

Four priority areas to advance invasion science in the face of rapid environmental change

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1 Four priority areas to advance invasion science in the face of rapid

2 environmental change

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56 Abstract

57 Unprecedented rates of introduction and spread of non-native species pose burgeoning 58 challenges to biodiversity, natural resource management, regional economies, and human health. Current biosecurity efforts are failing to keep pace with globalization, revealing critical gaps in 59 our understanding and response to invasions. Here, we identify four priority areas to advance 60 61 invasion science in the face of rapid global environmental change. First, invasion science should strive to develop a more comprehensive framework for predicting how the behavior, abundance, 62 and interspecific interactions of non-native species vary in relation to conditions in receiving 63 environments and how these factors govern the ecological impacts of invasion. A second priority 64 65 is to understand the potential synergistic effects of multiple co-occurring stressors – particularly involving climate change – on the establishment and impact of non-native species. Climate 66 adaptation and mitigation strategies will need to consider the possible consequences of 67 68 promoting non-native species, and appropriate management responses to non-native species will need to be developed. The third priority is to address the taxonomic impediment. The ability to 69 70 detect and evaluate invasion risks is compromised by a growing deficit in taxonomic expertise, 71 which cannot be adequately compensated by new molecular technologies alone. Management of 72 biosecurity risks will become increasingly challenging unless academia, industry, and governments train and employ new personnel in taxonomy and systematics. Fourth, we 73 recommend that internationally cooperative biosecurity strategies consider the bridgehead effects 74 75 of global dispersal networks, in which organisms tend to invade new regions from locations 76 where they have already established. Cooperation among countries to eradicate or control species established in bridgehead regions should yield greater benefit than independent attempts by 77 individual countries to exclude these species from arriving and establishing. 78

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Key words: biosecurity; climate change; ecological impact; invasive species; management; risk
assessment

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83 Introduction

84 Invasion science – the systematic investigation of the causes and consequences of biological invasions – is a rapidly evolving interdisciplinary field. Its explosive growth over the 85 past few decades mirrors societal concern over the upsurge in the global rate of invasions 86 (Seebens et al. 2017; Pyšek et al. 2020; Seebens et al. 2020) and reflects the fundamental and 87 88 applied importance of understanding how species spread into new regions, why some ecosystems are more vulnerable to invasions, and what factors govern the impacts of non-native species. To 89 date, research addressing these issues has yielded valuable insights into the forces that structure 90 ecological communities, the relationship between diversity and stability, mechanisms of 91 adaptation and rapid evolution, causes of extinction and biotic homogenization, and the 92 93 connectedness between socioeconomic and ecological systems, among other phenomena (Lockwood et al. 2013; Hui and Richardson 2019). More remains to be done to sharpen and 94 integrate these insights into predictive frameworks. In addition, pressure is increasing for 95 invasion science to adapt to emerging issues such as rapid advances in biotechnology, 96 accelerating global change, expanding transportation networks, abrupt landscape 97 98 transformations, and infectious disease emergence (Ricciardi et al. 2017; Nuñez et al. 2020). 99 Invasion science is a relatively young discipline (Ricciardi and MacIsaac 2008) that has embraced diverse domains in ecology and cognate fields (e.g., population biology, biogeography, 100 evolutionary biology, paleoecology, physiology) and has formed linkages with disciplines related 101

to biosecurity – such as epidemiology, risk analysis, resource economics, and vector science

- 103 (Vaz et al. 2017). This multidisciplinary expansion reflects the increasing complexity of
- biological invasions and their impacts (Richardson 2011; Pyšek et al. 2020).
- 105 Here, we consider how invasion science should adapt to the Anthropocene an era of
- 106 burgeoning human influence, novel stressors, and rapid environmental change (Steffen et al.
- 107 2015; Waters et al. 2016). We are an international team of ecologists, with diverse and extensive
- 108 experience in biological invasions in many parts of the world. Our team gathered in September
- 109 2018 to consider emerging scientific, technological, and sociological issues which, if addressed,
- should ensure that invasion science can more successfully contend with rapid global change.
- 111 Through consensus (see Supplemental Material), we arrived at four overarching issues, relevant
- to a broad range of taxa, environments, and geographic regions, and which encompass some of
- the most important challenges facing our field today (Figure 1).
- 114

115 1. Predicting ecological impacts of invasions under rapid environmental change

- 116 **1.1.** The need for greater predictive power: Major advances and ongoing challenges
- 117 *1.1.1. Environmental context-dependency of impacts*

While invasion science has made substantial progress in understanding how non-native species arrive in new locations and establish self-sustaining populations (Catford et al. 2009; Jeschke and Heger 2018), it has been less successful in forecasting when and where such species will substantially affect their recipient environments (Ricciardi et al. 2013; Simberloff et al. 2013; Kumschick et al. 2015). Non-native species can affect ecological, economic, cultural, and human health in diverse ways (Jeschke et al. 2014; Shackleton et al. 2018), but in this section we focus on ecological impacts. Here, 'impact' is defined broadly as a measurable change to the

125	environment attributable directly or indirectly to the presence of a non-native species (Ricciardi
126	et al. 2013), and includes their effects on individual performance, population size and
127	composition of ecological communities of native species, which in some cases may be
128	irreversible (IUCN 2020).
129	Impact prediction is a long-standing, complex challenge. While rates of non-native
130	species introductions are increasing across regions (Seebens et al. 2017, 2020), impacts have
131	been recorded for only a small fraction of these species and the sites they invade (Ruiz et al.
132	1999; Ricciardi and Kipp 2008; Vilà et al. 2011; Hulme et al. 2013; Simberloff et al. 2013;
133	Evans et al. 2018b). It is generally assumed that most invasions have negligible environmental
134	consequences (Williamson and Fitter 1996), whereas a small proportion has significant and
135	sometimes enormous effects – an inverse magnitude-frequency distribution similar to that
136	associated with natural disasters (Ricciardi et al. 2011). However, uncertainty exists concerning
137	which cases truly reflect an absence of impact rather than a lack of study (Latombe et al. 2019).
138	Even well-known impacts exhibit substantial variation over time and space; invaders may remain
139	innocuous for years or even decades prior to becoming disruptive when, for example,
140	environmental change triggers a new impact (Crooks 2005; Coutts et al. 2018). The impacts of
141	any given invader can vary greatly among ecosystems (Strayer 2020) and across environmental
142	gradients within ecosystems (Kestrup and Ricciardi 2009; Stritar et al. 2010; Hulme et al. 2013;
143	Sapsford et al. 2020). Context-dependencies of invasion – that is, interactions among propagule
144	pressure, the traits of the invader, the composition of the recipient community, and the
145	physicochemical environment – have hardly been addressed by any formal body of theory, but
146	some overarching frameworks are now being explored (e.g., Cronin et al. 2015; Iacarella et al.
147	2015a; Dickey et al. 2020; Sapsford et al. 2020).

Coupled with the challenge of context-dependency is the sheer complexity of mechanisms by which non-native species can interact with their environment (Ricciardi et al. 2013; Kumschick et al. 2015). Synergistic interactions, nonlinearities, time lags, threshold effects, regime shifts, and indirect effects of non-native species are difficult to predict (Ricciardi et al. 2013; Essl et al. 2015b; Kumschick et al. 2015; Aagaard and Lockwood 2016; Hui and Richardson 2017; Strayer et al. 2017). Consequently, accurate risk assessment tools for sound management decisions are still lacking.

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156 *1.1.2. Temporal variation and time lags of impacts*

Factors affecting temporal variation in impact remain a major research gap, in large part 157 because of the vast majority of impact studies are conducted over very short time scales (Strayer 158 et al. 2006; Stricker et al. 2015). Time-since-invasion has been found to be an important correlate 159 of the ecological impacts of non-native species (Iacarella et al. 2015b; Evans et al. 2018a; 160 Zavorka et al. 2018), but time lags between establishment and peak impact have thus far evaded 161 prediction and are increasingly recognized as hindering risk assessment (e.g., Coutts et al. 2018). 162 Predictions of spatiotemporal variation in impact direction and magnitude could be improved 163 164 through experimental and theoretical investigations of the relationship between an invader's per*capita* effect and its abundance (Yokomizo et al. 2009; Cronin et al. 2015; Sofaer et al. 2018; 165 166 Bradley et al. 2019; Strayer 2020). We must also consider the influence of spatial scale on *per* 167 capita effects or impacts measured in small plots and mesocosms; attempts to extrapolate these effects up to landscape scales relevant to management (e.g., by calculating the product of the per 168 169 capita effect, local abundance, and range size of an invader) might not adequately capture 170 changes to biodiversity, biotic interactions, and ecosystem function, and thus might

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171	underestimate some large-scale consequences of invasion (Hawkins et al. 2015; Bernard-Verdier
172	and Hulme 2019; but see Dick et al. 2017b). Greater effort is required to test factors that mediate
173	indirect and multi-scale effects, particularly where an invader's impact is transmitted across a
174	suite of interacting species (Feit et al. 2018).
175	Conservation interventions and ecosystem management must contend with significant
176	time lags between the onset of the environmental stressors and the expression of invader impacts,
177	and forecasting such phenomena is plagued by context dependencies and non-linearities (Essl et
178	al. 2015b, c; Coutts et al. 2018). An understudied issue is how to recognize and manage the
179	interactive and cumulative effects of time lags in ecological responses to invasion. Delayed
180	biodiversity responses (e.g., dominance shifts, species turnover, metapopulation dynamics,
181	extinction debt) to anthropogenic stressors such as invasion can lead to abrupt shifts in
182	ecosystem functioning (Essl et al. 2015b) and underestimation of rates of contemporary
183	biodiversity change (Essl et al. 2015c). Given the management implications of this phenomenon,
184	ecological responses to compounded and cumulative stressors are becoming an increasing focus
185	of theory, experiments, and time series analyses (Foster et al. 2016; Candolin et al. 2018;
186	Kleinman et al. 2019; Shinoda and Akasaka 2020).

