1 A novel semantic theory of the assembly rules of interaction

2 networks

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Abstract

47 In the web of life, every interaction between species tells a story of cooperation, 48 conflict, or chance. For centuries, ecologists have charted these stories to better 49 understand phenomena such as pollination and disease. We have been using the lens of 50 network science to distill them into topological patterns such as nestedness or 51 modularity. Yet, like in the old buddhist parable of the blind monks and the elephant, 52 these separate views have often missed the bigger picture. In this Perspective, we 53 review the history of topological studies of interaction networks, challenge the "black-54 and-white" paradigm, and point out a way forward: the Integrative Theory of Interaction 55 Networks (ITIN). This theory unveils the logic of how interaction networks assemble 56 through sequential, rule-bound processes shaped mainly by resource dissimilarity and 57 interaction type. We show that compound topologies are likely the norm in function-58 oriented, taxonomically-inclusive, well-sampled networks. ITIN is deductive, 59 predictive, and testable, mapping from first principles to real-world systems, thus 60 helping us grasp "the whole elephant". As ecologists confront the polycrisis, from 61 irreparable biodiversity loss to global health threats, ITIN offers a coherent framework 62 to understand, predict, and restore the threads of the web of life.

Introduction

65	Ecological interactions between individual organisms of different species (i.e., species
66	interactions) have drawn the attention of scientists, philosophers, and stakeholders for
67	millennia 1. This broad family of phenomena ranges from mutualistic (both sides gain
68	fitness) to antagonistic (one or both sides lose fitness), and result in crop pollination,
69	plant protection, seed dispersal, parasitism, and disease, among many other outcomes ² .
70	Species interactions have been studied from multiple perspectives and using numerous
71	approaches ³ . One approach that has become mainstream is network science, an
72	interdisciplinary field that provides not only tools for data analysis but also concepts
73	that may be integrated across scientific disciplines to obtain new insights ⁴ . The study of
74	interaction networks as we know it began in the late 19th century (or even earlier),
75	through a classical effort to understand complex relationships in lake food webs ⁵ . Only
76	more recently, between the late 1980s and early 2000s, the focus slowly shifted from
77	food webs to mutualistic networks ⁶ . The intriguing structural patterns of interaction
78	networks, also known as topologies, has become a fashionable research topic in itself,
79	irrespective of the effect they may, or may not, have on the community and ecosystem
80	dynamics 7. This focus, although being an important first descriptive effort, seems to
81	have led to worrisome epistemological biases, such as confounding the ecological
82	processes with the network patterns they generate ⁸ .
83	Biases in topological studies can be compared to the ancient Buddhist parable of the
84	blind monks and the elephant (Figure 1). Once upon a time, a group of blind monks
85	heard that a huge, strange animal had arrived near the temple, and, curious to know
86	what it was like, decided to examine it by touch. Each monk felt a different part of the
87	elephant and drew a different conclusion. To one monk, the trunk felt like a snake. To
88	another, the ear felt like a fan. To others, a leg felt like a tree trunk, the belly like a wall,
89	the tail like a rope, and a tusk like a spear. Each monk described the elephant based on
90	their separate experience, illustrating how partial perspectives can lead to incomplete or
91	conflicting understandings of a greater truth.



Figure 1: The blind monks and the elephant. As in the old Buddhist parable, we have been analyzing only network parts but drawing conclusions for the whole. Currently, mutualistic networks are generally assumed to be nested and antagonistic networks, modular. Gradient networks have been seldom considered, usually being associated with parasitism. Much fewer studies discussed compound topologies, although these have been drawing more interest lately. The image was modified from the original *ukiyo-e* print illustration by Itcho Hanabusa (Wikimedia Commons, public domain).

Not so different from those monks, most ecological studies have focused on interactions in only one context, ignoring that sites with different abundances will yield different networks and observed interaction preferences ⁸. Moreover, by focusing on isolated interaction types, while trying to draw conclusions for all types, those studies ignore how interactions create complex systems with emergent properties ⁹. Recent advances, however, show that networks require a more integrative approach to be fully understood ¹⁰. Integration is especially needed for understanding not only the topological patterns observed in nature, but also how they emerge as a result of ecological and evolutionary processes. In other words, the assembly rules of interaction networks.

