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1 Defining invasiveness and invasibility in ecological networks

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25 Abstract

The success and failure of a biological invasion is context based, and yet two key concepts – 26 the invasiveness of species and the invasibility of recipient ecosystems - are often defined and 27 considered separately. We propose a framework that can elucidate the complex relationship 28 between invasibility and invasiveness. It is based on trait-mediated interactions between 29 species and depicts the response of an ecological network to the intrusion of an alien species 30 drawing on the concept of community saturation. Here, invasiveness of an introduced species 31 with a particular trait is measured by its per capita population growth rate when the initial 32 propagule size of the introduced species is very small. The invasibility of the recipient habitat 33 or ecosystem is dependent on the structure of resident ecological network and is defined as 34 the total width of opportunity niche in the trait space susceptible to invasion. Invasibility is 35 thus a measure of network instability. We also correlate invasibility with the asymptotic 36 instability of resident ecological network, measured by the leading eigenvalue of the 37 interaction matrix that depicts trait-based interaction intensity multiplied by encounter rate (a 38 pairwise product of propagule sizes of all members in a community). We further examine the 39 40 relationship between invasibility and network architecture, including network connectance, nestedness and modularity. We exemplify this framework with a trait-based assembly model 41 42 under perturbations in ways to emulate fluctuating resources and random trait composition in ecological networks. The maximum invasiveness of a potential invader was found to be 43 positively correlated with invasibility of the recipient ecological network. Additionally, 44 ecosystems with high network modularity and high ecological stability tend to exhibit high 45 invasibility. Where quantitative data are lacking we propose using a qualitative interaction 46 matrix of the ecological network perceived by a potential invader so that the structural 47 network stability and invasibility can be estimated from literature or from expert opinion. This 48 approach links network structure, invasiveness and invasibility in the context of trait-mediated 49 interactions, such as the invasion of insects into mutualistic and antagonistic networks. 50

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52 Keywords: biological invasions, fluctuating resource hypothesis, invasiveness, invasibility,
53 ecological networks, interaction matrix, network stability, interaction strength

55 **1. Introduction**

The search for generality in invasion ecology has progressed largely through quantifying the 56 drivers behind two concepts separately and in concert: the invasiveness of alien species and 57 58 the invasibility of recipient ecosystems (Richardson and Pyšek 2006). The concept of invasiveness follows a species-centric view of the introduction-naturalisation-invasion 59 60 continuum (Blackburn et al. 2011). Much of the study of invasiveness has involved the identification and exploration of traits, barriers and drivers that determine location on this 61 continuum for a given taxon (Richardson and Pyšek 2012). Invasiveness, or the propensity of 62 invasive alien species (hereafter IAS) to invade, can be identified from comparative metrics 63 between invasive and non-invasive alien species, such as those related to translocation bias, 64 propagule pressure, and foraging/reproduction/dispersal traits (Pyšek and Richardson 2007). 65 Invasiveness is further related to the potential impacts of IAS on the function and service of 66 recipient ecosystems and thus dictates the prioritisation, prevention and control strategies in 67 response to biological invasions (Blackburn et al. 2014). Of particular importance are the 68 suite of traits of IAS that differ from those of native species and non-invasive alien species. A 69 trait-based priority list of potentially highly invasive species can then be developed (Peacock 70 and Worner 2008; Moravcová et al. 2015). 71

The second concept – invasibility – is a property of recipient ecosystems and involves 72 73 the elucidation of features that determine its vulnerability to invasion such a community diversity, composition and assembly (Lonsdale 1999). Community assembly rules outline 74 75 how species are "packed" in a community and how community composition is related to the occupied and available niche space in a given community. Early niche theories gave special 76 77 attention to the role of biotic interactions in structuring communities (Tilman 2004). More 78 recently, recognition that species assemblages in unsaturated local communities are at least in 79 part driven by neutral forcing via the continuous influx of regional and alien species (Hubbell 2001; Stohlgren et al. 2003). Despite contrasting opinions on the applicability of neutral 80 theory to real world communities (Chase 2005; Clark 2012; Rosindell et al. 2012), it is now 81 widely accepted that both deterministic and stochastic processes interact to structure species 82 assemblages (Bar-Massada et al. 2014; Nuwagaba et al. 2015). 83