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1.1.3. Impacts on ecosystem processes 188

Demand is growing for reliable assessments and predictions of the ecosystem-level 189 impacts of non-native species, especially those impacts that affect the provision of ecosystem 190 services in rapidly changing environments (Vilà and Hulme 2017). This need reflects the larger 191 challenge of understanding how ecosystem function is altered by the combined effects of species 192 gains (invasion, range expansion) and losses (extinction, range contraction), which are 193

simultaneously consequences and drivers of global change. With few exceptions (e.g., Mascaro 194 et al. 2012; Kuebbing et al. 2015), work on how these two forces affect ecosystem functioning 195 has developed largely in isolation (Wardle et al. 2011). Owing to this disconnect, ecologists are 196 unable to predict over the coming decades the net ecosystem consequence of these two opposing 197 forces – specifically, whether or not species that are gained at local scales through invasion will 198 199 affect ecosystem process rates in a comparable way to those native species that are lost. Moreover, despite the many ecosystem impacts revealed thus far (Ehrenfeld 2010; Vilà et al. 200 2011; Simberloff et al. 2013), few types of ecosystems and invaders have been studied relative to 201 202 those that exist (Crystal-Ornelas and Lockwood 2020). It is likely that an enormous number of non-native species have affected individual performance, population sizes, and community 203 structure, though direct and indirect effects on native species (e.g. via competition, herbivory, 204 predation, hybridization, and as diseases or their vectors), or by changing the physical, chemical 205 or structural characteristics of the environment (Blackburn et al. 2014; IUCN 2020), in ways that 206 207 have not been documented (Carlton 2009; Simberloff 2011). Ecosystem-level impacts must remain a major focus, with researchers taking advantage of available technological tools (e.g., 208 209 Asner et al. 2008). Further, research on how biodiversity loss affects ecosystem functioning must 210 be evaluated alongside effects of non-native species additions, to better understand how humandriven species change will affect ecosystem processes across scales. For example, given that 211 212 community composition can influence biosphere-atmosphere exchange of greenhouse gases 213 (Metcalfe et al. 2011), how non-native species influence processes that underpin this exchange 214 relative to native species extirpations can have significant, currently unrecognized consequences for climate change. 215

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217 **1.2.** New and future challenges

218 1.2.1. Impacts of interventions for restoring ecosystem function

Co-occurring environmental stressors are increasing pressures to use non-native species 219 for restoring ecosystem functions eroded by native species loss (Mascaro et al. 2012; Castro-220 Díez et al. 2019). The notion of restoring ecosystems that have lost important species by 221 222 substituting non-native species to perform key functions traces back at least to the 1980s (Atkinson 1988) and has seen growing interest in recent years (Seddon et al. 2014a; Galetti et al. 223 2017; Pires 2017). Of particular interest are proposals and ongoing projects to establish species 224 225 to replace seed dispersers of plant species that have lost their ancestral native mutualisms (Seddon et al. 2014a; Galetti et al. 2017), and large herbivores and carnivores to fulfill lost 226 trophic linkages (Svenning et al. 2016). These efforts are often listed under the rubric of 227 rewilding (Lorimer et al. 2015; Svenning et al. 2016). Calls for active rewilding to restore 228 ecological processes (Perino et al. 2019) have primarily focused on the reintroduction of native 229 species, but some practitioners have advocated a 'flexible' approach to restoration that entails 230 using non-native species (Ewel and Putz 2004; but see Sotka and Byers 2019) as well as the 231 reintroduction of species into parts of their native range from which they have been absent for 232 various lengths of time. 233

As with translocation to accommodate climate change (see section 2.2.3), proposals for translocations to restore ecosystem functions (e.g., IUCN 2013; Aslan et al. 2014) have been the subject of substantial discussion of potential risks and benefits (Nogués-Bravo et al. 2016; Rubenstein and Rubenstein 2016; Fernández et al. 2017; Pettorelli et al. 2018; Perino et al. 2019). Lunt et al. (2013) have compared possible risks and benefits of translocations to restore ecosystem functions and translocations to address climate change, pointing to the possibility of

240	addressing both goals simultaneously. To employ proposed decision tools and adhere to the
241	International Union for Conservation of Nature (IUCN) guidelines, both advocates and critics
242	increasingly agree that progress is required on more accurate risk assessments and on
243	characterization, categorization, and quantification of the environmental impacts of
244	translocations (Jeschke et al. 2014), as has occurred with the EICAT framework (Blackburn et al.
245	2014; Hawkins et al. 2015; Evans et al. 2016), which has been adopted as an IUCN standard
246	(IUCN 2020), and similarly for socioeconomic impacts, as has begun under the SEICAT
247	framework (Bacher et al. 2018).
248	Conversely, other efforts to conserve native species or restore ecosystems involve non-
249	native species eradication. Such interventions should be preceeded by a predictive risk
250	assessment of the indirect effects of invader removal (Bergstrom et al. 2009; Caut et al. 2009;
251	Ruscoe et al. 2011; Lindenmayer et al. 2017) and the legacy effects of invasion (Corbin and
252	D'Antonio 2012; Grove et al. 2015; Reynolds et al. 2017; Pickett et al. 2019). Eradication has
253	had demonstrable benefits to biodiversity (Baider and Florens 2011; Monks et al. 2014; Jones et
254	al. 2016), but targeting the removal of a single invasive species within an ecosystem that
255	contains several non-native species can be counterproductive. A predictive framework must
256	consider the topology of species interactions, both trophic and non-trophic, to determine when
257	single-species management may lead to unintended consequences (Glen et al. 2013; Ballari et al.
258	2016; Hui and Richardson 2019).
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260 *1.2.2. Burgeoning novel organisms*

Escalating risks are associated with the intentional and unintentional release of novel organisms (those with no analogue in the natural environment) through biotechnological

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advances that create transgenic or genetically engineered organisms. For example, some 263 proposals for rewilding entail *de-extinction* – i.e., creation of various sorts of proxies of extinct 264 species for release to the wild. Versions of de-extinction are expected to become increasingly 265 feasible (Stokstad 2015; Shapiro 2017). The process involves either backbreeding (Stokstad 266 2015) or the reconstruction of the genome of an extinct species from recovered strands of DNA, 267 268 which can then be used either to modify or to replace the genome of a suitable living relative or to genetically engineer embryos that can be implanted in a compatible host. Some 269 conservationists will advocate for such proxy species to be reintroduced to a suitable former 270 271 geographic environment (Seddon et al. 2014b), and perceived ecosystem management benefits may arise from doing so (Church 2013). Environmental differences between contemporary and 272 historic habitats (Peers et al. 2016) might encourage further genetic manipulation to create better 273 adapted species. Depending on the length of time the proxy species has been extinct and the 274 method used to produce the proxy, introducing such entities to the wild is tantamount to 275 introducing a non-native species (IUCN 2013; IUCN/SSC 2016; Genovesi and Simberloff 2020), 276 an action that in the absence of predictive knowledge increases the likelihood of unintended 277 ecological consequences. 278

Advances in biotechnology will also facilitate the creation of self-replicating synthetic cells designed for novel tasks such as contaminant remediation, carbon sequestration, and the production of biofuels (Menetrez 2012; Azad et al. 2014; Singh et al. 2016; Dvorak et al. 2017). As synthetic and transgenic organisms will contain combinations of ecological traits that are unlikely to be encountered naturally, recipient communities will be evolutionarily naïve to these organisms and could be predisposed to being altered by them (Saul and Jeschke 2015). Such impacts could be subtle but far-reaching, as has been demonstrated for macroscopic transgenic

species (Post and Parry 2011; Vacher et al. 2011; Oke et al. 2013). Among the larger risks is the 286 capacity for such organisms to evolve in the wild and to exchange genes with other organisms 287 (Dana et al. 2012). Given the exponential growth of molecular technology, the rate of 288 development of such organisms could outpace progress in developing effective risk assessments 289 of their ecological effects. This issue emphasizes a need for greater integration of evolutionary 290 291 and microbial biology into invasion science, and for developing impact theory and risk assessment methods that explicitly consider evolutionary change in both the invader and 292 interacting species. 293

294

1.3. The way forward: a theoretical framework and tools for impact management

296 *1.3.1. Developing and expanding a theoretical framework of impact*

To meet societal demands, invasion science must continue to build a body of theory for 297 understanding and predicting impacts from the level of populations to ecosystems (Ricciardi et 298 al. 2013; Blackburn et al. 2014; Bacher et al. 2018). Progress toward this goal requires that 299 hypotheses explicitly integrate abiotic and biotic context-dependencies, including biotic and 300 abiotic drivers of spatiotemporal variation in impact. This integration parallels and perhaps can 301 302 be informed by studies of how species loss affects ecosystem functioning in different environmental contexts (Ratcliffe et al. 2017; Baert et al. 2018; Kardol et al. 2018). One example 303 304 of an integrative hypothesis is Environmental Matching (Ricciardi et al. 2013), which posits that 305 the per capita effects of an invader vary along environmental gradients such that they are maximal where abiotic conditions more closely match the physiological optimum of the invader 306 307 (Kestrup and Ricciardi 2009; Iacarella et al. 2015a; Iacarella and Ricciardi 2015).

A second example that integrates context-dependence is the Ecological (or Functional) 308 Distinctiveness Hypothesis (Diamond and Case 1986; Vitousek 1990; Ricciardi and Atkinson 309 2004), which predicts that impact is most severe in communities missing species functionally 310 similar to the invader. This hypothesis is derived from two observed patterns with strong 311 empirical support. One such pattern is that a community's lack of eco-evolutionary experience, 312 313 or ecological naïveté, determines its vulnerability to non-native consumers, parasites, pathogens, and competitors (Sih et al. 2010; Saul and Jeschke 2015; Davis et al. 2019; Nunes et al. 2019; 314 Anton et al. 2020). The second empirically supported pattern is that the largest community-level 315 316 and ecosystem-level impacts are generated by invaders that use key resources differently or more efficiently than natives do and that can alter disturbance regimes, habitat structure, or food web 317 configurations (Vitousek 1990; Funk and Vitousek 2007; Morrison and Hay 2011). Given that 318 more closely related species tend to be ecologically similar (Burns and Strauss 2011), it follows 319 that phylogenetic distance, or simple taxonomic relatedness, is a proxy for functional 320 distinctiveness. Thus, an allied hypothesis predicts that invaders representing novel taxa, once 321 established in the community, are more likely to affect native populations negatively than 322 invaders that are taxonomically similar to natives in the recipient community (Ricciardi and 323 324 Atkinson 2004; Strauss et al. 2006; Davis et al. 2019). Despite longstanding recognition of ecoevolutionary experience as a driver of impact, most risk assessments do not consider 325 326 evolutionary context. The consequences of the contemporary evolution of non-native species 327 (e.g., Bertelsmeier and Keller 2018), and the effects of invaders on the evolution of native species, are underexploited but promising areas of research (Saul and Jeschke 2015; van Kleunen 328 329 et al. 2018) that point to the importance of integrating evolutionary biology in ways that enhance 330 the predictive power of invasion science.

