interactions. The utility of a particular concept of stability of a system depends on the level of detail in the system description. When a food chain is used to represent a series of stable, horizontally structured aggregates of species, then the chain will be sign stable if and only if omnivory is absent. In a sense, one should expect a sign-stable community of this kind to be selected because it will remain feasible despite variation in parameter values.

However, omnivorous links are frequently observed in trophic networks [54–56]. One must therefore look more closely to identify in community matrices a signature of non-adaptive systemic selection through stability constraints. Given two networks with omnivorous links that are both stable (will return to equilibrium following a small perturbation) the *Matrix Flower* (Box 4) can be used to determine which is more likely to be eliminated because of instability.

We can also try to determine how large the feasibility domain (parameter space compatible with coexistence) is and how this domain is modulated by network architecture and interaction strengths. In a competition system, for instance, we can show that the lower the level of interspecific competition, relative to intraspecific, the larger the domain of feasibility [57]. For mutualistic interactions, with a given number of species, links, and distribution of interaction strengths, feasibility domains increase with network nestedness [58]. A community pattern with a larger tolerance to perturbations in biotic and abiotic factors is, by definition, more structurally stable, and under the hypothesis of systemic selection, should be more frequently observed.

The idea that something analogous to natural selection acts on ecosystems has been around for almost as long as the idea that natural selection acts on species. For example, Lotka [59] claimed that natural selection maximizes the total energy flux through an ecosystem. Lotka’s argument was based on Boltzmann’s [60] statement that entropy, and not raw materials, is the limiting factor for all organisms. This intuition provided the foundation for a distinctive school of ecosystem ecology [e.g., 61,62], whose ideas are not currently widely accepted (although this does not imply they are wrong).

Taking a different approach, Tansley [63] argued that “a kind of natural selection” will tend to favor more stable ecosystems. This view has remained widespread in ecological theory. However, empirically evaluating the degree to which a particular selective mechanism might have shaped ecosystem properties requires knowledge of transient states that have been selected against. Our sample of observed ecosystems is unfortunately "length-biased": because most observations come from single time points, ecosystems that are more persistent are likely to be systematically over-represented, relative to the true distribution of historical persistence times. If ecosystem structure is shaped over time by stability constraints, we should expect unbiased samples (taken through time at fixed points of space) to differ systematically from extant ecosystems, which would represent the stable end products of historical processes.

There have occasionally been claims that natural selection can operate between adjacent “meta-ecosystems”, such that the most stable units leave the most descendants [e.g., 64]. However, it seems likely that such mechanisms will only work if ecosystems replicate themselves in their entirety by occupying new units of space. Non-adaptive selection does not require an analog of reproduction at higher levels of organization than the individual, but does depend upon persistence of definable configurations of species across time.

# Social systems

We have mainly focused on community ecology. Yet processes of non-adaptive systemic selection might also be relevant in explaining many other biological patterns. Consider for instance, structure within populations, in particular, the formation of animal social systems. In small groups in many species – from wasps to human children – dominance hierarchies often reveal linear structures. In a linear hierarchy one animal dominates all others – being aggressive toward them without receiving aggression in turn. A second dominates all but the top, and so on. All possible subgroups of three individuals (triads) in a group must have transitive configurations of attack (A attacks B, B attacks C, and A attacks C) for a hierarchy to be linear. If a hierarchy is not linear, then at least one triad has intransitive attack configurations (A attacks B, B attacks C, and C attacks A).

Traditionally, researchers have explained this pattern of organization using individual-based, adaptive models. They have proposed that differences in individual attributes (e.g., size, weaponry, aggressiveness) determine ranks in the hierarchy [65,66]. Or, they have suggested that feedback from winning or losing contests during group formation determines individual ranks [67,68]: an individual winning a contest increases its probability of winning another contest with a new individual, and an individual losing a contest increases its probability of losing. However, both experimental and analytical work [69,70] demonstrate that these simple biological mechanisms of differences in attributes, or winner and loser effects, are unable account for the observed structure of linear hierarchies. Instead, they are better explained by the differential stability rates of the building blocks of the structures.

In analyses of dynamical, peck-by-peck records of chickens forming hierarchies, Chase and Lindquist [71] discovered that intransitive configurations of attacks were unstable – usually lasting only for brief periods of time before changing to transitive ones. Transitive configurations, in contrast, lasted over 20 times as long as intransitive ones on average. They hypothesized that linear hierarchies are common across many species because, as in chickens, when intransitive configurations of attacks appear, they convert quickly to transitive ones, ensuring that hierarchy structures are almost always linear. This rule (transitive configurations are more stable than intransitive ones), plus the rule that all individuals interact, suffices to generate hierarchies that are almost always linear.

