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1	Species roles in plant-pollinator communities are conserved across native				
2	and alien ranges				
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#### 21 ABSTRACT

*Aim.* Alien species alter interaction networks by disrupting existing interactions, for example between plants and pollinators, and by engaging in new interactions. Predicting the effects of an incoming invader can be difficult, although recent work suggests species roles in interaction networks may be conserved across locations. We test whether species roles in plant-pollinator networks differ between their native and alien ranges, and whether the former can be used to predict the latter.

28 *Location*: worldwide.

29 Methods. We used 64 plant-pollinator networks to search for species occurring in at 30 least one network in its native range and one network in its alien range. We found 17 31 species meeting these criteria, distributed in 48 plant-pollinator networks. We characterized 32 each species' role by estimating species-level network indices: normalised degree, closeness 33 centrality, betweenness centrality, and two measures of contribution to modularity (c and z scores). Linear Mixed Models and Linear Regression Models were used to test for 34 differences in species role between native and alien ranges and to predict those roles from 35 the native to the alien range, respectively. 36

*Results.* Species roles varied considerably across species. Nevertheless, although species lost their native mutualists and gained novel interactions in the alien community, their role did not differ significantly between ranges. Consequently, closeness centrality and normalised degree in the alien range were highly predictable from the native range networks.

*Main conclusions.* Species with high degree and centrality define the core of nested networks. Our results suggest that core species are likely to establish interactions and be core species in the alien range, whilst species with few interactions in their native range will behave similarly in their alien range. Our results provide new insights into species role conservatism, and could help ecologists to predict alien species impact at the community level.

48 Key-words: biological invasions, centrality, conservatism, ecological networks, pollination,
49 predicting invasion

#### 50 INTRODUCTION

Predicting novel species interactions is a crucial challenge in today's rapidly changing world. 51 52 Alien species are an important driver of novel ecosystems (Hobbs et al., 2006) due to their 53 ability to outcompete native species (Chittka & Schurkens, 2001; Madjidian et al., 2008; Roy 54 et al., 2012), change the community structure (Albrecht & Gotelli, 2001; Memmott & Waser, 55 2002; Carpintero et al., 2005) and disrupt species interactions (Aizen et al., 2008; Traveset & Richardson, 2006; Tylianakis et al., 2008). Studies on alien species mostly focus on species 56 57 considered to be invasive, which means that rather little is known about those alien species 58 that remain at low population size or have fewer interactions with (and hence, impact on) the recipient community. 59

While many studies have tried to identify key features that predict which species will 60 61 become invasive and which communities are more likely to be invaded (Thuiller et al., 2005; 62 Richardson & Pysek, 2006; Pysek & Richardson, 2007) these remain of limited practical value. For example it remains difficult to predict whether a mutualistic interaction will 63 64 facilitate the establishment and dispersal of an alien species (Hulme, 2012). The limited 65 practical value of current work is partially due to the need for detailed information on each species involved in the potential novel interactions, which is usually very time consuming to 66 gather. Therefore, new methods to simplify predictions are required. An alternative could 67 68 be to assess the role a given species plays in the topology of interaction networks (e.g. 69 Stouffer et al. 2012; Martin Gonzalez et al., 2010; Albrecht et al. 2014). Species roles 70 summarize their ability to interact with, and potentially affect, other species in the 71 community in a way that is relatively easy to sample compared with measures of multiple

species and community traits. The application of species roles in ecological networks topredict invasion currently remains untested.

74 Ecological networks have been of considerable use when trying to understand how 75 alien species integrate into local communities (Memmott & Waser, 2002; Garcia et al., 2014, 76 Maruyama et al., 2016) and how they affect the overall mutualistic network structure 77 (Olesen et al., 2002a; Santos et al., 2012; Albrecht et al., 2014). In general, alien species are generalists, i.e. they interact with many species in the community in which they occur (Aizen 78 79 et al., 2008; Santos et al., 2012). Generalist species tend to occupy central positions in 80 ecological networks, and by interacting with other generalists and specialists (Memmott & Waser, 2002; Aizen et al., 2008) they contribute to the pattern of nestedness that 81 82 characterises many mutualistic networks (Bascompte, 2003; Bascompte & Jordano, 2007). 83 In addition to its number of direct interaction partners (termed 'degree'), a species' position allows it to connect different parts of the network and maintain network cohesiveness. This 84 85 helps to define its role in structuring the overall network topology (Martin Gonzalez et al., 2010), including elements of network structure such as clustering or modularity (Olesen et 86 al., 2007). Thus, the species' position in the network, i.e. its network role, captures key 87 information on its interactions with, and potential effects on, other species in the 88 89 community.