To further elucidate the concept of invasibility, it is essential to first determine how an ecosystem responds to perturbations such as biological invasions, an issue that has been debated at least since May's (1974) proposition that complexity begets instability (McCann 2000; Fridley 2011, Allesina and Tang 2012). As the recipient ecosystem often comprises

many interacting species, an ecological network provides an effective model for exploring the 88 inherent complexity. A key aspect of this debate thus relates to connecting the different 89 perspectives of network architecture to the stability of ecological networks. The architecture 90 of an ecological network can be measured as particular features of the interaction matrix, 91 92 depicting whether and how strongly two nodes interact. Typical features include connectance (the proportion of realized interactions among all possible ones; Olesen and Jordano 2002), 93 nestedness (specialists only interacting with a subset of species with which generalists interact; 94 Bascompte et al. 2003) and compartmentalization (a network can be grouped into delimited 95 modules, measured by the level of modularity, where species are strongly interacting with 96 species within the same module but not those from other modules; Newman 2006). Network 97 complexity normally refers to a combined factor of network size (the number of nodes) and 98 connectance (May 1974; Allesina and Tang 2012). Although consensus on the structure of 99 100 mutualistic networks has been reached (e.g. Bascompte et al. 2003; Olesen et al. 2007; 101 Guimarães et al. 2007; Thébault and Fontaine 2010; Mello et al. 2011), there is still considerable debate with respect to antagonistic networks (e.g., on whether antagonistic 102 networks are more compartmentalised than random expectation; e.g. Poisot 2013). 103

Network stability, in contrast to network architecture, concerns how networks respond 104 to perturbations (Yodzis 1981) and can be measured using different approaches (i.e., 105 Lyapunov asymptotic stability, resilience, persistence and robustness, among others; May 106 1974; Pimm and Lawton 1978; Dunne et al. 2002; Donohue et al. 2013). The analysis of 107 Lyapunov stability is a long established mathematical tool in dynamical systems for depicting 108 whether a complex system will return to its local equilibrium after weak perturbations; this is 109 typically assessed as the leading eigenvalue of the interaction matrix (May 1974; Allesina and 110 Tang 2012). Such perturbations in an ecological network are often manifested as changes in 111 population sizes caused by stochasticity or changing resources (Davis et al. 2000). Recent 112 progress in resolving the complexity-stability debate has involved exploring the causal 113 relationship between the architecture and stability of many mutualistic (e.g., plant-frugivore 114 and plant-pollinator), trophic (food web) and antagonistic (predator-prey and host-115 116 parasite/pathogen) networks (e.g. Memmott et al. 2004; Eklof and Ebenman 2006; Bascompte et al. 2006; Burgos et al. 2007; Estrada 2007; Bastola et al. 2009; Kiers et al. 2010; Thébault 117 and Fontaine 2010; Brose 2011; de Visser et al. 2011; Stouffer and Bascompte 2011; James et 118 al. 2012), and explaining emergent network structures using dynamic network models with 119 120 adaptive and random species rewiring (van Baalen et al. 2001; Kondoh 2003; Rezende et al. 2007; Vacher et al. 2008; Valdovinos et al. 2010; Zhang et al. 2011; Suweis et al. 2013; 121

122 Minoarivelo et al. 2014; Nuwagaba et al. 2015).

Here, we introduce *invasibility* as a new aspect of network instability. Full 123 comprehension of the proposed framework of invasiveness and invasibility in ecological 124 networks requires us first to establish the concept of *community saturation* in a network. This 125 concept was initially developed from the theory on competition and limiting similarity 126 127 (MacArthur 1972; Abrams 1983) where strong interspecific interactions preclude the establishment of IAS. A saturated ecological network can be defined as a particular 128 community assemblage (a suite of species with their particular traits and population sizes) that 129 cannot be invaded by an alien species given low propagule number, irrespective of that 130 species' life history or relevant traits. Certainly, when the propagule size is too large, system 131 behaviour will be overridden by the influx of propagules, making the concept of invasibility 132 irrelevant. Very few, if any, ecological networks are truly saturated, as local communities can 133 be strongly affected by regional species dynamics and stochasticity (Abrams 1998; Loreau 134 2000); rather the concept of community saturation serves as a theoretical benchmark by which 135 136 invasibility can be measured. Specifically, the deviation from a saturated community can be measured by the cumulative niche space that permits invasion for a given recipient ecosystem. 137

138 Invasibility and invasiveness are not isolated concepts but are strongly interwoven. As an example, as Darwin (1859) first posited in what has become known as his "naturalization 139 140 hypothesis", introduced species should be more successful (i.e., more invasive) when the recipient community lacks congeneric or ecologically similar species (Duncan and Williams 141 2002). The underlying logic of this statement relies on the assumption that closely related 142 species show greater life history, trait and therefore niche overlap such that an intact 143 community would be minimally invadable to congeners relative to more distantly related 144 species. A counter-argument which has received some empirical support is that trait similarity 145 146 among related species might predict habitat suitability and result in higher invasibility when congeners are considered (Duncan and Williams 2002). Either way, it is clear that the two 147 core concepts – invasibility and invasiveness – are context dependent and closely related. 148 Moreover, introduced species can often only invade certain native ecosystems following some 149 form of perturbation (Davis et al. 2000; Shea and Chesson 2002). Here, we attempt to 150 151 explicitly bridge these two concepts through trait-mediated interactions in ecological networks by visualising both in a single invasion fitness diagram. Since this is a new aspect of 152 153 network instability, we also examine how invasibility is related to other network stability measures (specifically asymptotic stability), and how it is related to typical network structure. 154 155 We exemplify this framework using a trait-mediated assembly-level model and discuss how invasibility can be practically assessed with poor data quality by using qualitative interaction 156 157 matrix, in cases for selected invasive alien insects in a variety of ecosystems.