331	Several distinct, and over a dozen overlapping, hypotheses explain invader impact
332	(Ricciardi et al. 2013), and additional hypotheses addressing invasion establishment success
333	could potentially be extended to understanding impact (Catford et al. 2009; Jeschke and Heger
334	2018). These hypotheses could be organized into a coherent body of impact theory by
335	eliminating redundancies and identifying commonalities (e.g., through consensus mapping of
336	hypothesis networks; Enders et al. 2020). We can envision a general predictive framework built
337	upon multiple axes that consider, among other things, 1) abiotic and biotic environmental
338	context; 2) functional distinctiveness between native and non-native species; and 3) time-since-
339	invasion (Figure 2). The generality of hypotheses needs to be tested within various ecological
340	and evolutionary contexts using, for example, spatially distributed experiments such as those
341	employed to examine plant responses to nutrient enrichment and exclosure of mammalian
342	herbivores (Borer et al. 2014). Experimental and survey designs that incorporate eco-
343	evolutionary context have rarely been applied to the study of non-native species (but see Wardle
344	et al. 2001; Colautti et al. 2014; Grimm et al. 2020). To address this gap, we advocate
345	comparisons of conspecific populations across invaded and native ranges, recognizing that
346	invasions and impact outcomes are population-level phenomena. Such experiments could be
347	coordinated by collaborative global networks (Packer et al. 2017), which are a potentially
348	powerful approach to understand the factors that govern large-scale variation in invader impact
349	across climatic gradients, disturbance gradients, biogeographic realms, and boundaries of
350	evolutionary significance.
351	Moreover, scientists would profit by looking to other areas of ecology and evolution,

disease biology, and the social sciences, for theory that could potentially explain many
 components of impact and seeking to integrate these approaches into invasion science. Several

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classical ecological hypotheses, metrics, and concepts that have been tested in various contexts 354 relevant to invasions (e.g., theories addressing biological control, island biogeography, metabolic 355 scaling, resource utilization, competition) have arguably been underexploited by invasion 356 scientists. Experimental approaches that have sought to incorporate principles of trophic ecology 357 have revealed important patterns (Dick et al. 2017a, b; Cuthbert et al. 2018, 2020). For example, 358 359 prey switching (frequency-dependent predation) is a classical concept that has until recently been virtually ignored by invasion science (Cuthbert et al. 2018, 2019). In recent years, the classical 360 functional response – the relationship between *per capita* consumption and resource density 361 362 (Solomon 1949; Holling 1959) - has been adapted and applied to forecasting and explaining non-native species impacts through multispecies comparisons (Dick et al. 2017a, b; Dickey et al. 363 2018; Faria et al. 2019). The rationale for exploring these experimental approaches is that 364 invasion success and impact are often mediated by resource acquisition, a concept at the 365 foundation of many hypotheses in invasion science (Catford et al. 2009; Ricciardi et al. 2013; 366 Jeschke and Heger 2018) and that is relevant for both animals and plants (Rossiter-Racher et al. 367 2009; Ehrenfeld 2010). Indeed, several high-impact invaders have been found to be more 368 efficient at using limiting resources than their native and non-invasive counterparts (Rehage et al. 369 2005; Funk and Vitousek 2007; Morrison and Hay 2011; Dick et al. 2017a; DeRoy et al. 2020). 370 Broadening analyses to a more comprehensive community context could also help predict 371 impacts in different environmental contexts (Smith-Ramesh 2017). An underexploited approach 372 373 is to treat invaded communities as complex adaptive networks (Lurgi et al. 2014; Valdovinos et al. 2018; Hui and Richardson 2019). Predictive information could potentially be gained from 374 375 modeling the dynamic responses of an ecological network, after developing appropriate metrics

of interaction strength, and thus identify resident species that are either facilitated or suppressedby the invasion (Hui and Richardson 2019).

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379 *1.3.2.* Toward more comprehensive quantifications of invader impact

There is growing interest in quantifying impacts beyond traditional ecological and 380 381 economic measures by using an ecosystem services framework that can capture information on provisioning (e.g., food, timber, fuel), regulating (e.g., climate, floods, nutrient cycling) and 382 cultural services (Perrings 2010; Simberloff et al. 2013). For example, in highly-degraded 383 ecosystems some established non-native species may offer beneficial services to some 384 stakeholders (McLaughlan and Aldridge 2013), although any benefits of local cultivation of such 385 species must be weighed carefully against risks of further spread. Such accounting would also 386 need to consider negative impacts, which are diverse and substantive, on ecosystem services 387 (e.g., Walsh et al. 2016; Vilà and Hulme 2017; Milanović et al. 2020). However, at present we 388 know remarkably little about how even the most high-profile non-native species affect ecosystem 389 services (Vilà et al. 2010; McLaughlan et al. 2014), a problem related to the challenges of 390 evaluating ecosystem-level impacts (Simberloff 2011; Ricciardi et al. 2013). More reliable 391 392 quantification of potential ecosystem services of invasive species, coupled with a deeper understanding of context-dependencies, would allow a more informed and comprehensive 393 394 impact assessment. To this end, the Millennium Ecosystem Assessment and, more recently, the 395 Intergovernmental Platform on Biodiversity and Ecosystem Services (IPBES), which have examined how humans have altered ecosystems and these alterations have affected ecosystem 396 397 services and human well-being (Millennium Ecosystem Assessment 2005; Díaz et al. 2019), 398 could provide a suitable framework for developing protocols for risk assessment, perhaps

399	informed by the EICAT and SEICAT classification schemes (Hawkins et al. 2015; Bacher et al.
400	2018).

Comprehensive impact quantification is challenged by knowledge gaps that may render 401 risk assessments incomplete or misleading (Kumschick et al. 2015). One major gap is predictive 402 knowledge of the role of species traits, combinations of traits, and trait-environment interactions 403 404 in impacts, particularly at the ecosystem level. It is not clear under what situations the same species traits that confer an ecosystem service can also damage an existing ecosystem service 405 (Vilà and Hulme 2017) or contribute to an 'ecosystem disservice' - properties or functions that 406 407 are disadvantageous to humans (Milanović et al. 2020). Another major context-dependency that could distort risk assessment of a given invader is the presence of other invaders. Predictions, as 408 well as post-hoc assessments, are potentially hampered by synergistic or antagonistic interactions 409 between invaders, including those that can contribute to invasional meltdown - in which one 410 invader facilitates another, leading to compounded impacts and potentially self-reinforcing 411 effects (Simberloff and Von Holle 1999; Ricciardi 2001; Green et al. 2011). Disentangling the 412 influence of various species involved in meltdowns requires detailed experimental planning (e.g., 413 Braga et al. 2020), whereas invader interactions in multiple invaded ecosystems are generally 414 415 poorly studied (Kuebbing et al. 2013). It therefore seems likely that most synergistic effects go unrecognized. Even where interactive effects do not occur, the cumulative effects of burgeoning 416 417 numbers of low-impact invaders on ecosystems have been virtually ignored. Approaches toward 418 quantifying and assessing the effects of multiple environmental stressors (Boyd et al. 2018; Hodgson and Halpern 2018; Hodgson et al. 2019) could potentially be adapted for multiple 419 420 invading species and, furthermore, might be enhanced by efforts to collate experimentally-421 validated invader interactions within global databases.

2. Addressing the challenge of global environmental change in invasion science

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The second overarching issue is how invasion science can adapt to the onslaught of 424 global environmental changes presently altering the rates, dynamics, and impacts of invasions 425 426 through myriad drivers including climate change, overharvesting, extinction, pollution, 427 landscape transformation, and shifting trade patterns. Ecosystems are likely to become more susceptible to invasions as these drivers degrade and modify food webs. For some native species, 428 global changes create physiologically intolerable or suboptimal conditions that lower relative 429 fitness (Catford et al. 2020) or provoke range shifts, further altering community composition and 430 431 susceptibility to invader impacts (Gallardo and Aldridge 2013; Wallingford et al. 2020). Environmental change often affects native and non-native species differentially, modifying their 432 interactions and selection pressures through shifting abiotic and biotic ecosystem conditions 433 434 (Xiao et al. 2016; Meyerson et al. 2020; Stern and Lee 2020). This issue is well recognized and has been widely investigated for several years, yet the need for research and management 435 436 solutions through the lens of invasion science is ongoing and increasing. Invasion science must continue to develop an understanding of key issues regarding global environmental change 437 including interactions between invasions and other environmental stressors, climate adaptation 438 and mitigation strategies, and evaluating and managing species range shifts and translocations. In 439 this section, we primarily focus on climate change (Figure 3) but note that many other forms of 440 human-induced environmental change facilitate invasions and the relative dominance of non-441

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444 **2.1.** Ecological synergies between invasions and climate change

native species (Catford et al. 2014; Seabloom et al. 2015; Liu et al. 2017; Essl et al. 2019).

445 *2.1.1. Non-native species performance*

Species distributions worldwide are mostly determined by climate, tectonic movements, 446 and orographic barriers (Ficetola et al. 2017). Climate change will therefore have a major impact 447 on species range and distributions irrespective of whether species are native or non-native to a 448 particular region. However, differences in the magnitude of potential range shifts predicted for 449 450 non-native and native species will be determined by differences in their biology, such as physiological tolerances and dispersal potential (Essl et al. 2019). The last decade has 451 accordingly seen major efforts to investigate the role of climate change in the introduction, 452 453 establishment, spread, and impact of non-native species (Hulme 2017). Various meta-analyses have shown that non-native species often outperform and adjust 454 better than native species to a rapidly changing climate (Sorte et al. 2013; Oduor et al. 2016; Liu 455 et al. 2017). For example, hotter, drier environmental conditions enable non-native Asian tiger 456 mosquitoes to outcompete native tree-hole mosquitoes in the United States (Smith et al. 2015), 457 Eastern mosquitofish (Gambusia holbrooki) persist more successfully than native fish species in 458 France (Cucherousset et al. 2007), and non-native Monterey pine (Pinus radiata) to grow faster 459 than native conifers in Spain (Godoy et al. 2011). Warmer temperatures in freshwater 460 ecosystems will favor non-native species as these frequently have a greater heat tolerance than 461 related native species (Bates et al. 2013); similarly, in the Mediterranean Sea, increases in 462 temperature have facilitated the establishment of non-native tropical species (Raitsos et al. 2010). 463 464 A key element of climate change is an increase in the frequency and magnitude of extreme climatic events, which can have greater effects on invasion than changes in average 465 466 conditions (Sheppard et al. 2012). Strong winds, floods, large waves, and storm surges can 467 transport organisms into new regions (Diez et al. 2012), as discussed below. Critically, extreme

climatic events like heat waves, fires, severe storms, droughts, and floods act as major 468 disturbances and will invariably destroy and damage resident native biota, reducing the uptake of 469 resources, and can also increase resource supply (Catford and Jones 2019). Such disturbances are 470 known to facilitate invasion (Davis et al. 2000), because many invasive species can take 471 advantage of fluctuations in resource availability caused by disturbances (Catford et al. 2012; 472 473 Singh et al. 2018). For example, European *Bromus* grasses that are highly invasive in North America can exploit available soil moisture more efficiently and thus recover more rapidly than 474 native vegetation after drought (Harris 1967), enabling them to invade areas formerly dominated 475 476 by native woody species (Kane et al. 2011). Similarly, a non-native freshwater phytoplankton species was able to invade and establish in a reservoir following the combined disturbance events 477 of macrophyte removal and extreme drought (Crossetti et al. 2019). 478

479

480 2.1.2. Non-native species range shifts

Shifts in temperature and rainfall patterns attributed to climate change can increase the 481 probability of establishment of non-native species, which were previously constrained by climate 482 (Walther et al. 2009; Hulme 2017) or climate-mediated interactions with native biota (Catford et 483 484 al. 2020). Increasing evidence indicates that non-native species tend to respond faster than native species to climate change, with spread rates an order of magnitude higher than the velocity of 485 486 climate change (Hulme 2012). For example, non-native plants have expanded upwards in the 487 European Alps twice as fast as native species in response to warming (Dainese et al. 2017). Nevertheless, climate change can lead to both increases (Kriticos et al. 2003; Barbet-Massin et 488 489 al. 2013; Gilioli et al. 2014) and declines (Bradley et al. 2009; Bellard et al. 2013; Xu et al. 2014) 490 in the geographical range of non-native species. A general finding is that, as a result of climate

change, the distribution range of non-native invertebrates and pathogens will expand, but range 491 contractions are mostly expected for non-native plants and vertebrates (Bellard et al. 2018). For 492 example, by the end of this century the suitable area worldwide for the red imported fire ant 493 (Solenopsis invicta) is predicted to be 21% greater (Morrison et al. 2014), whereas for the velvet 494 tree (*Miconia calvescens*) it is predicted that suitable habitat will be reduced in both its native 495 496 and introduced ranges (González-Muñoz et al. 2015). However, trends may differ between terrestrial and aquatic environments. For instance, the warming of North American lakes is likely 497 to increase thermal suitability for southern species of fishes that could expand their distribution 498 499 poleward into non-native regions, potentially as far as the Arctic (Sharma et al. 2007; Della Venezia et al. 2018). 500

Besides overall change in temperature and precipitation, extreme climatic events can also 501 help spread non-native species by overcoming dispersal barriers (Diez et al. 2012). For instance, 502 hurricanes promoted dispersal of non-native cactus moth (Cactoblastis cactorum) across the 503 Caribbean and into Mexico where it threatens native Opuntia species (Andraca-Gómez et al. 504 2015). Hurricane frequency was also positively correlated with the expansion of the non-native 505 grass *Phragmites australis* across wetlands along the Gulf Coasts of the USA (Bhattarai and 506 507 Cronin 2014). Likewise, flood events can increase pool connectivity and provide non-native freshwater species access to newly inundated areas (Vilizzi et al. 2014). For example, floods 508 enabled the escape of cultured black carp (Mylopharyngodon piceus) in the Missouri River, US 509 510 (Nico et al. 2005), and tilapia cichlids in southeast Asia (Canonico et al. 2005) and have facilitated the spread of zebra mussels (Dreissena polymorpha) in the Mississippi River 511 512 catchment (Tucker 1996). Nevertheless, the natural variability of climate makes it difficult to

attach high levels of confidence to some of the predicted changes, particularly those associatedwith extreme weather events (Bellard et al. 2013).