# Concluding remarks

A major focus of evolutionary biologists is upon processes that create and maintain diversity — how organisms become adapted to their various local environments. However, especially with regard to multispecies interaction systems, it is also important to understand the processes that result in similarity among systems. We have argued that, because of system dynamics, the deck is stacked in favor of some (stable and feasible) configurations and against others, so what one observes is a non-random distribution of system architectures, based on intrinsic properties shared across systems. There is a resurgence of interest in stability and selective processes in shaping ecological communities and their component structures (food webs, food chains, etc.). New theoretical approaches and computational resources reveal the potential significance of “non-adaptive” systemic selection processes and highlight key attributes promoting stability. At the same time, stability and feasibility are themselves complex concepts that have not been fully explored. Identification of stability criteria for communities and community modules might depend significantly on how researchers model smaller-scale consumer-resource interactions. Nevertheless, it is likely that systemic selection processes — processes of pruning away unstable configurations — will increasingly be seen to underlie many recurrent structural features of biological systems, within which the more familiar processes of local adaptation operate.

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# Boxes:

# The intraguild predation system (Box 1)

To illustrate the idea of non-adaptive selection, consider an intraguild predation community depicted in Figure I with four slightly different interaction strength configurations. Thicker arrows indicate stronger interactions as defined, for example, by the amount of energy flowing from a prey population to each predator. We could expect to obtain each of these configurations in several ways. For example, interaction strengths may change as species identities change through speciation, or the combination of extinction and invasion. Or they could change from adaptive coevolution among the species, due to variation over space, time or within populations, or due to behavioral changes in individual foraging decisions. However, while many such processes could lead to any of the four configurations, feasibility constraints dictate that we are nonetheless likely to observe one of these four configurations more often in nature than the other three [4,72].

That is, when the intraguild predator and prey are equally efficient competitors for the shared resource (Figure Ia) we expect the intraguild prey to be driven towards exclusion because the intraguild predator has the advantage of consuming its competition. Likewise, the intraguild prey will be driven towards exclusion when the intraguild predator is the superior competitor (Figure Ib), Increasing the strength of the intraguild predator’s effect on the intraguild prey (Figure Ic) thereby increases the degree to which the intraguild prey must be the superior competitor in order for it to maintain a viable population size. Although differences in mortality rates and the presence of alternative resources can weaken and to some degree counteract these feasibility requirements, it is when the intraguild prey is the superior competitor (Figure Id) that we are most likely to observe all species coexisting.

We can therefore say that communities *a-c* are systemically selected against. Further, because such selection is a result of the properties of the community (i.e. the configuration of interaction strengths) and not due to its particular environment, this selection is non-adaptive (see Box 2).

# Non-adaptive or adaptive? (Box 2)

Scientists use the term “adapt” and its derivatives in various ways. For our purposes the process of “adaptation” can be most usefully considered as *adaptation to a specific locally shared environment*. This is the most common usage in evolutionary biology. In emphasizing that for some systems there are systemic selective processes that impose the same selective forces regardless of variation across local environments, we highlight the fact that these processes are selective ones, but for reasons that do not involve adaptation to local conditions.

This perspective contrasts with a more conventional approach toward the evolution of macro-scale biological systems — accomplished by expanding hierarchically the theory of adaptation by natural selection by identifying a hierarchy of units of selection [73]. “Individual” and “population” are replaced with “population” and “community” (or “community” and “population-of-communities”), respectively, and analogous properties of these entities that correspond to key elements of organismic natural selection (e.g., reproduction, heritability, fitness, adaptation, environment) are identified. It has long been recognized that such a theory of the adaptation of higher-level entities to their environments is logically possible, though not without significant issues [73–76](J. Khai Tran, Ph.D. thesis, Stony Brook University, 2011).

In this view, feasibility and stability serve as an important component of community fitness and are considered to depend on the environment. For example, the environment can influence the strength and type of interactions between species. Moreover, overall community fitness also relies on “reproduction” in terms of a community colonizing new areas or influencing the composition of other communities through propagule pressure. Though the process is far more nuanced, in oversimplified terms, adaptive selection at the community level eliminates community types that on average do not persist long enough to produce a like community (absolute fitness less than 1). This is likely the fate of infeasible and highly unstable configurations, barring a reproductive rate high enough to compensate. Furthermore, community selection can favor one stable community type over another according to their relative fitnesses. Similarity among existing communities is explained as the consequence of convergence (limited ways to solve a problem), with the evolutionary process being higher level adaptive selection.

Despite its logical appeal, a consensus has not emerged about how to put in practice such a theory so that it both avoids conceptual difficulties and provides rewarding insights about community evolution. From the perspective of non-adaptive selection, a stable community is not necessarily a community that is “better adapted” to its specific environment. Exploring non-adaptive selection does not require resolving issues about defining local adaptation at the community level. A useful analogy from the physical sciences illustrates this point. Protons, the nuclei of hydrogen atoms, have never been observed to decay while the most stable isotope of the radioactive element francium is very unstable (short half-life). Hydrogen is the most abundant element in the universe, whereas francium is a laboratory curiosity. Yet we would be unwilling to say that hydrogen is better adapted to the universe. Eventually, a fruitful adaptive theory of community evolution, as a counterpart to the non-adaptive processes we discuss here, might be possible.