159 **2. Invasion fitness**

Assembly-level models have a long history in community ecology (e.g. Drake 1990; Morton 160 and Law 1997) and they normally assume infrequent colonization of new species from a pre-161 determined regional species pool. Some recent assembly-level models further allow limited 162 163 evolutionary processes (e.g. Drossel et al. 2001; McKane 2004) and adaptive response to disturbance (Kondoh 2003; Zhang et al. 2011; Suweis et al. 2013; Nuwagaba et al. 2015; 164 Minoarivelo and Hui 2015; Hui et al. 2015). In particular, the model proposed by Loeuille and 165 Loreau (2005) can depict the emergence of complex food webs through ecological and 166 evolutionary processes involving trait-mediated interactions. Here, we use a food-web model 167 with trait-mediated interactions to demonstrate the framework of invasiveness and invasibility. 168 Specifically, we consider a generalised version of the Loeuille and Loreau (2005) model 169 developed by Brännström et al. (2011) which depicts the per capita population change rate as 170 171 a function of population growth derived from prey consumption minus mortality from 172 senescence, predation and interference competition (Appendix A). This model is applied 173 because assembly-level data on functional traits, population sizes and interaction strengths are 174 often lacking, which precludes the study of a full network. Simply, targeting functional traits that appear especially important to the population demography of focal species (i.e., that 175 176 affect the strength of density dependence or influence biotic interactions) makes model parameterization more tractable. Certainly, the absence of such data hampers the clarity when 177 178 introducing the framework of network invasibility. For this reason, in what follows we will not discuss details of the model itself. Rather, we focus on introducing the framework with 179 this model only serving as a tool for generating required data. In the absence of 180 comprehensive knowledge of communities and community interactions, all is not lost – we 181 discuss the protocol for cases with insufficient data later. As we shall see, even only 182 characterizing the directionality of interactions among species without estimates of interaction 183 strength can be effective in quantifying invasiveness and invasibility (Rossberg et al. 2010). 184

Generating a resident network of multiple heterotrophic species as depicted in the model (Appendix A) can be accomplished in two ways. First, many studies have followed a simple procedure of randomly assigning trait values and parameters for all initial species, running the model until equilibrium is reached, and then removing those species with population sizes below a certain threshold (Holland and DeAngelis 2010). At this stage the network is considered to be at its equilibrium. Once the recipient community has reached its equilibrium, we could consider the invasiveness of a potential introduced species as its

invasion fitness, defined as its per-capita population growth rate when propagule size is trivial 192 (close to zero) and the community is at equilibrium (Fig.1a). Invasion fitness is a good proxy 193 of invasiveness for an introduced species – if the trait of an introduced species lies within the 194 green intervals along the zero invasion fitness line (Fig.1a), the introduced species will 195 196 experience positive invasion fitness and thus be able to establish and invade the resident community. If trait values land within the yellow intervals, the species will experience 197 negative invasion fitness and thus be repelled by the resident community (Fig.1a). Clearly, not 198 all species can invade the resident network (Fig.1a). 199

200 For a given introduced species with a particular trait, if there is a native resident species having an identical/similar trait (i.e. the trait of introduced species is close to any one 201 red dot [traits of resident species] in Fig.1a), the invasion fitness will then become close to 202 203 zero. Because of the zero population growth, such species are less likely to establish simply due to demographic stochasticity (the case of neutral coexistence). Even if these species 204 205 establish they will not become invasive but persist at low abundance until either eliminated via ecological drift or increasing opportunistically in response to disturbance. If the trait of an 206 207 alien species is quite different from those of any resident species (i.e. sitting between red dots in Fig.1a), it is then likely to become either highly invasive (peaks in green zones) or be 208 209 quickly expelled from the network (valleys in yellow zones), with a 50/50 chance for successful invasion in a species-rich network due to the constraints on any dynamic systems 210 given the continuity of the invasion fitness function (from the Fundamental Theorem of 211 Algebra and the Central Limit Theorem). To this end, the invasibility of the recipient 212 ecological network can be defined as the total width of opportunity niche in the trait space (i.e. 213 the summation of all the green intervals). 214