- 515
- 516 *2.1.3. Novel interactions and per capita impacts*

Climate change will, in many cases, increase the introduction rate, establishment 517 518 probability, and spread rate of non-native species (Bellard et al. 2013), while simultaneously facilitating extensive range shifts of native species (Inderjit et al. 2017; Pecl et al. 2017; Essl et 519 al. 2019), leading to novel ecological interactions and increased impacts. Range shifts are 520 521 expected to contribute to widespread biotic homogenization (where more species are shared among communities) in some regions and the formation of novel communities in others (García-522 Molinos et al. 2015). Diverse novel biotic interactions and assemblages will arise from divergent 523 responses of species and populations to climate change (Blois et al. 2013; Pecl et al. 2017). As 524 discussed previously, new biotic interactions often result in high impacts when resident species 525 have not co-evolved with newly arrived species (Ricciardi and Atkinson 2004; Cox and Lima 526 2006; Saul and Jeschke 2015). In some cases, range shifts of native species can cause impacts 527 similar to those involving non-native species (Sorte et al. 2013; Inderjit et al. 2017), although 528 529 impacts will be tempered by the eco-evolutionary experience of the resident species (sensu Saul and Jeschke 2015). Few studies have addressed range shifts of native and non-native species as a 530 531 joint issue (Gallardo and Aldridge 2013; Sorte et al. 2013; Dainese et al. 2017; Inderjit et al. 532 2017; Singh et al. 2018).

While many studies have linked climate change to the spread of invasive species (detailed above), the role of environmental factors in determining ecological impacts is understudied (Dickey et al. 2020). Climatic conditions that shift towards the physiological

optimum of a non-native species could promote increased feeding rates, growth, or reproduction 536 that amplifies its competitive or predatory effects (Hellmann et al. 2008; Iacarella et al. 2015a). 537 For example, an invasive bryozoan is expected to have enhanced growth rates at warmer 538 temperatures in the Northwest Atlantic, with greater modeled impacts on kelp beds under future 539 climate conditions (Denley et al. 2019). Similarly, higher growth rates enable an invasive plant 540 541 to outcompete a native plant in China along higher latitudes in the field and at warmer experimental temperatures (Wu et al. 2017). Predation rates of non-native species may also 542 increase when warming temperatures are within the physiological optima of the invader 543 (Iacarella et al. 2015a). For instance, the predatory response of an invasive freshwater amphipod 544 increases when exposed to elevated temperatures and infected by a common parasite (Laverty et 545 al. 2017). Given that non-native species are expected often to outperform native species in 546 response to environmental change, as discussed above, their competitive and predatory impacts 547 will likely also increase under these circumstances. A method has recently been developed that 548 incorporates the per capita and abundance effects of non-native species under altered variables 549 such as temperature, oxygen, salinity, and indeed any other variable in isolation or combination 550 (Dickey et al. 2020). This predictive method crucially also factors in the climate response of the 551 552 affected species (e.g., native prey), such that overall impact is holistically predictable. This method is in its infancy and ground-truthing is now limited only by data (Dickey et al. 2020). 553

554

555 2.1.4. Changes to ecosystem services and human well-being

Research on the interaction between invasions and global environmental change is 556 essential to identify effects on ecosystem services and human well-being (Dukes and Mooney 557 558 1999; Walther et al. 2009; Pecl et al. 2017; Vilà and Hulme 2017). Although tools such as

SEICAT (Bacher et al. 2018) and INSEAT ('INvasive Species Effects Assessment Tool'; 559 Martinez-Cillero et al. 2019) have been developed to classify non-native species within a 560 561 framework of ecosystem services and human well-being, these tools rely on expert elicitation as there are still surprisingly few quantitative data on the ecosystem services effects of even the 562 most prolific invasive species. This is, in part, owing to the context-dependent impacts of 563 invaders (see section 1) and because environmental change can alter the balance of positive and 564 negative effects (McLaughlan et al. 2014). For instance, disturbed river banks and roadsides in 565 Africa favor proliferation of the invasive tree, *Prosopis juliflora* (Shiferaw et al. 2019), which 566 increases local income from wood sales but reduces habitat suitable for livestock and results in 567 lower income from cattle sales (Linders et al. 2020). The predicted future effect of interactions 568 among climate, socioeconomic factors, and invasions on plant biodiversity hotspots constitutes 569 570 the greatest threat in emerging economies located in megadiverse regions of the Southern Hemisphere (Seebens et al. 2015). Invasions and climate change also pose a combined threat to 571 572 native species in protected areas and thus seriously compromise conservation of biodiversity and ecosystem services (Gallardo et al. 2017; Iacarella et al. 2020). Interactions between invasions 573 574 and climate change will also affect human health; for instance, climate change models predict an 575 increase in the life-cycle completion rate and extended periods suitable for development of the 576 invasive mosquito Aedes aegypti, a vector of arboviruses including dengue, zika, and yellow 577 fever, resulting in accelerated invasion in North America and China (Iwamura et al. 2020). 578 To investigate the effects of invasions on ecosystem services and human well-being, 579 models should integrate interactions among several components of global change, not only climate change (Walther et al. 2009). Furthermore, studies should also explore these interactions 580 581 in productive systems such as managed forests, agriculture, and aquaculture (Thomson et al.

2010; Ziska and Dukes 2014; Liebhold et al. 2017). A major concern for these resource sectors is that drought, warming, and elevated CO_2 will affect the performance of non-native species (i.e., pests, pathogens, and weeds) in complex and currently unpredictable ways. Research on their impacts requires, for example, quantifying not only how altered environmental conditions change weed and crop performance in isolation, but the magnitude of weed-crop competition on crop damage (Ramesh et al. 2017).

588

589 2.2. Human responses to climate change that favor non-native species

590 *2.2.1. Changes to invasion pathways*

Global change is also altering invasion risk by promoting new commercial trading routes 591 and corridors. Shifting global economic forces (e.g., tariffs, manufacturing trends, recession, 592 regional conflicts, climatic disasters) determine trade volume and thus the frequency with which 593 aircraft or oceanic vessels travel between airports or seaports (Seebens et al. 2015). Such shifts 594 drive temporal rates of species introduction and the range of taxa that invade (Levine and 595 D'Antonio 2003; Hulme 2015; Bertelsmeier et al. 2018). For example, commercial shipping at 596 polar latitudes of North America and Eurasia is either planned or already occurring, providing 597 novel opportunities for introducing non-native species to Arctic waters (Miller and Ruiz 2014; 598 Chan et al. 2019). The Southern Ocean is likewise becoming increasingly vulnerable to species 599 introductions owing to increased propagule pressure from vessel traffic and reduced physical and 600 601 physiological barriers (Aronson et al. 2015; Hughes and Ashton 2017; Smith et al. 2017; McCarthy et al. 2019; Cárdenas et al. 2020). Such human responses to climate change (Figure 3) 602 603 are altering the origins, taxonomic identity, and rate of introduction of non-native species in

terrestrial, freshwater, and marine habitats worldwide (Seebens et al. 2015; Early et al. 2016;
Della Venezia et al. 2018).

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607 *2.2.2. Climate adaptation: planting non-native species and adding infrastructure*

As governments increasingly develop adaptive strategies to address climate change, 608 609 many of these strategies are likely to entail using non-native species. Proposed interventions include initiatives to develop agricultural or aquacultural enterprises to deliver carbon-neutral 610 energy sources (e.g., macroalgae and plants for biofuels) using known invasive non-native 611 612 species (Barney and DiTomaso 2008). Pressure is also increasing to develop new varieties of pasture species that can better cope with changing climates, such as drought-tolerant and disease-613 resistant species, many of which are non-native in the countries in which they are sold and 614 planted (Driscoll et al. 2014). Increased development of green roofs, vertical gardens, and water-615 saving horticulture to mitigate effects of climate change (Perini and Rosasco 2016) carry the risk 616 of introducing non-native species by promoting drought-tolerant plants or breeding drought-617 resistant varieties, cultivars, or hybrids. Similarly, many large-scale tree-planting programs have 618 not led to the replenishment of degraded forests with native tree species, but rather to 619 620 afforestation of non-forest land, including biodiverse grsslands, with monocultures of non-native trees. Such efforts include massive tree-planting campaigns using non-native trees with the aim 621 622 of mitigating the impacts of climate change and for other poverty alleviation (Brundu et al. 623 2020). Such plantings might not help offset greenhouse gas emissions as much as expected, owing to unforeseen fluxes and complex system dynamics (Covey et al. 2012; Luyssaert et al. 624 625 2018; Popkin 2019). Indeed, inappropriate afforestation, especially in naturally treeless areas, 626 can have serious consequences for sustainable development, biodiversity conservation, and

ecosystem functioning (reviewed in Brundu et al. 2020). Furthermore, many species used in such 627 programs are highly invasive, which means that their impacts extend beyond areas identified for 628 afforestation (Brundu and Richardson 2016; Brundu et al. 2020). 629 Besides directly introducing species to sustain economic activities or to mitigate 630 emissions, governments at all levels are responding to environmental change by developing new 631 632 infrastructure. Strategies to combat sea-level rise have largely been addressed through engineered solutions (armoring, raising road-beds, flood control structures). Each of these 633 adaptation strategies presents an opportunity for existing non-native species to expand their 634 range or impact and can create new suitable habitat for non-native species that arrive via ballast, 635 hull-fouling, or the marine aquarium trade (Bulleri and Chapman 2010). Offshore wind farms 636 also provide novel fouling habitats and 'stepping stones' for invasions (Adams et al. 2014; De 637 Mesel et al. 2015). Similarly, frequent droughts lead to efforts to provide secure water sources to 638 urban populations, including construction of dams, canals, and other water-diverting mechanisms 639 that can spread non-native species (Strayer 2010; Zhan et al. 2015; Gallardo and Aldridge 2018). 640 However, infrastructure developments can be designed to reduce their suitability as novel 641 habitats or invasion routes for invasions by non-native species, by minimizing environmental 642 643 disturbances or emulating natural habitats (Dafforn et al. 2015).