In this Perspective, we aim to demonstrate how different views about network topologies and their assembly rules could be integrated, in order to help refocus our research efforts. First, we synthesize the history of network topology studies. Second, we show how the Integrative Hypothesis of Specialization (IHS) provides a comprehensive approach to see the whole elephant, in addition to shedding light on the processes behind the patterns. Third, we point to new directions, as after a series of developments, empirical tests, and additional synthesis, IHS is evolving into a semantic theory, the Integrative Theory of Interaction Networks (ITIN), which we first introduce.

Archetypal topologies

A seminal study ahead of its time in the mid-2000s proposed a taxonomy of network topologies ¹¹. From its perspective, there would be four archetypal topologies (<u>Figure 2</u>) that capture the essence of most real-world systems (but see an alternative taxonomy ¹²).

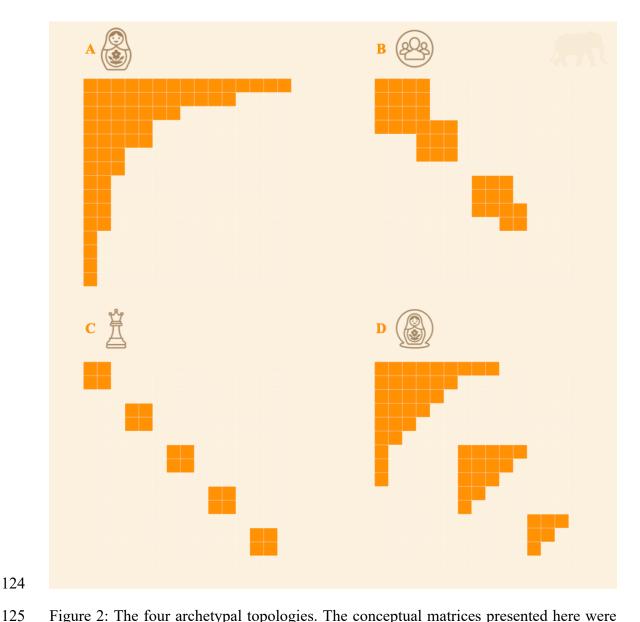


Figure 2: The four archetypal topologies. The conceptual matrices presented here were modified from those originally proposed in a seminal paper ¹¹. (A) A nested topology resembles a Russian doll: nodes with fewer links are connected to a subset of the links captured by the most connected nodes, sequentially, for the rows and columns separately. (B) The modular topology looks like the filter bubbles observed in social media, as the network is divided into cohesive subgroups of nodes (i.e., modules), more densely connected to one another than to other subgroups. (C) In a gradient topology, as in a chess board, there is little to no overlap in the links made by different nodes, and there are also much fewer links than in a modular network, leading to the formation of truly disconnected components. (D) A compound topology combines a modular structure in

135 the network, with a nested structure within its modules, resulting in "Russian dolls trapped 136 inside filter bubbles", and tends to be larger than a modular network. 137 138 The first archetype, nested networks (quantified as nestedness), emerges when the links 139 made by species with fewer interactions represent a subset of those made by species 140 with more interactions, sequentially. The second archetype, modular networks 141 (quantified as modularity), represents systems that consist of cohesive subgroups of 142 species (i.e., modules), which are densely interconnected but have fewer interactions 143 with other subgroups. Gradient networks are the third and least studied archetype, 144 usually observed when sampling in steep environmental gradients, leading to few links 145 and almost no overlap in the interactions made by different species, therefore forming 146 several small components (i.e., totally disconnected subgroups). Finally, the fourth 147 archetype, compound networks, combines modularity in the network and nestedness 148 within its modules. 149 Lately, different research groups have independently proposed topologies similar to the 150 compound archetype, with slight variations in conceptualization and operationalization 151 ^{13–15}. Their structure would in general resemble a polycentric heterarchy ¹⁶, almost equivalent to a dissortative network ¹⁷. That is, in such a network there are differences 152 153 in influence between nodes, with one or a few nodes dominating each module and 154 playing the role of module hubs. 155 Despite the multiple options provided by the four archetypal topologies, another 156 seminal study published a little earlier proposed the universality of nestedness in mutualistic networks ¹⁸. By comparing food webs, pollination, and seed dispersal 157 158 systems, the study invoked co-evolutionary arguments to conclude that mutualisms 159 would form highly cohesive, nested networks in which a core of generalist species 160 would drive the dynamics and maybe the evolution of the entire local community. In 161 contrast, antagonisms would be characteristically modular, as traditionally assumed ¹⁹. 162 The almost overlooked gradient topology, which was not considered then and still remains poorly studied ²⁰, would be found only in the most specialized systems with 163 164 high interaction intimacy ²¹ or along steep gradients with substantial turnover in species