As an alternative to the static trait approach taken above we could also generate a 215 model community as an adaptive network, where species within the network can co-evolve 216 according to adaptive dynamics (Brännström et al. 2011), or where species with different 217 traits can be continuously introduced into the community from a large species pool (i.e. a 218 meta-community; Gilpin and Hanski 1991; Hubbell 2001). This approach will potentially, but 219 not always, lead to a saturated ecological network (Fig.1b). No alien species can invade a 220 saturated network as the invasion fitness of any introduced species is equal to or less than zero 221 (Fig.1b). These two ways of generating community assemblages sitting either at the 222 equilibrium of ecological dynamics (Fig.1a) or the saturated assembly (Fig.1b) provide an 223 update to the standard naturalisation hypothesis (Duncan and Williams, 2002). Of course, 224 225 even if the saturated assembly does exist, a community under constant bombardment of IAS

intrusion is not likely to be either on the equilibrium or remaining saturated but somewhere
between the two extremes. Consequently, we examine how temporal and trait perturbation
creates opportunity niche in ecological networks.

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230 3. Network invasibility under temporal perturbation

Many factors can drive the change and cyclic fluctuation of resources including land use 231 232 change, alteration of fire regimes, seasonality and weather conditions. In what follows, we introduce temporal perturbation to a saturated ecological network. Specifically, we examine 233 the fluctuating resource hypothesis (Davis et al. 2000), where fluctuation in resource 234 availability has been identified as the key factor mediating the susceptibility of an ecosystem 235 to invasion by non-resident species. We set a cyclic dynamics to the primary producers 236 (autotrophs in the model; Appendix A) using a sine function to modify resource levels with a 237 particular pulse and magnitude and record the temporal invasion fitness caused by such 238 periodic perturbation in the saturated ecological network. 239

Results confirmed that fluctuating resources can create an opportunity niche with 240 positive invasion fitness when resource levels exceed original levels (Fig.2). The temporal 241 invasiveness of an introduced alien species will reach the maximum when its trait value is 242 optimal for consuming autotrophic resource. The invasibility (length of positive invasion 243 fitness zone along trait axis in Fig.2) appeared high for the first half of the perturbation period 244 (positive phase) and allowed a wide range of introduced species to invade (a long tail towards 245 high trait/trophic direction) but quickly dropped to zero for the second half of the perturbation 246 247 period (negative phase). This suggests that although perturbation might not create a persistent opportunity niche even for highly invasive alien species in a saturated community, 248 environmental stochasticity can markedly increase the network invasibility in otherwise 249 250 resistant assemblages.

251 Importantly, rapid resource increase (approaching the 1.0 perturbation period in Fig.2), created an opportunity niche for invasion especially for species with optimal traits for 252 resource consumption, even though resource availability was still below the pre-perturbation 253 levels. Similarly, rapid resource decline (happening at 0.5 perturbation period in Fig.2) also 254 eliminated a portion of opportunity niche around the optimal trait. Evidently, the rate and 255 direction of change in resources as well their absolute level can both affect ecological 256 invasibility, with both playing similar roles in influencing network invasibility. This provides 257 an interesting extension to the fluctuating resource hypothesis which posits that variability in 258 resources promotes invasion in plants (Davis et al. 2000). It further echoes predictions of the 259

260 paradox of enrichment (Rosenzweig 1971) where enhanced resource level can be

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263 **4. Network structure and stability**

accompanied by instability in a food chain.

To simulate an unsaturated ecological network, we randomly altered the trait of heterotrophic 264 species in a saturated network, by an increment following a normal distribution with zero 265 mean and a particular standard deviation (in this case, $\sigma = 0.3$). Theoretically, we can then 266 calculate the Jacobian for randomly-altered networks, with $a_{ii} = \partial (dn_i/dt)/\partial n_i$, defined at the 267 population equilibriums. However, this is impossible in practice; instead, interaction strength 268 is often measured as the observed rate of species j interacting with species i, $a_{ij} = P_{ij}n_in_j$, 269 where P_{ij} measures the per-capita interaction strength. Consequently, we calculated three 270 network structures for 1000 randomly altered networks depicted by the matrix of observed 271 interaction strengths. As these altered networks were generated from the same saturated 272 assembly, they were of the same network size. 273