Recent work suggests that species roles are conserved across different locations. Species interactions, either generalist or specialist, have been shown to be phylogenetically conserved across space and time (Jordano *et al.*, 2003; Rezende *et al.*, 2007; Gómez *et al.*, 2010), because intrinsic (inherited) characteristics of species can constrain who can interact with whom (Eklöf *et al.*, 2013) and can be related to native and alien species roles in

95 network topology (Maruyama *et al.*, 2016). If these traits show low intraspecific variability 96 across locations, this indicates that species roles in networks should also be conserved. For 97 example, species roles in predator-prey networks can be conserved from an evolutionary 98 perspective, such that dynamically-important species in one network will be important in 99 the other networks in which it occurs (Stouffer *et al.*, 2012). Similarly, species roles in host-100 parasitoid networks were found to be intrinsic characteristics conserved over different 101 temporal and spatial scales (Baker *et al.*, 2015).

102 Despite evidence of an intrinsic component of species network roles, species 103 interactions and network roles may also be affected by local environmental and biotic conditions (Tylianakis et al., 2008; Trøjelsgaard et al., 2015). Moreover, the number and 104 type of interactions a species has increase with that species' abundance (e.g., Trøjelsgaard 105 106 et al., 2015), and species abundance and interactions may change during different stages of invasion (Aizen et al., 2008). Finally, patterns of non-random association among species 107 108 based on their phylogenetic relatedness (Rezende et al., 2007) suggest that coevolved interactions may be important for structuring mutualistic networks. Therefore, it is currently 109 not clear whether species roles can be extrapolated from one location to another that 110 111 differs in its evolutionary history and local community traits.

Here we aim to understand whether species roles differ and can be predicted from the native to the alien range of their distribution. Specifically, we use measures of plant and insect species roles in plant-pollinator networks (normalised degree, closeness and betweenness centrality, and *c* and *z* scores) recorded in both their native and alien ranges to test whether they differ consistently or can be predicted between ranges. Based on the findings that species roles and ecological interactions can be temporally, spatially and

phylogenetically conserved (Rezende *et al.*, 2007; Gómez *et al.*, 2010; Stouffer *et al.*, 2012; Baker *et al.*, 2015) we predict that a species' network role will be similar in its native and alien ranges, such that the former can be used to predict the latter. By including both specialist and generalist species we can draw conclusions about both rare and common alien species.

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124

#### 125 METHODS

126 We searched for plant-pollinator networks where we could potentially find species recorded 127 in both their native and alien range. We found 48 plant-pollinator networks of which 42 128 were downloaded from the "Web of Life" database (Ortega, 2014), three are our own data sampled in New Zealand and three are unpublished data from Lopezaraiza-Mikel and 129 130 Memmott in Hawaii; Table S1). Our criteria of species/network inclusion in the dataset was 131 to have a target species occurring in at least one network as native and one network as 132 alien. Thus each network can contain more than one target species, each of which may be 133 either in its native or its alien range. As some of these networks contain only the 134 presence/absence of interactions and the sampling effort of these networks is mostly unknown, we analysed all networks as binary matrices. In addition, here a flower visitor was 135 considered to be a pollinator, irrespective of whether effective pollination was 136 demonstrated. To define species range as native or alien, we used the following online 137 information: Global Invasive Species Database (http://www.issg.org/database/welcome/), 138 139 Global Invasive Species Information Network (http://www.gisin.org), Delivering Alien

Invasive Species Inventories for Europe (http://www.europe-aliens.org/), GB Non-Native
Species Secretariat Website (http://www.nonnativespecies.org), Plant Pest Information
Network of New Zealand (http://archive.mpi.govt.nz/applications/ppin), Centre for Invasive
Species and Ecosystem Health (http://www.bugwood.org/), Weeds in Australia
(http://www.environment.gov.au/biodiversity/invasive/weeds/), and Invasive Species of
Japan (https://www.nies.go.jp ).