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166 Since nestedness was advanced, it became the dominant view in the study of interaction 167 networks, especially mutualistic ones. Most studies have been either trying to validate a 168 nested topology for the respective system of interest or aiming to link nestedness to the 169 stability of the entire community, sometimes resorting to circular reasoning ²³, despite how controversial this view has always been ²⁴. Nestedness has been pointed out as an 170 171 artifact ²⁵ or a simple byproduct of abundance ²⁶, and not a pattern derived from a 172 clearly defined process ²⁷. In addition, nestedness is also frequently found in 173 antagonisms ²⁸. A striking critique emerged from a study that revealed conceptual flaws in early nestedness indices and algorithms ²⁹. The high nestedness scores reported for 174 175 many systems were artificially inflated, and, as a more accurate index was proposed, it 176 became clear that nestedness was indeed highly variable and lower on average than 177 previously measured in many empirical networks ³⁰, and even lower than expected by chance from differences in abundance ³¹. 178 179 What should have been the final blow to the universality of nestedness came from a 180 naturalistic perspective. Different studies started to point out that the mutualistic 181 interactions sampled at a local scale and low completeness levels, included in popular databases, only represent fragments, not entire networks ^{32–34}. This deserves attention 182 183 considering that ecological functions, such as pollination, and its derived ecosystem 184 services, such as crop pollination, are not delivered by single taxa, but by guilds 185 composed of several taxa acting together at a local scale. 186 Therefore, variations in the definition of nestedness, conceptual limitations in its 187 proxies, and a narrow taxonomic focus have raised questions regarding its universality. 188 Nonetheless, nestedness remains a central concept in the study of mutualistic networks 189 and continues to provide a benchmark for assessing the structure of mutualistic interactions across pristine, degraded, and restored environments ³⁵. More recently, 190 191 however, studies started adopting conceptually refined approaches that offer fresh 192 perspectives on network topology ³⁶.

A way forward

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Given the challenges presented above, a new theoretical perspective was needed to make sense of patterns and processes. The Integrative Hypothesis of Specialization (IHS) was proposed as a way forward, originally presented as a graphical model ³⁷, and

197 subsequently mathematically formalized as an algorithmic model ³⁸. IHS was built as a 198 minimalistic mechanism that integrates sequential processes, which themselves 199 represent the mechanistic assembly rules of interaction networks. 200 In short, IHS proposes that interaction constraints of geographic, morphological, 201 physiological, or behavioral nature would lead to the formation of layers and modules in 202 interaction networks. Within these modules, the specialisation-niche breadth trade-off 203 predominates once the constraints are overcome. Consequently, we can deduce from 204 IHS that the assembly rules of interaction networks include a threshold of influence of 205 different ecological and evolutionary processes at a given hierarchical level (entire 206 network, layers, modules, motifs, and dyads), which determines which archetypal 207 topology predominates at that level. 208 To evaluate the four archetypal topologies with as much generality as possible, IHS 209 assumes that all interaction types can be viewed as pure consumer models, similarly to other network models ^{15,39,40}, but aiming at higher generality and parsimony. We know 210 211 that most interactions are more complex than this, but this simplification of reality has 212 proven to be surprisingly efficient in deducing topology. IHS is based on three first 213 principles that apply to evolving consumer-resource relationships in general 41: (i) each 214 type of resource has a set of characteristics that affect its use, and the types of resources 215 may be more or less similar to each other in these characteristics; (ii) a consumer's 216 mutation that increases its ability to use a given resource tends to improve its efficiency 217 on similar resources, but worsen its on different resources, generating clusters of usable 218 and non-usable resources; and (iii) a consumer's ability to use each resource in a given 219 cluster at a given time is the result of its previous adaptations and maladaptations (i.e. 220 mutations are small). Based on this, the free parameters of IHS are: (a) consumer 221 species richness, (b) resource species richness, (c) method used to generate the initial 222 innate performance matrix (innate method), (d) maximum dissimilarity between two 223 resource species (maximum dissimilarity), and (e) number of resource species clusters 224 (number of clusters). 225 The structure of IHS leads to the logical conclusion that the topology of a given 226 interaction network depends mainly on the dissimilarity between the resources 227 contained within it, from the point of view of consumer adaptations ⁴¹. In turn, this 228 conclusion leads to different secondary deductions, for example: the more dissimilar the 229 resources, say in traits, the more likely a modular topology emerges. Assuming that