644

645 2.2.3. Species translocations for conservation

Conservation scientists have introduced species to locations outside their native range for
three main reasons: (1) to avoid extinction caused by an introduced species, often an introduced
predator; (2) to restore an ecological function (as detailed in section 1.2.1); or (3) to allow
species' ranges to keep up with climate change (Corlett 2016). Introductions to accommodate

global climate change have increasingly attracted attention. As early as 1985, conservationists 650 recognized that the climate of current species ranges will change so that locations with climate 651 similar to that of today may be distant or separated by inhospitable habitat; they proposed several 652 measures including direct human assistance in the form of translocation to suitable habitat 653 unoccupied by the species of interest when adequate autonomous movement seemed unlikely 654 655 (Peters and Darling 1985; Peters 1988; Davis 1989; Peters 1992). This proposal received little interest for the next decade; a review of possible management responses to climate change listed 656 only 13 mentions of translocations (Heller and Zavaleta 2009). None of these acknowledged 657 658 possible negative effects of translocation. However, translocations had long been conducted in the name of conservation, notably of species threatened by introduced predators (Seddon et al. 659 2012, 2014a). For instance, endemic New Zealand birds threatened by non-native rats and 660 mustelids had been translocated to predator-free islands since 1894, with many well-publicized 661 projects (Clout and Craig 1995; Seddon et al. 2012); occasional concern about such efforts had 662 663 been expressed on the grounds of potential unanticipated ecological impacts (e.g., IUCN 1987; Conant 1988; Atkinson 1990; Craig and Veitch 1990; Towns et al. 1990). 664 Translocation as a management response to climate change began to gain substantial 665 666 attention with papers by McLachlan et al. (2007) and Hunter (2007), both raising the issue that this constitutes introducing a non-native species, which in turn might lead to a damaging 667

668 invasion. Hoegh-Guldberg et al. (2008) produced the first decision tree for application of

potential translocations in response to climate change, but they, and Hunter (2007), suggested

- 670 that intercontinental introductions have proven far more likely to be damaging, whereas
- 671 proposed translocations for climate change would be more restricted. A broader and more
- detailed criticism of climate change-motivated translocation (Ricciardi and Simberloff 2009a),

673	based on the possibility of non-target impacts, elicited an exchange with several respondents
674	(Ricciardi and Simberloff 2009b) and signaled a shift in the dialogue, with much more attention
675	paid to the possibility of unintended consequences by virtue of introducing new species. As such,
676	Richardson et al. (2009) expanded the decision-tree approach into a heuristic decision tool with
677	detailed considerations of both ecological and socioeconomic consequences of translocation or
678	failure to translocate; the difficulty lies in estimating the probability of various potential
679	outcomes (e.g., decline or loss of ecological functions in the recipient region) and quantifying
680	other risks, both ecological and socioeconomic, to inform comparisons and decisions. In the last
681	decade, translocation has received increasingly nuanced consideration of the relative risks and
682	virtues owing to the rapidly growing understanding of the enormous conservation challenge
683	posed by the scope and imminence of climate change and its likely effect on species ranges
684	(Hewitt et al. 2011; Thomas 2011; Schwartz and Martin 2013; Williams and Dumroese 2013;
685	Ricciardi and Simberloff 2014; Maier and Simberloff 2016; Simler et al. 2018).
686	The lines between translocation and biological invasion are becoming increasingly
687	blurred. Both events involve species expanding beyond their historical biogeographic ranges,
688	leading some authors to suggest that they differ only in public perception and value (Hoffmann
689	and Courchamp 2016; but see Ricciardi 2007; Wilson et al. 2016). In addition, views on how to
690	deal with the spectrum from 'desirable' self-migrating species, to translocations undertaken for
691	conservation (desirable to some, undesirable to others), to generally 'undesirable' biological
692	invasions, are yet to be reconciled. Further, determination of 'non-native', as defined by lack of
693	co-evolution with the invaded community (Ricciardi 2012), and 'desirable' or 'undesirable', as
694	defined by valuations of impact (Jeschke et al. 2014), will become increasingly challenging as
695	we seek to determine what to protect or manage in a shifting mosaic of species assemblages

(Gilroy et al. 2017; Hill and Hadly 2018). The current framework for managing non-native
species could yield protection of conservation-based translocated species despite potentially high
impacts, compared to management and mitigation of high-impact species that spread via selfdirected or direct, but accidental movement. Robust protocols for considering the entire range of
possible impacts of facilitated range shifts, as well as those of self-migrating species, must be
developed and integrated into policies and legislation with the engagement of stakeholders.

702

703

2.3. Government responses and global efforts

704 The global nature of biological invasions and their interactions with environmental change can strain the capabilities of governments to anticipate and respond to invasions now and 705 into the future. As discussed in detail above, the ecology of invasions under climate change is 706 707 complicated. The directed asymmetrical movement of certain species poleward (Winter et al. 2014), and to higher elevations (Pyšek et al. 2011; Dainese et al. 2017), can point to systems 708 requiring early-detection monitoring or intervention. On the other hand, the effects of climate 709 change could play out neatly along latitudinal or altitudinal gradients (Hanberry and Hansen 710 2015). A key unknown is the relative importance of introduction enhancement (e.g., colonization 711 712 pressure, propagule pressure) from changing trade patterns versus the influence of climate change factors in facilitating species' range changes. Policies that address invasions could also 713 714 be complicated by seemingly competing interests, including those associated with the economy 715 and trade versus biodiversity and human health. Despite devastating new species invasions and pleas for a comprehensive approach to biosecurity, some countries, such as the United States, 716 717 have even recently reduced coordinated federal leadership and investments to address invasions 718 (Meyerson et al. 2019; Simberloff et al. 2020). Current coordinated global efforts to document

invasions and impacts include the Global Register of Introduced and Invasive Species (GRIIS)
and an invasive species assessment by the IPBES. These substantial undertakings will
undoubtedly deepen our understanding of invasion trends, impacts, and management, but neither
will result in policies to prevent species introductions that participating nations are obliged to
adopt. Rather, it remains the role of national and local governments to identify, fund, implement,
and enforce policies to manage invasions under changing conditions and, where possible, to
coordinate with other nations.

726

727 **3. Resolving the Taxonomic Impediment**

728 **3.1.** The enduring problem of taxonomic identification

The third overarching issue is our capacity to distinguish non-native from native species 729 accurately. Scientific understanding of the processes that control the diversity, abundance, 730 distribution, and impacts of non-native species ultimately depends on the quality of taxonomic 731 data. The steady global erosion in training and expertise in systematics means that invasion 732 733 science often lacks the taxonomic support to accurately identify many taxonomic groups in terrestrial, freshwater, and marine habitats (Godfray 2002; and below). While this phenomenon 734 exists across biomes and taxa, the largest gaps in taxonomic knowledge are associated with some 735 of the most abundant species, including microorganisms and microfauna. Arguments (such as 736 those of Costello et al. 2013) that the field of taxonomy is robust appear to be based on, among 737 738 other fallacies, a misinterpretation that authorship inflation on taxonomic papers equates to an 739 increasing number of taxonomists (Bebber et al. 2013; Daglio and Dawson 2019). Molecular tools have made remarkable progress and offer great promise for illuminating 740 the overlooked scale of biodiversity in all habitats (Hebert et al. 2003; Dinca et al. 2011). The 741

application of modern sequencing techniques often reveals a vast array of unknown and often 742 cryptic species. Srivathsan et al. (2019) report that of 7,059 specimens of flies (Diptera, family 743 Phoridae) collected in a single Malaise trap in Uganda over an eight-week period, MinION 744 sequencing revealed more than 650 largely or entirely undescribed species, exceeding the total 745 number of phorid taxa described for the entire Afrotropical region. Only one of these 650 746 747 species, however, has to date been formally described, based on morphological characters, as a new species. Molecular techniques combined with advanced culturing methods have revealed an 748 enormous diversity of microbial taxa. Metagenomic sequencing of samples from only 68 ocean 749 750 locations revealed over 35,000 microbial 'species' (Sunagawa et al. 2015). Locey and Lennon (2016) predict that the Earth may support as many as a staggering 1 trillion (10^{12}) microbial 751 species. 752

Nevertheless, the use of molecular technologies to identify taxa to the species level by 753 genetic fingerprinting or 'barcoding' has often proven to be an insufficient and unreliable 754 755 response to the taxonomic impediment. The panacea that simply sequencing specimens and trusting that matching those sequences to databases will produce a reliable identification has 756 proven not to compensate for the growing gap in taxonomic expertise. Two principal problems 757 758 hinder molecular identifications: (1) accurate and complete barcoding of taxa across the taxonomic spectrum, and (2) accurate and complete reference databases against which taxonomic 759 assignments are made (Harris 2003). 760

These problems are hindering the compilation of inventories of non-native taxa, even in conspicuous and well-studied groups such as Australian *Acacia* species (Magona et al. 2018). Taxonomic biases may result from the markers used (Clarke et al. 2014), while many species for which sequences are obtained have no authenticated database reference (Briski et al. 2016). 765

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Further, all new sequence entries should (but do not) require that the sequenced taxon has been

705	i utilei, un new sequence entries should (out do not) require that the sequenced taxon has been
766	identified by a taxonomic expert based upon morphological evidence. Thus, a substantial fraction
767	of the species in these databases can be misidentified, at times egregiously so, potentially
768	producing erroneous matches that cannot be detected by non-specialists (Figure 4). For example,
769	DNA barcoding sequence information is missing from either the Barcode of Life Database,
770	GenBank, or both, for 60% of the 88 insect species listed in the Global Invasive Species
771	Database; 41% of the 88 species could be misidentified as another species, owing to
772	discrepancies between sequences and species identity (Boykin et al. 2012).
773	
774	3.2. Taxonomic impediments lead to under-estimations of invasion
775	Without changes to ensure the development of broad taxonomic expertise, invasion
776	science will continue to underestimate, often substantially, the number (and also, therefore, the
777	impacts) of non-native species across all habitats, regardless of the surveillance and detection
778	program (e.g., De Barro et al. 2011). For example, Carlton and Fowler (2018) recently estimated
779	that non-native species are under-reported globally for the majority of marine taxonomic groups,
780	owing to a lack of widely available taxonomic expertise. Conversely, what has been initially
781	viewed as an invasion by one widespread species is sometimes later discovered to be a group of
782	similar species, some or all of which are restricted to their native ranges (Darling and Carlton
783	2018). More broadly, the inability to detect what could be the most common new invasions (by
784	species and genotypes) across terrestrial, freshwater, and marine habitats undermines ecosystem
785	management and biodiversity assessment, and our capacity to detect changes in ecosystem
786	structure and function.