146

# 147 Species roles

Species roles in networks can be described by a variety of different, yet often correlated metrics. Our intent here was not to provide an exhaustive comparison of different potential measures of species roles, or to determine which metrics were best conserved and why. Rather, we focused on testing a 'proof of concept' that roles could be conserved, so we focused on five complementary metrics that could potentially capture different aspects of species ecology:

154 1) Normalised degree – the number of interactions per species (i.e. degree) divided 155 by the number of possible interacting partners, which controls for differences in network 156 size. Normalised degree is the most local centrality index that characterizes a species' 157 network position, such that species with high degree are core in the network structure and 158 enhance robustness (Solé & Montoya, 2001; Dunne *et al.*, 2002). Additionally, normalised 159 degree estimates how generalist/specialist a species is relative to other species in the same 160 trophic level of the community in which it occurs.

2) Closeness centrality (hereafter, closeness) – the average distance (path length) to 161 all other species in the network. Closeness incorporates the number of immediate 162 163 connections to adjacent nodes and the connections of those nodes, so is a more global 164 measure of location than degree. In bipartite networks, closeness and betweenness are measured for the unipartite projection of each trophic level based on shared interaction 165 partners, such that higher closeness indicates a greater number of interaction partners 166 167 shared with other species in the same trophic level that also share partners with many other species (Freeman, 1979; Martín Gonzalez et al., 2010). Thus, closeness is a measure of niche 168 169 overlap with other species at the same trophic level via shared pollinators and the potential 170 for either positive or negative indirect effects via short path lengths (Morales & Traveset, 2008; Carvalheiro et al., 2014). 171

3) Betweenness centrality (hereafter, betweenness) – the proportion of the shortest paths linking any pair of species in the network that cross through a given species. It estimates species importance for network cohesiveness (Freeman, 1979; Martín Gonzalez *et al.*, 2010). Species with high betweenness can potentially connect different parts of the network that could be otherwise sparsely linked or even isolated; thus alien species that tend to be highly generalist may be linking previously isolated species in plant-pollinator networks and affect the overall network structure.

4) and 5) *c* and *z* scores: the combination of these two metrics describes a species' role in the topology of the network as a hub, peripheral or connector within and among modules (Olesen *et al.*, 2007) based on the modularity of the network (Guimera & Amaral, 2005). The *z*–score calculates the standardized number of links a species has within a module, and the *c*–score calculates the among module connectivity, which is the number of

links a given species establishes among different modules. Therefore, high values of c and z 184 are related to generalist species that have many interactions throughout the whole 185 186 network, either as hubs connecting species within modules, or as connectors linking 187 different modules. On the other hand, low values of c and z describe peripheral species that tend to be specialists. Alien plant species that invade a new range may act as network hubs 188 by attracting many different pollinator species through providing high amounts of nectar, 189 190 for example, Himalayan balsam (Impatiens glandulifera Royle) acts as a "magnet species" in its alien range (Chittka & Schurkens, 2001, Lopezaraiza-Mikel et al. 2007), whilst alien 191 192 pollinator species may act as network connectors while searching for floral resources in 193 different modules.

To allow comparisons across networks with different size, closeness and betweenness were each scaled to sum to 1. Species role metrics were calculated using bipartite (Dormann *et al.*, 2009) and rnetcarto packages (Doulcier, 2015) for R; correlations among these metrics are shown in Table S5.

198

199 Statistical analysis

#### 200 Are there differences in species roles in their native vs. alien range?

To answer whether species roles differed from native to alien ranges we used Linear Mixed-Effects Models (LMMs) in the Ime4 package (Bates *et al.*, 2014). Individual models were fitted for normalised degree, closeness, betweenness, and *c*- and *z*-scores. The first four metrics were logit transformed to solve the issue of being bounded from zero to one (Warton & Hui, 2011). Range (native vs. alien) was modelled as a fixed factor, whilst