most resources in nature tend to be organized in clusters ⁴², very large networks with 230 231 high heterogeneity would tend to have a compound topology: modular across clusters, 232 but nested within. In turn, small and homogeneous networks, which would actually 233 represent clusters, would tend to be nested. At the extreme of this continuum, small 234 networks with very high resource dissimilarity, probably resulting from very high 235 interaction intimacy or steep environmental gradients, would be organized as a gradient. 236 These deductions are valid mainly for interactions studied at the community level, 237 although further expansions of IHS to interactions between individuals have already 238 been made ⁴³. Although many studies propose that coevolution is an important driver of interaction networks 44, IHS assumes that their structure can emerge without mutual 239 adaptation between consumers and resources. This does not mean that coevolution is 240 241 not important for understanding species interactions in general. Rather, within the scope 242 of IHS, coevolution is not *necessary* to explain the assembly rules that generate the four 243 archetypal topologies. 244 A novel theory 245 Starting from the realization that IHS already has all the characteristics of a semantic theory ^{45–47}, all that remains is to formalize it as: the Integrative Theory of Interaction 246 247 Networks (or ITIN, for short). 248 To build a novel semantic theory of the assembly rules of interaction networks, it is 249 essential to understand the concepts and assumptions that guide this approach. Models 250 of this type are constructed theoretically and do not directly depend on real-world data, 251 which allows for a more flexible and adaptable approach to the conceptualization of the 252 expected patterns in natural phenomena. The development of these models involves the 253 application of a mathematical or logical formalism to deduce the models from the 254 axioms, allowing a meaningful comparison with the observed phenomena. This crucial 255 step has already been taken in building the algorithmic model used to make the proof of 256 concept of IHS 41.

Since its proof of concept, IHS has undergone a series of empirical tests of several

multiplicity of network topologies in real-world networks of pathogen infection ³⁷, parasitism ⁴⁸, seed dispersal ⁴⁹, pollination ⁵⁰, resin collection ⁵¹, burrow sharing ⁵²,

predictions deduced from its central axiom. IHS proved to be capable of explaining the

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predation ⁵³, resource partitioning ⁵⁴, multilayered interactions ⁵⁵, and food crop 261 262 management ⁵⁶, among many other examples, widely extending its original domain of 263 validity. 264 Later studies also demonstrated that compound and modular topologies are actually much more common than previously thought ¹⁴, as expected from theory. Most 265 266 interaction networks, in the end, resemble filter bubbles filled with Russian dolls, and a 267 good example of this kind of compound topology was observed in a system formed by small mammals and their ectoparasitic fleas ⁴⁸. So, if modular and compound topologies 268 269 are the most common structures observed in well-sampled, taxonomically 270 comprehensive interaction networks studied at different spatial scales, from local to 271 continental, what about nested and gradient topologies? Explaining the logical 272 relationship between archetypal topologies is the next step in the development of our 273 novel theory (Figure 3).