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787	The deficit of taxonomic expertise associated with microorganisms is especially
788	worrying. In general, species richness and density of organisms are inversely related to size. Not
789	only do small-bodied creatures dominate the world, but the magnitude of non-native species
790	transfers is often greatest for small organisms, many of which have life histories that facilitate
791	colonization (e.g., asexual reproduction; resting stages) (Ruiz et al. 2000). While invasions of
792	microorganisms are increasingly recorded (Seebens et al. 2017), the extent of these invasions,
793	and their impacts, remain poorly described outside of forestry, agriculture, and aquaculture
794	(Desprez-Loustau et al. 2007; Lohan et al. 2020).
795	Several marine disease outbreaks (such as those in oysters, sea urchins, and fishes) have
796	been attributed to non-native pathogens. For example, MSX is an oyster disease caused by a
797	protozoan (Haplosporidium nelsoni) that is native to Asia but was detected on the Atlantic coast
798	of North America in 1957. The native eastern oyster (Crassostrea virginica) proved highly
799	susceptible to MSX, leaving local populations substantially depleted from Chesapeake Bay to
800	Nova Scotia (Bushek and Ford 2016). The same is true for terrestrial and freshwater habitats
801	where non-native pathogens cause diseases such as ash dieback, crayfish plague,
802	chytridiomycosis, and sudden oak death (Skerratt et al. 2007; Grunwald et al. 2012; Roy et al.
803	2017). Many of these non-native pathogens spill-over, colonizing native host species in the
804	invaded range, whereas non-native hosts may harbor native parasites that then spill-back to
805	native hosts (Roy and Handley 2012; Blackburn and Ewen 2017). Both effects complicate
806	parasite identification (Morand 2017). Given recent work on the role of microbial communities
807	in ecosystem processes (Worden et al. 2015) and their importance in microbiomes, host-parasite
808	interactions (Egan and Gardiner 2016), and plant mutualisms (Traveset and Richardson 2014),
809	the potential importance of microorganism invasions is enormous. Thus, evaluation of

microorganism biogeography is a high priority if we are to understand the full scope and impactof invasions in all ecosystems.

812

813 **3.3.** Lack of taxonomic expertise limits our ability to test and develop invasion theory

The taxonomic impediment also impairs our ability to evaluate and understand the 814 815 spatiotemporal dynamics of invasions and their impacts. Much of the theory and current knowledge of invasion science has arisen from syntheses and analyses of secondary data drawn 816 from regional checklists and distribution atlases of floras and faunas (van Kleunen et al. 2015, 817 2019; Dyer et al. 2017; Pyšek et al. 2017). However, such checklists and databases can be 818 seriously compromised by the quality of species identifications (McGeoch et al. 2012). 819 Identifying plant hybrids, in particular, requires professional taxonomic expertise and is crucial 820 for management, given that hybridization often facilitates establishment (Yamaguchi et al. 2019) 821

and stimulates invasiveness, where the new taxon is more vigorous than either parent (Ellstrand

and Schierenbeck 2000; Vilà et al. 2000). An example is provided by *Fallopia* taxa (knotweeds)

824 in the Czech Republic, for which redetermination of plants in the field revealed

misidentifications for up to 16% of the records reported in the literature or deposited in herbaria

for *Fallopia japonica* and *F. sachalinensis*, and 20% of records of the hybrid *F.* \times *bohemica*,

827 (Pyšek et al. 2001). Only after the complicated patterns of increased ploidy variation and rapid

post-invasion evolution in the invaded range of Europe were disentangled was it possible to

conduct ecological studies that revealed the elevated invasiveness of the hybrid compared to that

of the parents (Pyšek et al. 2003).

831 Other taxonomic challenges in plant invasion research include apomictic groups,

karyologically variable complexes, genera with specific reproduction systems, or those for which

horticulturalists have bred many cultivars and varieties (e.g., *Centaurea*, *Cotoneaster*, *Heracleum*, *Lupinus*, *Myriophyllum*, *Phragmites*, *Rhododendron*, *Rubus*, *Spartina*, and *Tamarix*).
Some of those taxa are among the most widespread plant invaders, and ecological studies aimed
at understanding their invasion have profited substantially from detailed taxonomic knowledge
(Pyšek et al. 2013).

838

839 3.4. Lack of taxonomic expertise limits our ability to manage invasions

Taxonomic expertise is fundamental to management and policy efforts, from border 840 control to early detection (and both encouraging and justifying rapid response based on expert 841 identification) to post-invasion management. In several cases, misidentifications and failures to 842 recognize cryptic species complexes have delayed the discovery and introduction of suitable 843 biological control agents (Anderson and Wagner 2016). This is illustrated by biological control 844 of Cactaceae in South Africa that was delayed because the wrong species of herbivorous insect 845 846 was collected. After taxonomic problems were resolved and the appropriate insect was released, the population of the non-native cactus declined (Paterson et al. 2011). Similarly, a carnivorous 847 beetle, Laricobius naganoensis, was inadvertently imported to eastern North America with a 848 849 closely-related species, L. osakensis, introduced from Japan to control an invasive insect – the hemlock woolly adelgid. The U.S. Department of Agriculture subsequently permitted further 850 851 introduction of L. naganoensis, requiring no risk assessment or monitoring, simply because it 852 was too difficult to distinguish it readily from its congener (Leppanen et al. 2019).

Food security is also compromised by taxonomic problems. Inability to determine species identity in imported live seafood can result in widespread substitution by cheaper species in many countries, some of which include invasive non-native species. For example, in South

Africa several species of *Clarias* catfish are native to the continent and are used in aquaculture as
a local food source. However, the walking catfish (*C. batrachus*) – a southeastern Asian species
known to cause detrimental impacts where it has established – is prohibited for aquaculture. The
walking catfish is difficult to distinguish from its African congeners based on morphology alone,
making it an easy species to label inappropriately, import, grow, and sell (Grobler et al. 2015).
Equally worrisome is that, with the rapidity with which vectors and pathways are
changing in today's globalized economy, we may be unaware of – and unprepared for – many

future invasions. The widening gap between our desire to assess changing biodiversity and our ability to identify species implicates all taxa in all habitats and thus compromises our evaluation of the consequences of invasion. The need to narrow this gap through enhanced taxonomic expertise is crucial if we are to keep pace with the constantly expanding numbers of non-native animals and plants being introduced across the planet (Seebens et al. 2018, 2020).

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3.5. The way forward: Training the next generations of researchers to identify species

The way forward requires a new international emphasis on the value of taxonomy. The 870 foundations of the scientific community's ability to recognize biodiversity, including the 871 presence and impacts of non-native species, have been crumbling for decades. Rebuilding these 872 foundations requires consensus that we need to do so, accompanied by agreement of the scale of 873 874 restoration required, a plan to undertake renovation, and the commitment and capital to see it 875 through. Each of these stages, except for commitment and capital, has been discussed exhaustively to little avail. The challenge of old and oft-repeated clarion calls is that they fall on 876 deaf ears, or worse. And yet without this commitment, the global number of scientists who are 877 878 trained in the basics of taxonomy (including expert field identification) and possess skills in

measuring biodiversity will continue to diminish (Lücking 2020). Failures to identify organisms
correctly will lead to spurious conclusions in ecological studies and ultimately to inappropriate
and ineffective legislation, management, and policy (Pyšek et al. 2013). We note that recent
championing of *taxonomic sufficiency* or the Higher Taxon Approach, which is designed to
circumvent either the absence of, or the need for engaging, expert resources (de Oliveira et al.
2020; Gerwing et al. 2020), is inapplicable to invasion science – which requires the highest
quality and accuracy of species-level identification.

Thousands of protist, animal, and plant phylogenies produced every year contain 886 innumerable clades deprived of binomial nomenclature (Darling and Carlton 2018). Students 887 engaged in such work should be trained, supported, and encouraged to provide taxonomic 888 descriptions of clades as new species. The great satisfaction and pride of describing and 889 publishing new species, including the honor of choosing a name, could inspire a measurable 890 fraction of future generations of biologists and ecologists to become recognized taxonomic 891 892 experts while at the same time remaining experts in other specialties. A key advance will be the dissolution of the enduring myth that simultaneously being an expert taxonomist and an expert 893 ecologist (or neurobiologist or molecular biologist) is impossible. Building pride in contributing 894 895 to global biodiversity knowledge is a critical step in addressing the taxonomic impediment in the 21st century. While we champion the rapidly growing concept of *integrative taxonomy* (Daglio 896 897 and Dawson 2019; Zhang 2020) – what Boxshall (2020) describes as the "reciprocal illumination 898 of morphological systematics and molecular sequence-based systematics" - we emphasize that no integration is possible if only one partner is on the stage. The central role of taxonomists in 899 900 resource management, biodiversity conservation, and biosecurity must be affirmed (Hutchings 901 2020). The decline in funding and the startling erosion of taxonomic positions in museums and

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902 other institutions must be addressed through novel collaborations, underscoring societal903 significance.

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4. Enhancing international biosecurity and multi-stakeholder cooperation

4.1. Shifting international trade and travel patterns mediate invasions

907 A final overarching issue is the need for invasion science to provide better guidance for biosecurity programs, at both national and international levels. The suite of species transferred 908 between regions varies as global trade patterns wax and wane (Dyer et al. 2017; Seebens et al. 909 2018). An emerging example is the vast Chinese 'Belt and Road' initiative, which can potentially 910 911 elevate invasion risks greatly among the more than 120 countries through the development of a series of land-based economic corridors between core cities and key ports (Liu et al. 2019). 912 Historically, changes to biosecurity policies that focus on specific pathways have been motivated 913 914 by the impacts of species arriving via those pathways, but the effectiveness of such reactive approaches to policy development is hampered by long lags between the establishment of 915 916 pathways and the onset of invasion. Novel, forward-looking approaches to pathway risk analysis are needed. For example, internet commerce of plants and animals is an expanding global 917 pathway that can radically transform the composition and introduction routes of species in trade 918 (Humair et al. 2015). Structural changes to the horticultural industry, such as the shift to off-919 shore production, have major implications for plant health and trajectories of biological 920 invasions (Dehnen-Schmutz et al. 2010). Another emerging pathway is ecotourism; well-921 meaning nature enthusiasts unwittingly introduce non-native species even to remote regions 922 (e.g., Nash 2009). Research is needed to develop educational and social engineering tools that 923 can be used to alter tourist behavior to reduce risks of future invasions. 924

Contemporary problems with non-native species reflect economic, societal, and trade 925 drivers and patterns that prevailed over the past few centuries (Essl et al. 2015a; Hulme 2015; 926 Dyer et al. 2017; Zieritz et al. 2017). This means that interventions to regulate pathways and 927 their effects on invasions are out of sync and that time horizons of decades must be considered in 928 strategic planning. The time lags inherent in many biological invasions imply that many 929 930 additional non-native species are destined to become established and cause problems in the coming decades, even if biosecurity measures are radically improved (Essl et al. 2011). The 931 dimensions and implications of this invasion debt are yet to be clearly incorporated into strategic 932 933 biosecurity planning anywhere in the world (Rouget et al. 2016). To this end, Wilson et al. (2018) included indicators pertaining to four components of invasion debt (introduction debt; 934 establishment debt; spread debt; and impact debt) among 20 indicators for reporting on 935 biological invasions at the national level. These indicators form the basis for regular reporting on 936 the status of biological invasions developed for South Africa – the first country to have instituted 937 such a comprehensive reporting protocol (van Wilgen et al. 2020). Uptake of such measures for 938 all countries is a priority. 939

Trends observed in past invasions, most of which have played out over the last five 940 decades, provide imperfect insights for planning of biosecurity interventions, since many aspects 941 of future invasions (e.g., taxa involved, pathways, drivers of progression along the introduction-942 943 establishment-spread continuum, interactions between drivers) will differ from those of previous 944 invasions. Invasion science must develop more detailed understanding of how international trade and travel are altered by national and international socioeconomic changes, and how these 945 946 changes in turn influence invasion trends (Hulme 2015). Such insights can greatly enhance the 947 development of scenarios and allow for improved risk categorization. A major priority for

invasion science is thus to advance beyond pattern recognition to embrace mechanistic socio-948 ecological models; for example, the Global Trade Analysis Project model was used to assess the 949 economic and trade impacts of required phytosanitary treatments of wood packaging (Strutt et al. 950 2013), and it was later applied to estimate the ultimate economic benefits of this policy (Leung et 951 al. 2014). An improved understanding of the links between global socio-economic trends and 952 invasions will ensure more effective targeting of national and international biosecurity efforts. 953 Such knowledge is also needed to inform the development of incentives and educational tools to 954 alter the behavior of importers, travelers, and others whose activities pose significant invasion 955 956 risks (Colton and Alpert 1998; Perry and Farmer 2011; Springborn et al. 2016).