network and species were fitted as random effects to account for multiple observations 206 from the same network and to group native and alien measures from the same species. 207 208 Residual plots were used to check model adherence to assumptions. The overall variance 209 explained by the model, and the proportion that could be attributed to the fixed factor (range) and the random factors were estimated by calculating: i) conditional Pseudo R-210 squared (R<sup>2</sup>GLMM<sub>(fix+rand)</sub>), to estimate total variance explained by the fixed and random 211 212 effects combined, ii) marginal Pseudo R-squared (R<sup>2</sup>GLMM<sub>(fix)</sub>), to estimate the variance 213 explained by range, and iii) the difference between the two  $(R^2GLMM_{(fix+rand)} - R^2GLMM_{(fix)})$ to estimate the contribution of the random effects only  $(R^2GLMM_{(rand)})$  (Nakagawa & 214 215 Schielzeth, 2013), using the MuMIm package (Barton, 2013). Then, to determine if any difference in species roles between native and exotic range could have occurred due to 216 217 biogeographical patterns from tropical to temperate zones (Olesen & Jordano, 2002; 218 Schleuning et al., 2012), we re-ran the above models including the absolute latitude as a 219 fixed effect interacting with range. Likewise, we re-ran the models with trophic level (plant or pollinator) and its interaction with range to determine whether any differences between 220 221 native and alien range only applied to one trophic level.

222

#### 223 Does a species' role in the native range predict its role in the alien range?

To test whether a species' role in the native range can predict its role in the alien range, we fitted five linear regressions relating species' mean normalised degree, closeness, betweenness, and the *c*- and *z*-scores in the alien range to the mean values in their native range. Normalised degree was strongly influenced by an outlier, which was removed and consequently improved model fit (Appendix S1). Model validation to check for 11

homoscedasticity and normality of the residuals was performed following Crawley (2013) 229 and Zuur et al. (2009). As previously, we re-ran these regressions including, separately, 230 231 absolute latitude and trophic level and their interactions with species' role in the native 232 range to determine whether the predictive power depended on these variables. Latitude was determined for each species as the absolute difference between latitudinal mean in the 233 native range and the latitudinal mean in the alien range. The latitudinal mean was obtained 234 235 by averaging the absolute latitude of all occurrences each species has in its native and alien 236 ranges.

237 Subsequently, we jack-knifed the linear regression models to provide an unbiased assessment of how accurately species roles could be predicted in alien networks based on 238 their mean role in the native networks (Efron, 1983). Each species was removed from the 239 240 linear regression in turn, the regression re-fitted, and predictions of the role metrics were generated for that species in the alien networks based on its mean value across its native 241 242 networks. The observed mean values in the alien range were then compared against the predicted values using Pearson's correlations. Individual species roles and mean species 243 roles were tested for correlation (presented as the Spearman coefficient in Table S5) and a 244 245 Bonferroni correction was used in both LMMs and LMs. All statistical and network analyses 246 were run in R v. 2.15.3 and v. 3.1.1 (R Core Team, 2014).

247

248

249 **RESULTS** 

We compiled information on 12 plant species and five pollinator species that occurred in at least one network in a native range and one network in an alien range (Table 1). These 17 species, from 19 different countries, were distributed in all continents except Antarctica (Fig. 1, Table S1); this translates into a large range of different habitats, climatic conditions and species richness. In total, we worked with 167 occurrences of the 17 target species (i.e. one occurrence corresponds to the occurrence of a species in either its native or alien range; note that multiple target species can occur in the same network) (Table S2).

257

#### 258 Are there differences in species roles in their native and alien range?

259 There was no significant difference between native and alien ranges in any of the measures 260 of species' role (Table 2). In other words we found no evidence that, for example, species consistently interact in a more generalist way in their exotic vs. native range. Rather, the 261 variance explained by the models was primarily attributable to the random factors 262 263 (R<sup>2</sup>GLMM<sub>(rand)</sub> was 94%, 40%, and 20% in the closeness, normalised degree and betweenness models respectively), which were the network and the species identity, whilst 264 265 range, the fixed term, was not statistically significant for any of the metrics tested (Table 2). 266 Similarly, the random structure explained around one third of the variance in the z-score (29%) and the *c*-score models (37%). The large variance retained by the random structure 267 suggests that species differ considerably in their network roles and that, unsurprisingly, 268 269 species roles depend on the local network (e.g., network size constrains the range of possible roles), and this large variance within native or exotic ranges of a species blurred any 270 significant differences between them. 271

Even though network architecture can change across regions (Olesen & Jordano, 272 2002), we found no systematic change in species roles with latitude, neither significant 273 274 range x latitude interaction (Table S3). However, a significant range x trophic level 275 interaction for closeness (Table S3) revealed that the native range had lower closeness for pollinators but not for plants. This indicates that pollinators may move into a more central 276 role in their alien range by pollinating generalist plants that are also pollinated by many 277 278 other species and share those pollinators with many other plants. Given that in our analyses 279 there were more plant species than pollinator species, this interaction effect captured the 280 difference between ranges for pollinators that was otherwise masked by the lack of 281 difference on plant species. Moreover, pollinator species had higher *c*-scores than plant species independently of range, suggesting that the pollinators included in our analyses may 282 be better network connectors (Table S3). In fact, most plant and pollinator species played 283 peripheral roles in our networks (73%) but pollinators were the main connectors (88%), 284 285 module hubs (75%) and the only network hubs (100%) (Table S4).