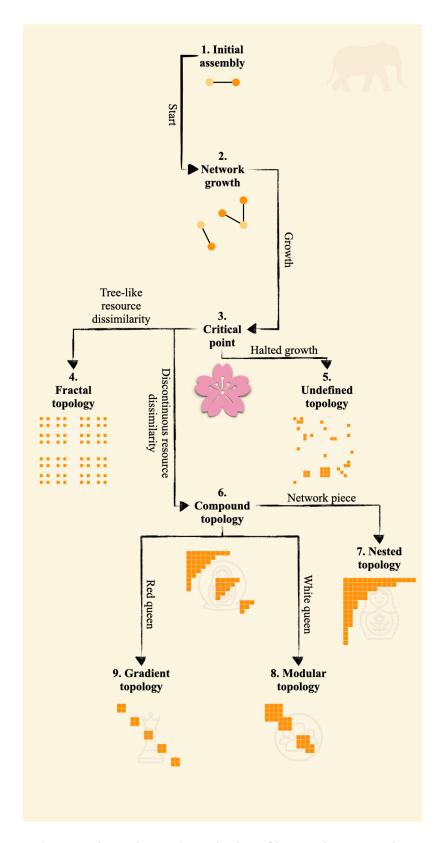


Figure 3: A roadmap to the archetypal topologies of interaction networks according to the Integrative Theory of Interaction Networks (ITIN). (1) As an example (see section Three archetypal stories), two species arrive at an open area after a disturbance, such as a forest fire, and begin to interact. (2) Ecological succession follows, bringing more

280 species and interactions, forming a shifting topology that may resemble a random 281 network. (3) A critical point is reached, a phase transition, after which the network may 282 or may not blossom, giving rise to different stable topologies. (4) When resource 283 dissimilarity follows a tree-like, hierarchical structure, the network becomes fractal, 284 resembling a Cantor matrix. (5) If network growth is halted by external factors, such as 285 hunting, its topology remains undefined. (6) When resource dissimilarity is 286 discontinuous and the network continues to grow, a compound topology emerges, with 287 modularity across scales and nestedness within modules. (7) Nested topologies, 288 therefore, represent the internal structure of modules, not the whole mature network. (8) 289 In mutualistic systems (White Queen Hypothesis), resources strengthen associations 290 with better consumers, leading to modular or compound topologies. (9) In antagonistic 291 systems (Red Queen Hypothesis), escape dynamics promote specialization, resulting in 292 gradient topologies. 293 294 Conceptual examples might help understand our rationale (Box 1). 295 Box 1: Three archetypal stories The Integrative Theory of Interaction Networks (ITIN) is a description of how real-296 297 world networks develop over time. The following "stories" exemplify possible paths 298 towards an observed network topology. 299 The niche sequence along a steep gradient story 300 Imagine several plant species from the same family, growing along a mountain's 301 altitudinal gradient. Evolved from the same ancestor, they still share most of their niche 302 characteristics, but differ, say, in frost tolerance. Each species has thus a position along 303 the growth-frost tolerance gradient. A set of herbivore species adapted to feeding on this 304 family will feed on all species. The network therefore will be fully connected. Over 305 time, slow-growing, high-altitude species will invest more in chemical defense, as each 306 herbivore attack leads to tissue loss, which they cannot readily compensate for by fast 307 growth given that they experience a short growing season. In response to the increased 308 chemical anti-herbivore defense, some herbivores will start avoiding these species, 309 while others will adapt and even specialize on them. The resulting topology will depend

310 on the degree of separation: if almost every herbivore becomes a specialist, we end up 311 with a modular network. If most herbivores exhibit some lower preference for the 312 species a bit higher up and lower down the gradient, we end up with a gradient 313 topology. 314 The abundance story Most species are rare ⁵⁷. The causes are multifaceted, from neutral ecological drift to 315 competitive hierarchies ⁵⁸ or statistical inevitability of random processes ⁵⁹. Either way, 316 317 the evolutionary pressures to evade an abundant predator are much stronger than those 318 to evade a rare one. Antelopes run (almost) fast enough to evade a cheetah, not only fast 319 enough to outrun the much slower jackal. As a consequence, all antelopes face 320 predation by common predators, and common antelopes face all predators, simply as a matter of probability. We can compute this as the matrix product of the abundance 321 322 vectors of antelopes and predators, normalised to sum to 1. The resulting network is 323 nested. Interestingly, two common predators may compete for common prey, leading to 324 specialisation for the most common prey only. This will be visible as anti-nestedness for 325 the most common prey, but still a strong nestedness signal overall. This makes 326 "nestedness" an awkward concept: the observed competition signal leads to a reduced 327 nestedness, while the overall topology is still highly nested. 328 *The introduced species story* 329 The effect of adding a new species to a network is likely to be extremely case-specific. 330 Imagine a pollination network to which a non-native plant species is added ⁶⁰. If that 331 species is similar in traits to species already present in the network, we can expect it to 332 "slot in", maintaining whatever topology existed before. However, if the plant's 333 attractive traits vastly exceed those of the existing species (e.g., flowers that are more 334 nectar rich), it may actually withdraw flower visitors from other plants. The network 335 topology is dependent on what happens next. If the pollinators keep their visitation 336 pattern, but prefer the new plant species, then the network becomes a bit more nested. 337 If, however, the new plant leads to a strong re-alignment of trait preferences, it may lead 338 to network fragmentation, and a more modular structure. The point here is that the 339 topology, in concert with the relevant traits, will in the end help us to understand what

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happened.