# Modeling species interactions (Box 3)

Usually, interactions within food webs are modeled with generalized Lotka-Volterra equations. However, although Lotka-Volterra equations can be acceptable in some conditions, they do not apply universally. The Lotka-Volterra functional response model assumes random and homogeneous mixing of prey and predator individuals, and that predators are rare enough to avoid interference. Empirical evidence and theoretical considerations suggest that it is necessary to use different models, particularly with incorporation of predator interference. Extra care must be taken when using non-Lotka-Volterra interactions in food webs models to ensure that the generalized mathematical expression remains logically consistent [77]

An alternative to Lotka-Volterra functional response models, the donor-control model, is a simple case of ratio dependence that has often been observed. This type of interaction might result from non-adaptive selection, but it can also emerge from complex mechanistic models that incorporate conventional adaptive selection [78]. We can generalize the donor-control model to a multispecies case, where several predator and/or prey species interact. For two prey species having a common predator, one prey species can lead to indirect exclusion of the other species (apparent competition). In the case of two predators competing for a single prey species, one species is always excluded by the other, often even in the presence of a top predator. That is, a keystone effect is never observed and the configuration collapses to a three-level food chain. Given donor control, simple food chains will persist or not because of basal production, not because of top-down effects. Intraguild predation can, in some instances, help maintain more complex structures.

# The Matrix Flower (Box 4)

In addition to the famous concept of *Lyapunov* stability with its well-known matrix criterion, there are several other, less well known stability concepts defined in matrix terms: *D*- and *aD*-stability, *sign*-stability, *Volterra dissipativeness*, and others. These stability concepts deal with stability despite various kinds of uncertainty or perturbation. However, few such concepts have clear characterizations via the matrix structure and/or elements, and some just have definitions, with only necessary or sufficient conditions for their verification known. Logical inclusion relations (Venn diagrams) among the special stability subsets in a general space of real matrices *n*×*n* help verify particular properties of a given matrix. Adding a few more conventions to a standard Venn diagram turns it into a 3D *Matrix Flower* [79] whose “petals” represent the subsets of matrices with particular stability properties (Figure I).

Two, relatively small, petals of sign-stable matrices are located in the very “core” of the Flower, while the other stability petals signify the lower levels in non-adaptive selection.

We should keep in mind that having the most stable pattern in interspecific interactions is only one side of the problem. The other side is the feasibility of the community, i.e., the existence of an equilibrium point where all species have strictly positive abundances. Illustrating that feasibility is as crucial as the stability constraint, consider the simple example given by a set of species competing for resources. Moreover, assume that the interspecific interaction matrix has been derived from species overlap in a niche space [80]. In this case, the matrix belongs automatically to the petal of being *Volterra dissipative*. This implies that the community is automatically globally stable as long as there exists a feasible equilibrium point. Therefore, the only remaining question is whether the system will be feasible or not. We can show that there exists a specific domain in the space of carrying capacities (or intrinsic growth rates), in which the community is feasible [53,81]. An open question is how these ideas translate into systems where coexistence arises because of nonlinear feedbacks (e.g., Armstrong-McGehee effects [82,83]).

# Glossary

**Adaptive selection:** a selection process (such as neo-Darwinian natural selection) that can cause adaptation to a local environment, because fitness differences result from an interaction between phenotypes and that shared local environment.

**Environment (of a system):** all local factors external to the system, biotic and-or abiotic (depending upon the system definition).

**Feasibility:** the existence of an equilibrium point or attractor such that all species in the system can coexist.

**Non-adaptive selection:** any selective process that does not act to adapt the members of the population on which it acts to a specific local environment.

**System:** the particular unit of interest, typically involving multispecies interactions but dependent upon the ecological scale of the study. At macro-ecological scales we typically refer to systems as multispecies interaction networks (e.g., food webs, plant-pollinator webs).

**Systemic selection:** non-adaptive selection on systems based on their intrinsic system properties.

**Intrinsic system properties:** the attributes and qualities of the system as a whole, such as the pattern and magnitudes of species interactions, and the densities of participating species.

# Figures

**Legends**

Figure 1: The empirical relationship between handling time and attack rates. (A) Published attack rates (day-1 ind-1) and handling times (day-1) for 57 species, see [11] for details. (B) Empirical relationship between the species-specific handling times (days) and per capita attack rates (prey eaten pred-1 prey-1 m-2 day-1) of two predatory whelk species feeding on their prey (n=181 (M. Novak, Ph. D. thesis, University of Chicago, 2008)). Inset shows the relationship at low attack rates.

Figure 2: A bar plot of the mean relative frequency of the thirteen three-node subgraphs (pictured at the top of the plot) in 50 empirical food webs ordered by their stability. For more details on methods and data see Ref. [26].

Figure 3: Predator-prey richness ratios in freshwater invertebrate communities after [42], compared with predicted ratio of 1/3

Box 1 Figure I: Four hypothetical intraguild predation communities. Arrows point in the direction of energy flow and their width represents interaction strength.

Box 4 Figure I: Logical relations among different types of matrix stabilities (adapted from [79], Fig. 4)