286

# 287 Does a species' role in the native range predict its role in the alien range?

Two measures of species roles, closeness and normalised degree, in the alien range could be predicted from the native range data ( $F_{1,15} = 27.32$ , p = 0.0001,  $r^2 = 0.62$  and  $F_{1,14} = 13.56$ , p= 0.0025,  $r^2 = 0.46$ , respectively; Fig. 2). The coefficients for closeness and normalised degree were 0.98 (SE ± 0.187) and 0.71 (SE ± 0.192), respectively, and both had intercepts that did not differ significantly from zero (closeness: t = 0.25, p = 0.809; normalised degree: t = 0.67, p = 0.512), suggesting that a species' role in the native range is associated to that in the alien range. In contrast, the positive trend in the relationship between native and alien 14 295 range when estimating betweenness (slope = 0.208 SE ± 0.109) and the z-score (slope = 0.412  $\pm$  0.204) was marginally non-significant (F<sub>1,15</sub> = 3.63, p = 0.076, r<sup>2</sup> = 0.14 and F<sub>1,15</sub> = 296 4.07, p = 0.062,  $r^2 = 0.16$ , respectively; Fig. 2) and lacked any significance for the *c*-score 297 298 model ( $F_{1,15} = 0.22$ , p = 0.649). Although the testing of correlated variables (Table S5) 299 increases the probability of type I error, the effects for closeness and normalised degree remained significant when a Bonferroni correction was applied (corrected alpha = 0.01). 300 301 Moreover, out of five variables tested, the probability of finding two significant at an alpha 302 below 0.0025 is extremely low (6.2 x 10<sup>-5</sup>, calculated using the Bernoulli process described in 303 Moran 2003), indicating that overall the suite of species roles in the exotic range could be 304 predicted better from roles in the native range than would be expected by chance.

305 The predictive effects of closeness and normalised degree were consistent when 306 latitude and trophic level were included in the models (Table S4). Neither latitude (normalised degree:  $F_{3,13} = 0.355$ , p = 0.787; closeness:  $F_{3,13} = 1.61$ , p = 0.235; betweenness: 307 308  $F_{3,13} = 0.938$ , p = 0.450; c-score:  $F_{3,14} = 2.00$ , p = 0.173; z-score:  $F_{3,14} = 0.56$ , p = 0.652) or trophic level (normalised degree:  $F_{3,13} = 0.262$ , p = 0.851; closeness:  $F_{3,13} = 1.708$ , p = 0.214; 309 betweenness: F<sub>3,13</sub> = 1.044, p = 0.406; *c*-score: F<sub>3,14</sub> = 2.00, p = 0.173; *z*-score: F<sub>3,14</sub> = 0.56, p = 310 311 0.652) showed any significant interaction with range when tested for predictive effects of species roles from the native to the alien range of a species distribution (Table S4). 312 Congruent with the LMM results, after model selection we detected that the mean c-score 313 314 was also higher for pollinators than for plants independently of range ( $F_{2,14}$  = 12.02, p = 315 0.0009).

316 In the jack-knife validation of our predictions, predicted values of closeness in the 317 alien range were highly correlated with the corresponding observed values (t = 15.339, p <

318 0.0001, r = 0.777), suggesting that the species closeness in the native range is a good 319 predictor of the species closeness in the alien range. The predictive power of native range 320 was lower but still a good predictor for more than half of the species when estimating 321 normalised degree (t = 9.040, p < 0.0001, r = 0.583), z-score (t = 8.0445 p = < 0.0001, r= 322 0.53), and c-score (t = 8.587, p < 0.001, r = 0.56), though not as good for betweenness (t = 323 5.621, p < 0.0001, r = 0.401).