Metrics of network architecture are diverse. Essentially, these metrics descript 274 different aspects of network structure based on the matrix of interaction strength. These 275 metrics can be divided into two categories: those portraying the role of particular species in 276 the network (e.g. centrality) and those portraying the structure of entire networks. As 277 invasibility is related to the latter, we chose three metrics from this category that depict the 278 three most important features of a matrix – its matrix sparsity, asymmetry and symmetry. In 279 particular, these features are normally measured by three widely used metrics for quantitative 280 networks: connectance, nestedness and modularity. The quantitative connectance metric was 281 computed as the quantitative linkage density divided by the number of species in the network 282 (Tylianakis et al. 2007). A highly connected network is formed largely by generalists with 283 strong and evenly distributed interactions, whereas a less connected network is formed by 284 specialists. We used the metric WINE (weighted interaction nestedness estimator) to quantify 285 the level of nestedness (Galeano et al. 2009). Nested communities are often formed when both 286 specialist and generalist species are present, and where specialists primarily interact with a 287 subset of the partners of generalists. Finally, the level of modularity was measured using a 288 new algorithm QuanBimo (Dormann and Strauß 2014; adapted from Claused et al. 2008). A 289 290 community with high modularity is compartmentalised into multiple species modules or motifs, with species within the same module interacting strongly with each other but not with 291 species from other modules. All these network metric measurements were implemented in the 292 R library *bipartite* v2.05 (Dormann et al. 2008). 293

We examine how network metrics are associated with the invasibility (the total width 294 of opportunity niche) and the maximum invasiveness (height of the peak invasion fitness). 295 How the system is altered/deviated from its saturated assembly is depicted by the absolute 296 297 deviation of traits from the traits of the saturated assembly (hereafter, distance to assembly 298 saturation). The asymptotic instability of a system can be measured by the leading eigenvalue of its interaction matrix. When the leading eigenvalue is less than zero, the system will return 299 to its local equilibrium after small perturbations; otherwise, the fluctuations in population 300 abundance will be amplified. 301

302 As illustrated in Fig.3, there is a strong positive correlation between invasibility and maximum invasiveness (Pearson's r = 0.418, p < 0.001), suggesting an invasive species often 303 performing more invasively in highly invadable ecosystems. Invasibility showed a positive 304 305 correlation to the distance to assembly saturation (r = 0.106, p = 0.0005). In contrast, invasibility showed a negative correlation to ecological stability measured by the lead 306 307 eigenvalue (r = -0.267, p < 0.0001). Evidently, since distance to assembly saturation and asymptotic stability reflect different aspects of interaction networks, they have different 308 309 implications for understanding network function. Such an opposing relationship between the distance to assembly saturation and asymptotic stability is also evident in literature. For 310 311 instance, Allesina and Tang (2012) reconfirmed that asymptotic stability is negatively affected by nestedness in bipartite mutualistic networks, and as such at ecological time scales, 312 an ecosystem dominated by mutualistic interactions is likely unstable and species poor. In 313 contrast, mutualistic communities can maximise structural stability through potentially 314 enhanced nestedness (Rohr et al. 2014); that is, at long-term time scales, mutualistic 315 interactions can act as a stabilizing force and restrict diversification (Raimundo et al. 2014). 316 Invasibility also shows strong positive correlations (p < 0.0001) with all three measurements 317 of network architecture (with quantitative connectance, r = 0.266; with nestedness, r = 0.179; 318 with modularity, r = 0.324). When invasibility is unknown, we could predict it from 319 asymptotic stability and the three measurable network architectures. A generalized linear 320 model of the generated data showed that connectance and modularity are two strong 321 322 predictors, with predicted invasibility from only these two network structures showing a strong correlation with observed ones (r = 0.325, p < 0.0001). 323

Although these results only reflect non-causal correlations between network
architectures and invasibility, we could still contemplate the following ecological
explanations for these positive correlations. First, May (1974) devised a necessary condition,
further generalised by Allesina and Tang (2012), to ensure the stability in a complex network,

 $\alpha(SC)^{1/2} < \beta$, where α stands for the standard deviation of interaction strength, S species 328 richness, C connectance and β self-regulating force (e.g. negative density dependence). It 329 suggests that the stability of a network requires stronger self-regulating force (large β) than 330 potential reinforcing feedbacks from interspecific interactions, captured by the left side of the 331 332 inequality. A highly connected network (large C) could encompass more reinforcing feedbacks between species, violating this condition and thus rendering network instability. 333 Unstable networks, either due to reinforcing feedbacks or disturbance, could create 334 opportunity niches for invasion, thus augmenting invasibility. Second, a highly nested 335 network suggests a strong hierarchy and asymmetry, potentially from sorting species through 336 multiple ecological filters, with the most extreme specialists only interacting with the most 337 extreme generalists. This asymmetry could potentially create unbalanced energy/material flow 338 from specialists to generalists, creating opportunity niches for introduced specialists that can 339 exploit, perhaps more efficiently than resident generalists, the resident specialists. That is, 340 341 network asymmetry creates opportunity niches for specialists and thus enhances invasibility. However, as the correlation between nestedness and invasibility is weak, further 342 investigations are needed, especially for different types of ecological networks. Finally, a 343 highly compartmentalized network is formed by clearly bounded modules, with species 344 between modules rarely interacting. This suggests that these modules could have spatially or 345 temporally partitioned available niches and habitats. This nevertheless provides opportunities 346 for the invaders that can explore two or more modules. Species possessing traits with high 347 plasticity or tolerance, and those with complex life cycles (through ontogenetic niche shift), 348 could invade highly compartmentalized networks. 349