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4.2. Global cooperation among national biosecurity programs

959 Most countries operate biosecurity programs that are designed to prevent the arrival, establishment, and spread of non-native species inside their national borders (Meyerson and 960 961 Reaser 2002; Hulme 2011). In some cases, unexpected prioritization of biosecurity measures can result from independent policy actions. For example, the EU Regulation (1143/2014) on non-962 963 native species has resulted in stakeholders focusing on biosecurity programs that limit the export 964 of live animals and plants but which neglects new introductions. Moreover, contemporary 965 national biosecurity programs are generally designed to protect the interests of individual 966 countries (Black and Bartlett 2020), with relatively little consideration given to the 'greater good' - i.e., protecting all nations from invasions. The mission of most national plant protection 967 968 organizations, for example, includes regulating imports that pose high risks of harmful introductions, while simultaneously promoting exports from their own countries; scant attention 969 is given to minimizing risks associated with such exports (MacLeod et al. 2010). Cooperation is 970

urgently needed among countries to craft biosecurity programs that are more cost-effective than 971 those where countries act largely in isolation (Latombe et al. 2017). Despite long-standing calls 972 for a binding internationally convention on invasive alien species (Perrings et al. 2010; Stoett 973 2010), there has been no progress towards this goal in over a decade. Within individual 974 countries, there are often political and economic obstacles to adopting truly cooperative 975 976 biosecurity. Thus, rather than a top-down multilateral approach to regulation, it is likely that closer integration of national biosecurity strategies will occur through a coalition of the willing. 977 A fine example is the Consultative Group on Biosecurity Cooperation established by Australian 978 979 and New Zealand ministers in 1999 under the terms of the Australia -New Zealand Closer Economic Relations Trade Agreement. This group has led efforts to harmonize animal and plant 980 health measures affecting trade between the two countries as well as coordinating biosecurity 981 responses. The Plant Health Quadrilaterals is a strategic coalition composed of the national plant 982 protection organizations of Australia, Canada, New Zealand, and the United States that enables 983 the respective plant health and biosecurity officials to address plant health and biosecurity issues, 984 particularly as they affect international trade of plants, plant products, and other regulated 985 articles. In 2016, a similar quadrilateral group involving the same four nations was established to 986 987 coordinate efforts to address marine biosecurity. These coalitions do not have any regulatory power, but through dialogue and cooperation they can address emerging issues in biosecurity in 988 989 an open and collaborative manner. For multilateral initiatives, a useful model for research on 990 cooperative biosecurity would be studies on cooperation between different countries to optimize harvest from shared fisheries (Bailey et al. 2010). These studies apply game theory, which could 991 992 also be applied to biological invasions to explore how cooperative biosecurity might yield higher 993 benefits to all countries by collectively reducing the flow of species globally, rather than just

994	preventing invasions at the national level. Lampert (2020) applied a dynamic game-theoretic
995	model to identify a Nash equilibrium corresponding to optimal contributions that various
996	countries or other entities could adopt for managing invading species with shared adverse
997	impacts. This approach could be expanded to explore cooperation by countries to fund various
998	pre- and post-border biosecurity activities.
999	
1000	4.3. The role of the bridgehead effect in managing invasions
1001	A key consideration driving the need for internationally cooperative biosecurity strategies
1002	is the tendency of organisms to invade new regions from locations where they have already
1003	established, a phenomenon referred to as the bridgehead effect (Lombaert et al. 2010;
1004	Bertelsmeier and Keller 2018) or 'hub-and-spoke' invasion topology (Carlton 1996). This
1005	phenomenon has been documented in historical global patterns of invasions for several plant and
1006	animal species (e.g., Bertelsmeier et al. 2018; Correa et al. 2019; Javal et al. 2019). The term was
1007	first coined by Lombaert et al. (2010), who used molecular analyses of the global spread of the
1008	harlequin beetle Harmonia axyridis and found that even though the species is native to east Asia,
1009	its invasions of Europe, Africa, South America and western North America all originated from
1010	eastern North America (Figure 5). Evidence exists that invasions from bridgehead regions may
1011	be promoted by genetic changes, demographics, or simply by the topologies of trade networks
1012	(Bertelsmeier and Keller 2018). More work is needed on the drivers of bridgehead dynamics to
1013	determine whether management-relevant generalizations exist.
1014	From the perspective of designing biosecurity programs, an important implication of such
1015	bridgehead dynamics is that benefits will accrue from preventing a species from establishing
1016	within a hub or bridgehead region – that is, an invaded location from which spread to other

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regions is more easily facilitated. Furthermore, cooperation among countries to eradicate or
control species established in bridgehead regions could yield greater benefit than attempts by
individual countries to exclude these species from arriving and establishing.

The current unilateral approach that dominates national biosecurity has roots in the close 1020 relationship between trade and import quarantines; quarantine is an effective and important tool 1021 1022 for excluding arrivals of new species, but there is a history of quarantine being abused to justify protectionist trade policies (Castonguay 2010). For example, the World Trade Organization 1023 (WTO) recognizes the International Plant Protection Convention (IPPC) as the authority for 1024 1025 setting standards for plant quarantine, and the WTO uses its Appellate Body for settling quarantine-related trade disputes. However, while the IPPC identifies quarantine practices and 1026 harmonized standards that individual countries should follow, it generally does not implement 1027 actions to minimize the movement of species worldwide; however, the IPPC has developed a 1028 National Phytosanitary Capacity Development Strategy that facilitates investment by member 1029 1030 countries in the development of biosecurity capacities in economically under-developed countries. Interdisciplinary research between invasion scientists and international trade 1031 economists is required to develop frameworks and justifications for globally collaborative 1032 1033 biosecurity efforts (Horan and Lupi 2005). Among the topics this research could address is how 1034 countries with varying economic resources can share resources for preventing the global 1035 movement of non-native species (Early et al. 2016).

1036 This research could also focus on developing strategies to identify bridgehead regions 1037 and initiate cooperative biosecurity negotiations with governments responsible for such regions. 1038 Border inspection data provide information on the identity of the geographical sources of species 1039 arriving at ports and are thus valuable resources for identifying bridgehead regions (Bertelsmeier

et al. 2018). Biosecurity agencies often consider inspection data as confidential (because of their 1040 possible significance in trade dispute litigation). Given the potential value of such data for 1041 identifying and delimiting bridgehead regions and global invasion risk (Turner et al. 2020), a 1042 challenge for invasion science is to ensure that such data are made more widely available and in 1043 a timely way to prevent regions that have received an invasive species from serving as sources 1044 1045 for new invasions even before the bridgehead population has been discovered. Ultimately, such data sharing could help inform biosecurity practices in individual countries, thereby reducing 1046 risks of future invasions. The world has recently witnessed an unprecedented case of 1047 1048 international sharing of spatiotemporal spread data for SARS-CoV-2 from its earliest stages, which should serve as an example for tracking other invasive organisms (Bertelsmeier and Ollier 1049 1050 2020).

1051

1052 4.4. Managing conflicting interests in biosecurity

1053 A related problem is that of cooperative approaches to transboundary biosecurity. The establishment of non-native populations can span regions managed for varying purposes, often 1054 with conflicting priorities (Epanchin-Niell et al. 2010). Conflicts of interest frequently bedevil 1055 1056 attempts to manage non-native species, especially when the focal species is simultaneously perceived as both beneficial and harmful by different sectors of society or in different areas of 1057 1058 the landscape (e.g., van Wilgen and Richardson 2014). For example, the ornamental horticulture 1059 industry benefits from importing and propagating non-native plant species while its actions conflict with other societal segments (e.g., ranchers, farmers, conservation managers) who suffer 1060 1061 from the impacts of plant invasions (Niemiera and Von Holle 2009). Invasion scientists must 1062 collaborate with economists and other researchers to devise approaches to engender cooperation

among stakeholders who are differentially impacted by the same non-native species and to 1063 explore how to optimize diverse management interests. In responding to changing perceptions of 1064 non-native species, their impacts, and their value to society, invasion science is facing challenges 1065 similar to those confronting other disciplines including the medical profession with regard to 1066 how best to communicate information about risk (Alaszewski and Horlick-Jones 2003). Social 1067 1068 science research must also develop effective strategies or models for systematic engagement of stakeholders seeking sustainable solutions to invasions (Shackleton et al. 2019). 1069 Conflicting interests among stakeholders that affect management of invasions sometimes 1070 1071 manifest as 'wicked problems'. These are characterized by diverse, opposing perspectives, objectives, and management goals that make them almost impossible to characterize or frame, let 1072 alone resolve, to the satisfaction of all stakeholders (Woodford et al. 2016). Woodford et al. 1073 1074 (2016) suggest that systematic framing of 'wickedness' by mediators can lead to negotiated solutions – either by reaching agreement on the dimensions and implications of unavoidable 1075 conflicts, or by circumventing the conflict by seeking alternative management perspectives. To 1076 this end, Novoa et al. (2018) developed a 12-step process designed to place stakeholders at the 1077 center of the development and implementation of decisions relating to conflicts of interest in 1078 1079 invasive species management. Fundamental requirements for achieving such aims are to 1) 1080 ensure that decisions and management actions are co-designed, co-produced, and co-1081 implemented to promote social learning and provide feedback to stakeholders, and 2) increase 1082 levels of collaboration and partnerships beyond the natural sciences and academia (Shackleton et al. 2019). Further work is clearly needed to achieve integration of broad stakeholder engagement 1083 1084 and co-operation in invasion research and management. Opportunities abound to apply existing

economic theory on governing common-pool goods (Ostrom 2010) to solve problems related to
the increasingly complex conflicts between stakeholders relating to invasive non-native species.