324

325

#### 326 **DISCUSSION**

327 Two consistent patterns emerged from our analyses of the 48 datasets: 1) although species 328 differed considerably in their roles, the roles of species generally did not differ consistently between their alien and native ranges, and 2) two metrics of species roles, closeness and 329 330 normalised degree, in the alien range could be predicted from the native range. Betweenness and z-score predictions from the native to the alien range were marginally 331 332 non-significant, but showed a trend toward positive correlation, which was unsurprising in 333 the case of betweenness, given its high correlation with normalised degree and closeness 334 (Table S5b). Despite this overall predictive ability, we found that pollinators (but not plants) 335 had a higher closeness in their alien range, probably due to their ability to exploit a wide range of resources and thus interact with generalist plants. Still, trophic level (pollinator vs. 336 337 plants) did not interact significantly with range, except for *c*-score, which showed higher 338 values for pollinators, suggesting they may play a better role in connecting the whole 339 networks than did plants. Our results suggest that species role conservatism may occur,

340 such that species that are generalists or play a central role in their native network are likely341 to play a similar role in their alien range.

342

343 Limitations

344 In an ideal situation, the networks studied would have been collected using the same methods, aiming for quantitative data collected over similar periods of time. The dataset 345 used comes from different sources that used different sampling methodologies, spatial and 346 347 temporal scales. Moreover, it contains only species that successfully established in the alien range thus it lacks information for those species that failed to establish in the alien range. 348 349 Moreover, our models do not consider species abundance, which is known to drive some 350 network patterns (Blüthgen et al., 2007; Dorado et al., 2011; Staniczenko et al., 2013; Fort et al., 2016) as well as the effects of invasive species (Dostal et al., 2013; Carvalheiro et al., 351 352 2014; Traveset & Richardson, 2014). Furthermore, the conservation status of the areas from 353 which the networks were sampled is mostly unknown. Thus, the native range should not be 354 necessarily interpreted as a pristine environment given that we are likely working with 355 altered environments in both ranges. This high heterogeneity in the dataset generated high 356 variance across different networks (even within a species' native or alien range), which would have reduced the probability of detecting differences across 'treatments'. In that 357 358 sense, the absence of evidence for differences in species roles in native vs. alien range 359 cannot be viewed as evidence of absence. That said, the positive correlations we observed between native- and alien-range values of closeness and normalised degree were robust 360 361 enough to be seen despite the data being averaged across these heterogeneous replicate 362 networks and spanning species with a range of roles from specialists to generalists.

363

# 364 The intrinsic roles of alien species in pollination networks

365 The correlation between species roles in their native and alien range in the five network 366 statistics concurs with other authors who report that species have intrinsic properties in ecological networks that persist over temporal and spatial scales (Jordano et al., 2003; 367 Gómez et al., 2010; Stouffer et al., 2012; Baker et al., 2015). From the roles estimated here, 368 369 high degree and high closeness define the core of the nested network (i.e. those generalists that interact with both specialists and generalists), and our results suggest that core species 370 371 will tend to maintain this role even when they enter novel communities. Species with high 372 degree, i.e. generalists, are expected to be good invaders because they can increase their 373 chance to establish and spread through the population by interacting with many of the 374 "available" species. Conversely, specialist species with few interactions in the native range 375 will also have only few interactions in the alien range, and this may lower their chance of 376 establishing into the novel community if, for example, the resource is scarce and 377 competition strong (Aizen et al., 2008; Aizen et al., 2012), as shown in previous work that simulated invasion of food webs (Romanuk et al., 2009). In turn, high closeness can be seen 378 in species that interact with other central species in the community, even if the focal species 379 380 is not a generalist itself. In fact, in our dataset the average normalised degree and average 381 closeness were not significantly correlated (r = 0.24, Table S5b), such that a species could 382 occupy a consistently central position in networks by interacting with central species, rather 383 than by being a generalist itself. Therefore, the combination of degree and closeness can potentially be good indicators of species with high risk of introduction success in terms of 384 invasion. On the other hand, the poor prediction of betweenness and the c- and z-score, 385

386 which indicate the role a species plays as connecting different parts of the network, 387 suggests that the role of species as connectors may depend on the distribution of species 388 into modules.