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351 **5. Invasibility assessment**

The above demonstration with the trait-based assembly model suggests that to be capable of 352 353 directly assessing the invasibility of a recipient ecosystem we need to rely on a comprehensive picture of functional traits that affect biotic interaction strengths and thus 354 population dynamics, as well as overall densities. In practice, however, sufficient data are 355 rarely available for estimating invasibility directly, even when global monitoring and web-356 based data sharing are actively seeking to make full records of species densities and traits in 357 prioritised ecosystems available. For a rapid assessment of invasibility, we need an indirect or 358 alternative way of capturing the interaction matrix. As shown above, an effective and 359 conceptually and logistically tractable approach is to construct the interaction matrix of the 360 recipient ecosystem (Fig.4a, top). In the absence of a quantitative matrix, a qualitative matrix 361

would suffice. Conservation agencies could work with local experts to compile a semi-362 quantitative interaction matrix (Fig.4a, middle) with weak interactions indicated by strength 363 0.1, intermediate interactions by 1, and strong interactions by 10. In cases where data were 364 even scarcer, a binary interaction matrix can be used (Fig.4a, bottom). While accurate 365 366 interaction strengths make enhance predictive power, qualitative matrices are largely sufficient to understand network behaviours. That said, incorrect designation of the 367 directionality of interactions (+, - or 0) may strongly bias the assessment of stability (Quirk 368 and Ruppert, 1965; May, 1973; Jeffries, 1974). Ideally, the matrix should reflect the full 369 species list of the recipient ecosystem. This can also be relaxed by only considering the 370 potential networks that an introduced species will likely to impact or interact with, referred to 371 here as an invader-centric ecological network. 372

373 Using an estimated interaction matrix at three different levels of acuity (quantitative, semi-quantitative and binary) we infer and compare invasibility from network architecture we 374 375 use the well-studied biocontrol agent cum invader, the Harlequin ladybird, Harmonia axyridis (Pallas) (Coleoptera: Coccinellidae) which is predicted to be a major threat to other species 376 377 within the aphidophagous guild (Roy et al. 2016). Native to Asia, H. axyridis has been introduced to many countries around the world as a biocontrol agent of aphids but it has 378 379 spread to countries in which it was not intentionally introduced. It is now widespread and abundant in many regions and many habitats (Roy et al. 2016). Here, we compare the 380 invasibility of two recipient habitats (agricultural and boreal systems in Europe; see Fig. 4b) 381 to the invasion of *H. axyridis*. Many studies have demonstrated the potential interactions 382 between *H. axyridis* and other aphidophagous species through laboratory and, to a lesser 383 extent, field studies (Pell et al. 2008; Hautier et al. 2011; Brown et al. 2015; Roy et al. 2016). 384 Harmonia axvridis engages in intraguild interactions with many species, including other 385 aphid predators such as other ladybirds (Ware et al. 2009), lacewings (Santi and Maini 2006), 386 and hoverflies (Ingels et al. 2015), as well as aphid parasitoids (Chacón et al. 2008) and 387 aphid-pathogenic fungi (Roy et al. 2008). In general these interactions favour H. axyridis, 388 though interaction strength and even direction can be influenced by the life stage of the 389 390 interacting species (Felix and Soares 2004) and the environment (Gardiner et al. 2009).

Based on literature and expert opinions, we compiled the semi-quantitative interaction matrices of the European agricultural and boreal systems that are currently being invaded by *H. axyridis* (Fig. 4c and d). The lead eigenvalue before the invasion (removing the entries related to *H. axyridis* in the matrix) is effectively zero for both the agricultural system and boreal forests (absolute value less than 10^{-17}), suggesting that both systems are at weak ecological equilibriums (asymptotically stable). After invasion by *H. axyridis*, both systems
become ecologically unstable, with the boreal forests more unstable than the agricultural
system (lead eigenvalue: 5.51 vs. 4.12), suggesting a stronger impact of *H. axyridis* on the
boreal forests from the perspective of stability.