342 ITIN's mechanism sets the thresholds that define when one topology ends and another 343 begins. In the simplest and least realistic scenario, considering only a tree-like 344 distribution of resource dissimilarity, such as in a dendrogram, with progressive 345 dissimilarity between hierarchical levels, the fundamental topology of interaction 346 networks should be fractal, like in a Cantor matrix ⁶¹. Nevertheless, this should not be 347 expected in the real world, as virtually all trait distributions are discontinuous ⁴². 348 Considering even more realistic scenarios, sometimes network assembly is halted by a 349 process, such as overhunting, that prevents the addition of new nodes or the 350 establishment of links between them. In this case, the network might not reach an 351 archetypal topology, remaining frozen in an undefined transitional topology. 352 If the network continues to grow, it might develop to a point where it becomes 353 compound. First, because not all consumers are empirically able to use all resources ⁶², 354 this leads to the formation of layers and modules. Second, because after constraints are 355 met, there is more freedom of association within a module. Consequently, neutral processes leading to differences in abundance 63 or universal network assembly 356 357 processes such as preferential attachment ⁶⁴ should lead to a nested structure within the modules. Nested topologies, therefore, would represent pieces and not entire networks. 358 359 Two processes could further shape interaction networks into the remaining archetypal 360 topologies (Figure 4). On one hand, if the interaction type is antagonistic, according to 361 the Red Queen Hypothesis ⁶⁵ the resources would tend to escape the consumers, 362 reducing niche overlap and even the total number of links observed, maybe even 363 resulting in the extinction of some consumers, leading to a gradient topology. On the 364 other hand, if the interaction type that defines the links is mutualistic, then the resources 365 would tend to strengthen their association with the most advantageous consumers, while 366 escaping the worst consumers, which could be called a "White Queen Hypothesis". This 367 would lead to a truly modular topology, with smaller modules that are not internally nested, because freedom of association is still limited after the main constraints are 368 369 resolved.

The Integrative Theory of Interaction Networks Each type of resource has a set of characteristics that affect its use by each type of consumer, and the types of resources may be more or less similar to each other in these characteristics: A consumer's mutation that increases its ability to use a given resource tends to improve its use of similar resources, but worsen its use of different resources, generating clusters of usable and non-usable resources in different degrees; A consumer's ability to use each resource in a given cluster at a given time is the result of its previous adaptations and maladaptations; If resource dissimilarities are distributed discontinuously, there are clusters of resources and consumers in the network (v) If the interaction is mutualistic, associations between consumers and resources tend to be strengthened, but if the interaction is antagonistic, resources tend to escape the consumers. Free parameters: (a) consumer species richness (b) resource species richness: (c) method used to generate the initial innate performance matrix (innate (d) maximum dissimilarity between two resource species (maximum dissimilarity); (e) number of resource species clusters (number of clusters); (f) interaction type.

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Figure 4: A novel semantic theory of the assembly rules of interaction networks. As a development of the Integrative Hypothesis of Specialization (IHS), the Integrative Theory of Interaction Networks (ITIN) has been expanded by the addition of two first principles and one free parameter. This enhances its ability to explain which processes determine where one archetypal topology ends and another begins. See also Figure 3.

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Therefore, in addition to the first principles considered in the original IHS, ITIN also includes the following assumptions: (iv) if resource dissimilarities are distributed discontinuously, there are clusters of resources and consumers linked within the network; and (v) if the interaction is mutualistic, associations between consumers and resources tend to be strengthened, but if the interaction is antagonistic, resources tend to escape the consumers. Consequently, ITIN includes an additional free parameter: (f) interaction type.