Most plant species depend on animal species for pollination (Waser & Ollerton, 389 390 2006; Ollerton et al., 2011), thereby any characteristic that enhances interactions with 391 pollinators would likely be favourable when colonizing a new area. Central alien plants may have an advantage in the new range in terms of gene flow if local pollinators show high 392 393 fidelity. A greater number of pollinator species constantly visiting different conspecific 394 flowers may promote greater deposition of conspecific pollen grains, therefore increasing pollination (Brosi & Briggs 2013; Huang et al., 2015). Nevertheless, the benefits of this 395 increased visitation frequency may be partly offset by an increase in heterospecfic pollen 396 397 transport (Fang & Huang 2013) if, instead, the alien plant interacts with a generalist pollinator that visits different plant species therefore increasing heterospecific pollen 398 399 transfer, potentially reducing seed set (Ashman & Arceo-Gómez, 2013). Still, heterospecific pollen transfer has been shown to be generally low and have none, low or species-specific 400 effect on plant reproduction (Bartomeus *et al.*, 2008; Montgomery & Rathcke, 2012; Fang & 401 402 Huang, 2013; Emer et al., 2015). Moreover, central pollinator species may have an advantage over less connected species when arriving in an alien community due to their 403 ability to visit different flower species, thereby obtaining different food resources (Traveset 404 405 et al, 2013). Pollinators were the main connectors in our networks and that was more 406 frequent in their alien range. Given that the main pollinator connectors in our network were social insects (i.e Apis mellifera and Bombus spp.), which are usually highly abundant in 407 408 invaded areas (e.g. Aizen et al., 2008; Santos et al., 2012), and whose foraging individuals

409 reflect the colony needs (Willmer & Finlayson 2014 and references therein), it may be that 410 these species' roles vary according to their population density and foraging behaviour. Yet, 411 central pollinator species may face high competition with the local pollinators with which 412 they share interactions, a constraint that may make it difficult for pollinators to establish in 413 a novel community with low nectar/pollen resources, for example.

414 Our findings also have implications for network persistence. Rewiring, i.e. the reshuffling of interaction links among species, can enhance network resilience and 415 416 robustness to disturbance (Staniczenko et al., 2010; Kaiser-Bunbury et al., 2011; Olesen et 417 al., 2011). Given that both plant and pollinator links can be transferred from native generalist to alien generalist species (Aizen et al., 2008), and that the probability of a native 418 pollinator interacting with an alien plant increases with its degree and nestedness 419 420 contribution (Stouffer et al., 2014), the introduction of a highly generalist alien species may 421 affect not only the local generalist species but also the more specialized ones that connect 422 to it via interaction rewiring (Aizen et al., 2008). The consequences of this will depend on the centrality of the introduced species in combination with that of the native species, e.g. 423 highly-connected alien species will likely promote local species rewiring, whilst the arrival of 424 425 a poorly-connected species (i.e. a specialist) may have a mild or even neutral effect on local species interactions. Moreover, a species that remains in its home range in which the 426 community has changed due to local extinctions and alien species invasion will find itself in 427 428 a novel network of interactions. Given that species roles are conserved, rewiring of 429 interactions will be needed for the local species to fit into the novel community (Gilljam et al., 2015). 430

431

#### 432 Conclusions

In summary, there seems to be an intrinsic component of species roles in plant-pollinator 433 434 networks that is conserved across species native and alien ranges. Our results suggest that 435 the core network position that a species occupies when introduced in a novel community 436 will resemble how generalist or specialist it is in its native community. Our results provide 437 new insights into the recent literature about interactions and species role conservatism, and have implications regarding the potential links that alien species may be able to create or 438 439 disrupt once introduced into novel communities. Further studies incorporating community 440 traits and the phylogenetic relationship between species with species network roles will advance our understanding of how alien species interact with, and potentially drive the 441 formation of, novel communities. 442

443

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# 677 SUPPORTING INFORMATION

678 Additional Supporting Information can be found in the online version of this article:

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680 **APPENDIX S1.** Outlier detection analyses.

TABLE S1 – Description of the networks used for the analyses of the species' roles of plants
and pollinators in the alien and native range.

TABLE S2. List of the target species and the networks in which they were recorded. Network
ID follows Figure 1 and Table S1 in which details of each network are provided.

TABLE S3. Results of the Linear Mixed-Effect Models (LMMs) and the Linear Regression
Models (LMs) testing whether latitude and trophic level interact with species range to
determine species' roles.