To calculate the three network structures, we replaced all negative entries with zeros 400 as the calculation of these network metrics would otherwise report errors (note, for a 401 predation interaction between species *i* and *j*; if $a_{ii} > 0$, we could either record $a_{ii} < 0$ which is 402 the convention or $a_{ii} = 0$ which is the format for typical network analysis [used here]; 403 Newman 2010). Prior to the invasion of H. axyridis in our estimated networks, the 404 agricultural system has slightly higher connectance than the boreal system (0.38 vs. 0.34), as 405 well as low modularity (<0.01) and a high level of nestedness (>0.97). After the invasion, 406 407 connectance dropped slightly in both habitats (0.35 vs. 0.32) while modularity remained low (<0.01) and the nestedness high (~ 1) in both. 408

409 Comparing these results with general expectations from our models, specifically the 410 lead eigenvalue and connectance (Fig.3), the following interpretations can be proffered. First, before the invasion of *H. axyridis*, the agricultural system is more invadable than the boreal 411 412 forest. As such, the invasion of *H. axyridis* would be more likely to happen first in the agricultural system. Second, after the invasion of *H. axyridis*, invasibility of both systems was 413 414 reduced (reducing the risk of future invasion by similar invaders) although the agricultural system is still quite open for future invasions. The impact of the invasion of H. axyridis is 415 416 more strongly felt by the boreal forest with its stability more disturbed (experiencing greater 417 changes in the relative and absolute abundances of species after the arrival of H. axyridis). It is worth noting that the above exercise can be easily done before any invasions; a quick 418 picture of the invasibility and the potential impact of the invasion quickly drawn and the 419 invasibility between different habitats and ecosystems fairly compared. This can be 420 accomplished using expert opinions, though the quality of the predictions is of course 421 dependent on ecological realism in the interaction matrix. 422

423

424 **6.** Conclusions

We have demonstrated that ecological networks provide a good model for capturing the complexity of recipient ecosystems, and that the invasiveness of potential invaders and the invasibility of the recipient ecological networks can be defined using the concepts of invasion fitness and assembly saturation as a reference points. In this framework, invasiveness of a potential invader is defined as its invasion fitness (= per-capita population growth rate when 430 the propagule size is trivial) and the invasibility of the network is defined as the width of

431 opportunity niche available for potential invasions. This method is robust to the use of

- 432 simplified interaction matrices for rapid assessments of network invasibility. Importantly, we
- 433 argue that invasibility can be inferred from network stability, and that this is more closely
- 434 linked to assembly saturation than ecological stability. The trait-based approach allowed for
- the testing of the fluctuating resource hypothesis, thus emphasizing its heuristic value.
- 436

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659 Figure Legends:

Fig.1: Invasion fitness of an introduced species as a function of its trait value relative to the 660 trait values of the resident species in the ecological networks. Red dots indicate the trait 661 values of native resident species. (a) A randomly generated network at its ecological 662 equilibrium. Only introduced species with traits lie in the green intervals can establish and 663 invade the recipient ecological network; introduced species with traits lie in the yellow 664 intervals will be repelled by the network (i.e. wiped out from the resident species/ecosystem). 665 Invasibility is thus defined as the total width of the green intervals along the zero-fitness line. 666 (b) A saturated assembly is defined as the recipient network with zero invasibility. Parameter 667 values are the same as in Fig.1 of Brännström et al. (2011). 668

Fig.2: Invasion fitness of an introduced species as a function of its trait value in an ecological network. The resources of the saturated assembly in Fig.1b, n_0 , are disturbed by adding a periodic perturbation (sine form with pulse 10 and magnitude 100). An alien species with the trait and timing of introduction in the positive invasion fitness zone can invade the ecological network; otherwise it will be repelled from the network. The peak of positive invasion fitness corresponds to the first quarter of the period and the optimal trait for resource consumption (r $= \mu = 2$).

Fig.3: Relationships between invasibility, maximum invasiveness, distance to assembly
saturation (DAS) and the lead eigenvalue of the interaction matrix, as well as network
architectures (quantitative connectance, levels of nestedness and modularity) in 1000 random
ecological networks around the saturated assembly. Block tone corresponds to the frequency
of networks.

681 Fig.4: Interaction matrices in practice. (a) An illustration of quantitative, semi-quantitative

and qualitative interaction matrices (top, middle and bottom), with decreasing demands for

683 data quality. (b) A picture of one European agricultural system with boreal habitat in

background (Photo: H.E. Roy); both invaded by invasive alien ladybird *Harmonia axyridis*.

- 685 Panels (c) and (d) are semi-quantitative interaction matrices for agricultural and boreal
- 686 systems in Europe, based on literature and expert opinion. Acronyms: HA: Harmonia axyridis;
- 687 AB: Adalia bipunctata; CS: Coccinella septempunctata; CC: Chrysoperla carnea; EB:
- 688 Episyrphus balteatus; PN: Pandora neoaphidis; DC: Dinocampus coccinellae; AD: Adalia
- 689 *decempunctata*; HS: *Halyzia sedecimguttata*; PF: Phorid fly; AP: aphids.