From description to explanation

As discussed in the previous sections, the study of interaction networks has been guided by a multitude of metrics. Connectance, modularity, nestedness, specialization: each captures a piece of the puzzle. Yet, as in the parable of the blind monks and the elephant, these fragmented perspectives did not converge into the bigger picture. The result has been a proliferation of metrics that quantify patterns but do not explain

391 mechanistically how they arise or what they mean for ecological functions and 392 ecosystem services 8. 393 Without a mechanism that ties pattern to process, our ability to predict and intervene 394 remains limited. Conservation biologists struggle to anticipate how seed dispersal networks will respond to human disturbances ⁶⁶. Agroecologists seek to design 395 396 sustainable crops but are left with taxon-biased network fragments that overlook the 397 functional guilds actually delivering vital ecosystem services ⁶⁷. Epidemiologists face 398 difficulties estimating the risk of pathogen spillover without understanding the assembly 399 rules that govern host-pathogen networks ⁶⁸. The black-and-white paradigm of network topology has constrained both scientific inference and practical action, leaving us only 400 401 with numbers yet still in search for meaning. 402 ITIN provides this much needed meaning. By rooting network topologies in first 403 principles and free parameters mappable into operational variables, it shifts the focus 404 from static descriptors to dynamic assembly rules. It explains not only what structures 405 we observe, but why they emerge and where the thresholds lie between one topology 406 and another. 407

Practical applications

- 408 As an empirically tested semantic theory, ITIN might be used to help solve problems. 409 Not only theoretical puzzles, but also practical challenges within the frameworks
- provided by the UN's Sustainable Development Goals ⁶⁹ and the One Health Initiative 410
- ⁷⁰ (Figure 5). 411

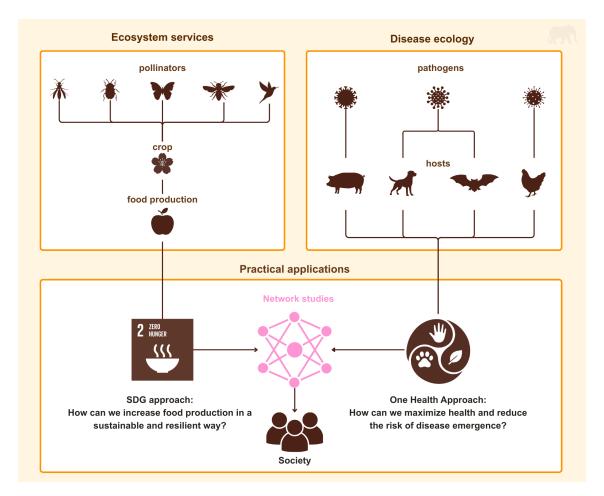


Figure 5: Conceptual framework illustrating how network studies can help translate knowledge about ecosystem services and diseases into solutions to real-world problems. On the left, pollinators contribute to ecosystem services such as crop pollination, supporting food production. On the right, animal hosts and pathogens represent the ecological interactions underlying disease emergence. By using network science, including the new framework provided by ITIN, both research areas can be aligned with the global agendas of Sustainable Development Goals and One Health.

The central driver of network topologies according to ITIN is resource dissimilarity. In order to develop practical solutions, it is of paramount importance to sample entire local networks responsible for a given ecosystem service of interest. A function-oriented modelling of interaction networks and the ecosystem services they deliver requires collaboration beyond taxonomic boundaries ⁹. For example, if the aim is to restore or improve pollination services delivered to a given crop, all functional groups (e.g., buzz pollinators and oil-collectors) that compose the local guild should be considered ⁷¹.

429 After replacing the black-and-white paradigm and focusing on functions and services, 430 another step would be to carefully consider what kind of phenomenon is actually being 431 represented as a network. In other words, what is flowing from node to node: matter, energy, or information ¹⁰? A network approach may help us better predict how 432 pathogens flow from one host to the other, leading to spillover or outbreaks 72. Such a 433 434 network approach could be guided by ITIN adapted to transient flows 73. This area of 435 application highlights the need to invest in initiatives targeted at biosurveillance and big data integration ⁷⁴. 436 437 There are critical interfaces across crops and diseases as well ⁶⁸, uniting the ecological and social layers of real-world networks 75. It is vital to better understand how the 438 439 relationships between humans, wildlife, crops, and livestock determine economic and 440 health gains and losses ⁷⁶. ITIN can be also helpful in this human-domestic-wildlife 441 interface, considering the huge resource dissimilarity observed in extensively changed 442 social-ecological systems ⁷⁷. 443 Moving from the entire network to its layers and modules, another practical application 444 of ITIN is the use of centrality metrics as proxies for the concept of keystone species 78. 445 Within this family of metrics, some have been pointed out as very useful not only for 446 identifying the most important species responsible for maintaining different ecosystem 447 services ⁷¹, but also to help select species to improve agroecosystems and crops in general ⁷⁹, and identify pathogen superspreaders ⁸⁰. Now it is also possible to assess the 448 449 topology of multilayer networks 81, as well as the multilayer centrality of a species in such a network 82, especially considering thresholds in resource dissimilarity that create 450 451 constraints for consumers between and within layers.