688**TABLE S4.** Species roles on pollination networks following Olesen et al. (2007): Peripheral  $z \le$ 6892.5,  $c \le 0.62$ ; Connector  $z \le 2.5$ , c > 0.62; Module hub z > 2.5,  $c \le 0.62$ ; Network hub z > 2.5

**TABLE S5.** Correlation between normalised degree, closeness, betweenness, *c* and *z* scores measured with (a) individual entries, i.e. the value of the role of each species in each network is taking into account, as used in the Linear Mixed Models, and (b) when the averages for each species are considered, as used in the Linear Regressions of the manuscript. Values correspond to the Spearman correlation coefficient  $\rho$ .

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# 699 BIOSKETCHES

Carine Emer is a community ecologist interested on understanding how anthropogenic
 disturbance affect animal-plant interactions. Her research includes both mutualistic and

antagonistic processes in tropical and temperate habitats. Recently she has studied the
effects of invasive species, habitat loss and fragmentation on ecological networks. She is
currently a postdoctoral researcher at the Universidade Estadual Paulista (UNESP) in Brazil.
The authors are part of a collaboration established during her doctorate at the University of
Bristol, UK.

Authors contributions: CE and JMT developed the study framework. CE gathered the data, ran the analyses, and wrote the manuscript. IPV provided statistical advice. DM contributed with the study design and discussion. JM advised on the collection of the field data, and JM and JMT commented and edited the versions of the manuscript.

#### 712 TABLES AND FIGURES LEGEND

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**Table 1**. The 17 plant and pollinator species analysed in this study (see Table S1 for further
information about each network).

**Table 2.** Results of the Linear Mixed-Effects Models (LMMs) testing whether species roles differ from the native to the alien range. Pseudo R-squared values were calculated to estimate the variance explained by the fixed and random structure of each model:  $R^{2}_{\text{fix+rand}}$ estimates total variance explained by the fixed and random effects combined;  $R^{2}_{\text{fix}}$  estimates the variance explained by range;  $R^{2}_{\text{rand}}$  estimates the contribution of the random effects only.

**Figure 1.** The location of the 48 plant-pollinator networks. Panels A-G show the location of those networks that overlap in the full map. Numbers are the individual codes of each network identity (see Supplementary Material).

**Figure 2**. Results of the linear regression models testing whether a species' role in the native range predicts its role in the alien range. (a) Normalised degree; (b) Closeness; (c) Betweenness; (d) *c*-score; and (e) *z*-score. Results of normalised degree are shown after the removal of an outlier.

		Number of networks present		
Plant species	Family	Native networks	Alien networks	
Achillea millefolium L.	Asteraceae	4	5	
Cirsium arvense (L.) Scop	Asteraceae	3	6	
Cytisus scoparius (L.) Link	Fabaceae	1	1	
Eupatorium cannabinum L.	Asteraceae	1	2	
Hieracium pillosela L.	Asteraceae	2	4	
Hypochaeris radicata L.	Asteraceae	5	6	
Leucanthemum vulgare Lam.	Asteraceae	2	4	
Lotus corniculatus L.	Fabaceae	3	1	
Taraxacum officinale F.H. Wigg	Asteraceae	4	1	
Trifolium pratense L.	Fabaceae	2	4	
Trifolium repens L.	Fabaceae	3	10	
Verbascum thapsus L.	Scrophulariaceae	2	3	
Total plants` occurrences		31	47	
Insect species	Order			
Apis mellifera L.	Hymenoptera	9	28	
Bombus hortorum L.	Hymenoptera	7	4	
Bombus terrestris L.	Hymenoptera	9	6	
Eristalis tenax L.	Diptera	5	11	
Pieris rapae L.	Lepidoptera	3	6	
Total insects` occurrences		33	46	
Total		64	102	

# 730 Table 2

	Linear Mixed-Effects Models					
	Est	t	р	$R^2_{\text{ fix-rand}}$	$R^2_{fix}$	$R^2_{rand}$
Normalised degree	0.305	1.227	0.226	0.408	0.011	0.397
Closeness	-0.108	-1.188	0.237	0.939	0.003	0.936
Betweenness	0.116	0.326	0.747	0.201	0.000	0.201
z – score	-0.029	-0.158	0.875	0.285	0.000	0.285
<i>c</i> - score	0.028	1.076	0.285	0.378	0.010	0.377