Fig.1:





Fig.2:







(a)	spp1	spp2	spp3	spp4	spp5	5	(b)										
spp1	-0.96	-0.44	-0.11	0.58	-0.35	5												
spp2	0.33	0.58	0.65	-0.13	-0.85	5												
spp3	-0.23	-0.98	8 -0.24	-0.37	0.45													
spp4	0.64	-0.44	0.36	-0.01	-0.43	3												
spp5	-0.03	-0.46	0.40	-0.19	0.62													
r						-												
spp1	-10	-1	-0.1	1	-1													
spp2	0.1	1	1	-0.1	-10		Sec. and	42						-				
spp3	-0.1	-10	-0.1	-1	1				La Marca	-	No. all	. and	2				a unit	-
spp4	1	-1	1	-0.1	-1									.	alles.			
spp5	-0.1	-0.1	1	-0.1	1				Care S S	1.1	Reise Cont							
r						-		Para .	4 J A	Last of	1		-	The				
spp1	-1	-1	0	1	-1		A LINE				(and	-				-		X
spp2	0	1	1	0	-1					C.		and and						
spp3	0	-1	0	-1	1					No.		N.Y.						
spp4	1	-1	1	0	-1		1								20101-2	/ 148.66		
spp5	0	-1	1	0	1			a)	HA	AB	AD	HS	CC	EB	ΡN	DC	PF	AP
(c)								HA	-1	0	0	0	-0.1	0	0	-0.1	-0.1	10
(C)	HA	AB C	cs cc	EB	PN	DC	AP	AB	-10	-0.1	0	0	-0.1	0	0	-0.1	-1	10
HA	-1	0 -0	.1 -0.1	0	0 -0).1	10	AD	-10	0	-0.1	0	-0.1	0	0	-0.1	-1	10
AB	-10 -	0.1 -0	.1 -0.1	0	0 -0).1	10	HS	-10	0	0	0	-0.1	0	0	-0.1	0	0.1
CS	-0.1	0 ·	-1 -0.1	0	0	-1	10	CC	-10	0	0	0	-0.1	0	0	0	0	10
CC	-10	0	0 -0.1	0	0	0	10	EB	-10	0	0	0	0	0	0	0	0	10
EB	-10	0	0 -0.1	0	0	0	10	ΡN	0.1	0.1	0.1	0	0	0	0	0	0	10
ΡN	0.1	0.1 0	.1 0	0	0	0	10	DC	0	0	0	0	0	0	0	-0.1	0	0
DC	0	0	0 0	0	0 -0).1	0	PF	-0.1	0	0	0	-0.1	0	0	-0.1	0	0
AP	-10 ·	-10 -1	0 -10	-10	-10	0	0	AP	-10	-10	-10	0	-10	-10	-10	0	0	0

706 **Online Appendix:**

- 707 Appendix A: A trait-based assembly model.
- 708 We consider a generalised version of the Loeuille and Loreau (2005) model developed by
- 709 Brännström et al. (2011) which depicts the per capita population change rate as a function of
- population growth derived from prey consumption minus mortality from senescence,
- 711 predation and interference competition:

712
$$\frac{1}{n_i} \frac{dn_i}{dt} = \lambda \sum_{j=0}^{s} \frac{P_{ji}n_j}{\exp(\Delta r_{ij})} - \left(D_i + \sum_{j=1}^{s} P_{ij}n_j\right) - \sum_{j=1}^{s} C_{ij}n_j$$

- 713 where λ is the conversion efficiency for the consumptive interactions. The per-capita
- predation rate of species *i* preying on species *j* is set to be trait-mediated, $P_{ji} = \gamma_0 N(\Delta r_{ij} \mu, \sigma_{\gamma})$,
- where N(a, b) is the probability density function of normal distribution with mean *a* and
- standard deviation b, Δr_{ii} the trait difference of species i and j, r_i - r_j (considering the logarithm
- of body size relative to that of the autotroph). The coefficients μ and σ_{γ} represent the optimal
- trait ratio of predator to prey and the dietary breadth of the predator. The natural mortality is
- also assumed to be trait-mediated, $D_i = d_0 \exp(-r_i/4)$ (Peters, 1983). The intensity of
- interference competition is at its maximum when the two competing species have identical
- traits, $C_{ij} = k_0 N(\Delta r_{ij}, \sigma_k)$. In addition, the dynamics of the autotroph (resource base) is
- governed by, $dn_0/dt = n_0(g k_0n_0 \sum_{j=1}^s P_{0j}n_j)$, where g is the intrinsic population growth rate
- (see detail model description in Brännström et al. 2011). The resources n_0 are disturbed by
- adding a periodic perturbation (sine form with pulse 10 and magnitude 100).

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