Testing for archetypal topologies

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Traditionally, ecologists have used different topological indices of avoidance ⁸³, connectance ⁸⁴, complementary specialization ⁸⁵, nestedness ⁸⁶, or modularity ⁸⁷, just to give a few examples, to measure the structure of empirical or synthetical networks.

Most commonly, the next step is to test whether the observed index is significantly different from expected by chance ³¹. However, neither a topological score nor a P-value alone can mechanistically explain the structure of empirical networks if the analysis is not guided by a hypothesis-oriented, strong-inference framework ⁸⁸.

460 Attempts to provide such a framework have been recently made in the case of 461 nestedness ⁸⁹ and compound topologies ⁴⁸. Similar hypothesis-driven efforts are still 462 lacking for the other two archetypal topologies, modular and gradient, as is an 463 information-theoretical way to compare the fits of these four archetypal topologies to 464 the data. Together, these advances and gaps signal that we need to work further on 465 conceptualization and operationalization to better understand network topologies and, most importantly, their assembly rules, so we can improve our mechanistic 466 467 understanding of interaction networks. 468 To help fill these gaps, we provide an analytical workflow built as an R tutorial to test 469 for network topologies, using the best solutions currently available (Supplementary 470 Code 1). Our workflow is based on five main axioms that create a framework to 471 interpret the results of each test. First, a network's topology is a matter of degree. 472 Second, no single metric is able to tell the whole story of a network's topology. Third, 473 null model analysis should be used to test hypothesis-derived predictions. Fourth, 474 networks can have mixed topologies. Fifth, weighted interaction networks have more 475 statistical power.

Concluding remarks

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477 Grounded in both logic and empirical evidence, the Integrative Theory of Interaction 478 Networks brings together a coherent set of mechanistic hypotheses from which testable 479 predictions can be drawn. It aims for generality while focusing on the assembly rules 480 that give rise to the four archetypal topologies observed in interaction networks. 481 Because it maps directly onto operational variables, it offers a practical framework for 482 studying real-world systems. Over time, our field is slowly moving from a black-and-483 white paradigm toward a more balanced view of network assembly, something akin to a 484 Buddhist "middle way." Like the elephant whose parts we have long examined, the 485 whole continues to reveal more than the sum of its parts. The moment calls for 486 widening our view, deepening our understanding of species interactions, and applying 487 that knowledge to the urgent challenges set by the Sustainable Development Goals. 488 There is still much to learn, and we can only move forward together.

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520 Author contributions

- MARM has since 2007 coordinated the research program that led to the novel theory.
- MARM, SES, and CFD discussed the final synthesis of the theory, wrote the first draft
- of the manuscript and the supplementary material, and coordinated the writing. CAK
- and RLM worked on the literature review and figures. MARM, CFD, and RLM worked
- on the supplement. All other authors contributed by discussing the ideas that form the
- 526 corpus of ITIN and reviewing drafts of the manuscript and supplement.

Competing interests

None to declare.

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Disclosure of generative AI-use

- We used ChatGPT (OpenAI, https://chatgpt.com) in combination with GitHub Copilot
- (GitHub, https://github.com/features/copilot) to optimize the code provided in the
- 532 supplement.

Supplementary materials

- Supplementary Code 1: We have prepared a repository on GitHub with an analytical
- workflow that includes data, code, and tutorials aimed at helping assess the topology of
- empirical and synthetical interaction networks, within the framework proposed by the
- 537 Integrative Theory of Interaction Networks:
- 538 https://github.com/marmello77/new-theory.

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