1088 **Conclusions**

1089 Invasion science is an increasingly interdisciplinary field that addresses questions and 1090 hypotheses of fundamental and applied importance to ecology, conservation biology, ecosystem management and restoration, and biosecurity (Ricciardi et al. 2017; Pyšek et al. 2020). We have 1091 identified four overarching issues that are critically important for the field to further adapt to 1092 1093 societal demands in the face of rapid global change. Reflected in these issues are burgeoning 1094 challenges posed by new sources and pathways (e.g. evolving trade routes and transportation systems) of invaders. Understanding and predicting invasions and their consequences are 1095 scientific endeavors, whereas managing them successfully largely rests with society: the former 1096 1097 informs the latter, and both tasks are complicated by context-dependencies that are becoming increasingly significant as rapid environmental change ensues. 1098

1099 Solutions to these challenges require innovations in theory and methods that potentially could be found through linkages with other disciplines. For example, factors promoting the 1100 1101 emergence and spread of novel infectious disease could be better understood and managed through collaborative research involving medical science and invasion science, to the benefit of 1102 both fields (Nuñez et al. 2020). In addition, within the broad discipline of ecology there are 1103 1104 disparate concepts and methods that have not yet been well integrated into invasion science (e.g. 1105 species interaction networks; Hui and Richardson 2019), or that are only now becoming broadly applied (e.g. the use of functional response metrics in risk assessment; Dick et al. 2017a, b; 1106 Dickey et al. 2020). 1107

New approaches are needed to forecast candidate invaders, probable invasion success, 1108 and consequent invader impacts under future terrestrial, freshwater, and marine conditions that 1109 have little or no analogue reference point in the past. A key growth point for the field would be 1110 to develop a better understanding of temporal invasion dynamics, including invasion debt and 1111 time lags. The concept of invasion debt (Essl et al. 2011; Rouget et al. 2016), in which invasions 1112 1113 are the end result of processes currently at play (e.g., increasing propagule transport and introduction in the face of reduced environmental resistance) is analogous to the emergence of 1114 disease symptoms following viral or bacterial exposure resulting from lapses in hygienic 1115 measures or failed social behaviors. A more predictive understanding of invader impact could be 1116 advanced, in part, through research on interacting and cumulative time lags in biodiversity and 1117 ecosystem responses to invasions (Essl et al. 2015b, c). 1118 Fundamental taxonomic skills are essential for biosecurity and a deeper understanding of 1119 biogeography and evolutionary history – the foundations of invasion science. The application of 1120 1121 invasion science to early detection is compromised without expertise suitable to identify nonnative species rapidly. Misidentifications have and will lead to spurious conclusions in 1122 ecological studies and, ultimately, to inappropriate and ineffective management and policy, when 1123 1124 such are called for. The necessary expertise could be cultivated through application and enhancement of infrastructure support (e.g. cyber-tools, specimen collections linked with 1125 1126 permanent custodial care), and re-establishment of training of both classic and advanced 1127 taxonomic skills in biology programs.

Finally, invasion science must address transcultural sociopolitical challenges including how best to communicate information and uncertainty about risk, how to engage diverse stakeholders who are differentially impacted by the same non-native species, and how to inform

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transboundary biosecurity policies. There is still much work required to harmonize the definition 1131 and application of biosecurity policies across different multilateral organisations such as the 1132 Convention on Biological Diversity, the International Plant Protection Convention and the World 1133 Organisation for Animal Health. Invasion science must continue to inform the rapidly evolving 1134 landscape of international biosecurity agreements designed to control pathways that create 1135 1136 bridgehead populations, which can drive widespread invasions. International data-sharing will be needed to reduce invasion risk at regional and global scales. The remarkable example of the 1137 rapid cooperative sharing by most countries of spatiotemporal spread data for SARS-CoV-2 from 1138 its earliest stages should inspire global efforts to track other invasive organisms. 1139

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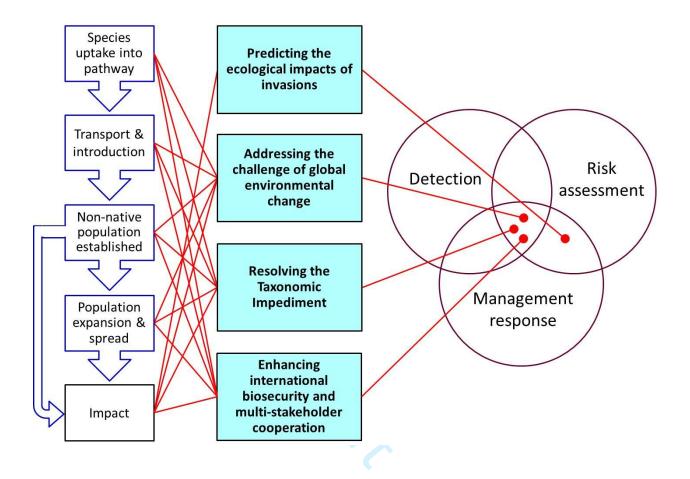


Figure 1. Four priority issues (center column) that must be addressed by invasion science to meet burgeoning challenges in an era of rapid environmental change. Through multiple connections, each issue is implicated in one or more stages of the invasion process (left column), as well as in the impact of the invader (which can occur at any stage from introduction to establishment to spread) and in the detection, risk assessment, and management response of invasion threats. For example, scientific understanding of the processes that control the diversity, abundance, distribution, and impacts of non-native species ultimately depends on the quality of taxonomic data; therefore, resolving the Taxonomic Impediment (the erosion of our capacity to recognize biodiversity and distinguish non-native from native species accurately) would enhance our ability to detect non-native species, assess their impacts, and respond to new invasion threats.

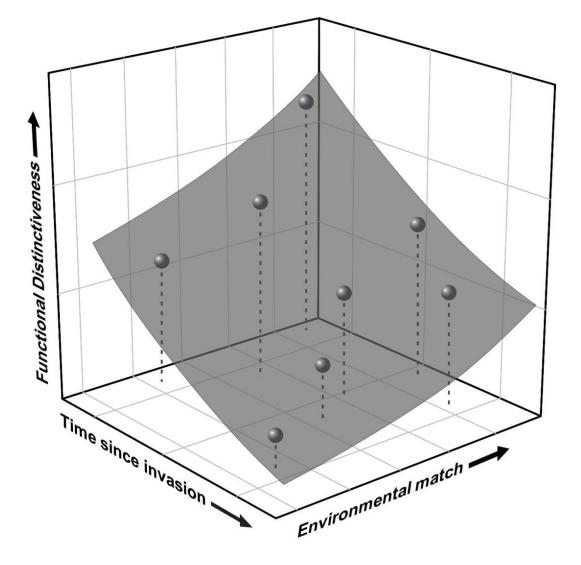


Figure 2. An example of integration of impact hypotheses. The 3-dimensional plot represents the predicted variation in an invader's ecological impact in relation to three factors, shown as axes: 1) the functional (or phylogenetic) distinctiveness of the invader among resident species; 2) the degree of environmental match – i.e., the inverse of the distance between mean abiotic conditions in the invaded environment and the invader's physiological optimum; and 3) time since invasion. Functionally novel invaders, especially those that exploit key resources, are

predicted to have greater impacts on the invaded ecosystem (Functional Distinctiveness Hypothesis). Invaders that are more physiologically matched to abiotic conditions in the invaded environment should have greater per capita effects (Environmental Matching Hypothesis). Further, in this example, impact is hypothesized to attenuate over time, based largely on the premise that given suitable time resident species (predators, prey, parasites, competitors) will adapt to the invader and dampen its influence. These factors are shown here to be mutually independent, but interactions are possible (e.g. physiological match may interact with time since invasion, owing to local adaptation or directional shifts in abiotic conditions).

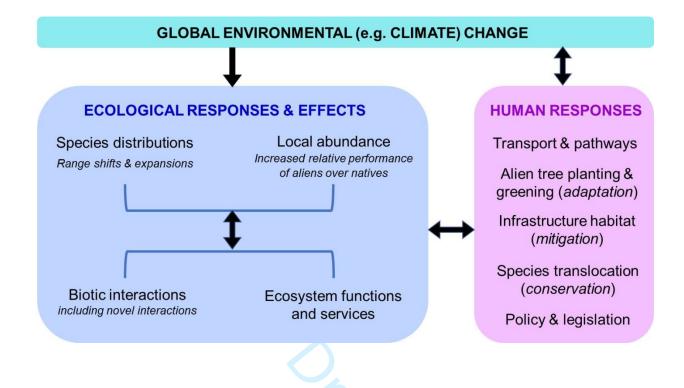


Figure 3. Global environmental change (in particular, climate change) directly and indirectly elicits ecological and human responses that promote invasions. Environmental change can trigger shifts in the distributions and abundances of native and non-native species, leading to novel biotic interactions and altered ecosystem functions and services, which can themselves prompt further ecological responses. Human responses include climate change adaptation and mitigation, as well as species conservation; many of the current human responses will likely facilitate invasions. These ecological and human responses also affect each other, compounding the direct impacts of environmental change.

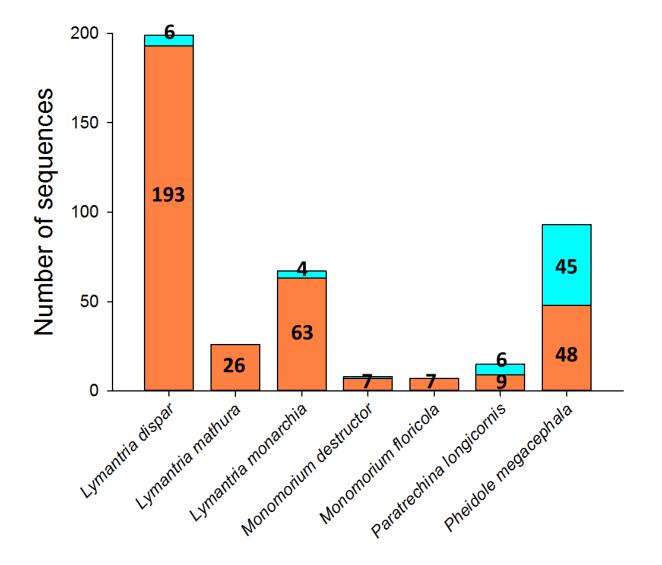


Figure 4. Examples of invasive insect species for which a discrepancy exists between the number of sequences available in GenBank v3.0 when using the two primary search query tools they provide: a taxonomy-based search of GenBank records (green) and a broader search using sequences or taxonomy of other publicly available data sources linked to GenBank (orange). Such discrepancies in search results across databases increase the risk that these species will be incorrectly classed as 'unidentified' when metabarcoding approaches are used to identify non-native insects. Data from Boykin et al. (2012).

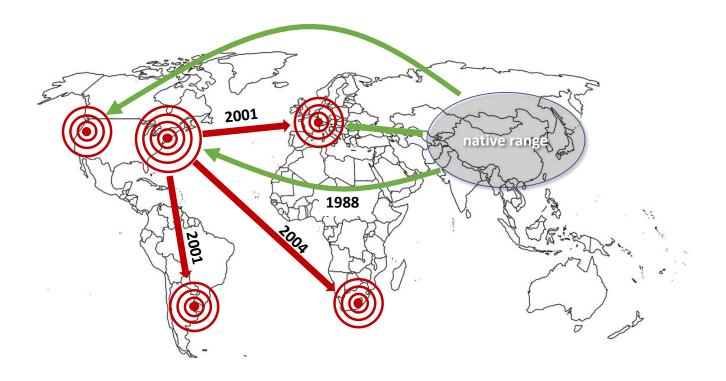


Figure 5. The Bridgehead Effect illustrated by the global spread of the Harlequin ladybird beetle *Harmonia axyridis*, based on genetic analyses by Lombaert et al. (2010). Intentional biocontrol introductions are shown in green, whereas accidental invasions are shown in red. In this example, most of the global spread of this species has originated from non-native populations established in Eastern North America, which has functioned as a bridgehead region (adapted from Lombaert et al. 2010).