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Introduced parasites in food-webs: new species, shifting structures?
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Abstract

Introduction of free-living species also result in co-introduction of their parasites. Since recent advances have shown that native parasites dramatically alter food web structure, I evaluate here how introduced parasites might reorganise food webs. Empirical evidence suggests that introduced parasites alter food webs qualitatively through topological changes and quantitatively through shifts in trophic relationships arising from modified host phenotypic traits. I argue that predicting the extent of food web reorganisation is, however, difficult due to underlying ecological and evolutionary processes that could provide contrasting food web outcomes, including enemy release, biotic resistance and parasite spillover and spillback. Nevertheless, I suggest these food web reorganisations represent a further aspect of human-mediated global change resulting in irreversible consequences across multiple trophic levels.

Introduced species and their parasites

Introduced species have adverse consequences for native biodiversity and raise global concerns over biotic homogenization [1-3]. Introductions of free-living species can also result in the co-introduction of their parasites [3,4]. Although the introduction process might filter out many of these [5], the consequences in the receiving ecosystem of those parasites that are co-introduced vary according to a number of factors including the complexity of their lifecycle, their ability to spillover to native species, and the resistance and tolerance of these new hosts to infection [5-7]. Although high mortality rates might be incurred, these tend to be a consequence of the emergence of an infectious disease [8-10] or be symptomatic of additional underlying stresses, such as poor environmental conditions [11]. Sub-lethal host consequences can include pathological, physiological and/or behavioural changes, with likely adverse consequences for growth, survival and fitness [7,11].

In considering the consequences of infection by non-native parasites, information from native parasites can be very revealing. Native parasites can profoundly shape the dynamics of their host populations and communities, alter competition strength and influence trophic relationships, and are important drivers of biodiversity [12-16]. Although their inclusion in food web studies has tended to be overlooked [17,18], recent studies suggest this has lead to dramatic underestimates of food web connectivity and complexity [19,20]. This is because these studies on 'infected food webs' have revealed parasites alter food web structure and stability through, for example, substantially increasing connectivity, nestedness and linkage density [18-29]. As the consequences of infection by native parasites [13-15,24] are relatively similar to those of introduced parasites [7,11,24,30,31] and given the dramatic changes in structure and complexity that occur when native parasites are included in food webs [18-29], this raises the question of how introduced parasites might influence food web structures. Here, I explore this question through: (i) identifying how introductions of free living species and their parasites could influence food web topology; (ii) examining how parasite infections might influence introduction outcomes and food web topology; (iii) examining the processes and implications of parasite spillback and spillover between native and introduced species; and (iv) determining how parasite lifecycles and host species' characteristics influence food web structure. Case studies highlight relevant examples and opportunities for further research (Boxes 1-3). The influences of introduced parasites on food web structure are considered through qualitative changes, such as topology [21,27], and through more quantitative perspectives, such as the magnitude of shifts in the intra- and inter-specific trophic interactions that result from host phenotypic modifications and might disrupt patterns of energy flow [6-8,24]. For simplicity and as pathogens are also parasitic by nature, parasites are defined here as organisms that feed on a single host individual per life history stage [32].

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Introduced species influence food web topology

The introduction of a free-living species into a food web should increase food web diversity and cause some topological reorganisation through the formation of a new node that establishes new links [33]. Although the co-introduction of their parasitic fauna (both meiroand macro-parasites) should provide further new nodes and links [4], introduced species often bring a relatively low number of parasites into the recipient food web (Table 1). This is because they tend to lose their parasites during the introduction process [34] through such factors as only a small sub-set of propagules being introduced from the native range that might be uninfected or comprise uninfected life-stages; the parasite having a complex lifecycle that requires multiple hosts with some of these missing in the receiving food web; and strong selective pressures, founder events and population bottlenecks in both parasites and hosts that lead to their early extirpation [5,7,35]. Nevertheless, some parasites will be cointroduced [36], with Torchin et al. [5] suggesting that where an introduced species in their native range might have a mean of 16 parasites, three will remain in their introduced range. Introduced species do, however, gain an additional four native parasites through parasite acquisition during the introduction process [5,6]. Thus, while the number of new nodes resulting from co-introduced parasites might be low those introduced will still result in some shifts in food web topology (Table 1).

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Parasites with complex lifecycles that have infective, free-living lifestages can also be introduced in the absence of their non-native host species. For example, the nematode parasite *Anguillicoloides crassus* is native to the Japanese eel *Anguilla japonicus* but as a consequence of movements in the global aquaculture trade has spilled-over into the European eel *Anguilla anguilla* and is now widely distributed in their range [36,37]. The initial introduction of *A. crassus* into rivers in the UK was through water discharges from

aquaculture transport lorries that released only their eggs and juveniles into adjoining water courses [38]. These infective free-living lifestages then completed their lifecycle through infecting native crustacean intermediate hosts, with subsequent transmission to fish paratenic hosts and *A. anguilla* [38]. For food web topology, this would also have provided a number of new nodes and multiple new links across a range of trophic levels.

Parasites influence introduction outcomes that affect food web topology

The outcome of the introduction of a free-living species will strongly influence how their cointroduced parasites will subsequently affect food web topology. The establishment and subsequent invasion of the free-living species might be enhanced through enemy-release or inhibited by biotic resistance.

The enemy release hypothesis (ERH) relates to the parasite loss experienced by introduced species that was outlined in the previous section and predicts that the loss of their natural 'enemies' enhances their ability to establish and invade [24,39,40]. Indeed, with introduced species escaping at least 75 % of their parasites from their native range [32], there are likely to be substantial benefits in terms of their fitness and survival [5]. Despite the ERH having been used to help explain the invasion success of species as diverse as slugs [41], mosquitoes [42] and frogs [43], evidence is not unequivocal. This is because whilst introduced species can experience enemy release, they might also incur significantly higher levels of prevalence than in their native range and this could be detrimental to their fitness, impinge on their ability to invade and diminish their functional role in the receiving ecosystem [5,7]. Further, Colautti *et al.* [40] suggested many studies over-estimate the role of ERH in successful invasions as few experimentally test the differential effects of enemy release versus alternative factors.

By contrast, biotic resistance inhibits the survival, establishment and invasion of introduced species through, for example, the presence of native predators, strong competitors and/or parasites that impede their survival and reproduction (Table 1) [44]. For example, whilst introduced bivalves in the North Sea, such as *Crassostrea gigas* and *Ensis americanus* were believed to be free of parasites, providing competitive advantages over native bivalves, up to 80 % were found to be infected with native trematode parasites that were providing some natural biotic resistance against their colonisation [45].

The contrasting outcomes of enemy release and biotic resistance for introduced species and their parasites are important in determining the consequences for food web structure. Should invasive populations of free-living species develop, they can reach higher densities than both populations in their native range and functionally-similar native free-living species in the receiving ecosystem [7]. This might result in the displacement of native species, particularly those that exploit similar resources [41]. This is important as the parasite diversity of the invasive species can be substantially reduced compared to the displaced native species [32]. For example, Torchin et al. [46] revealed a native snail with 10 native trematode parasites in an aquatic food web was displaced by a functionally similar invasive snail with only one trematode; similar findings are apparent in Northern Ireland for the parasites of native and invasive amphipods [47]. Thus, the increased number of nodes and links in the food web formed by the invader and its parasites in food web topology might fail to compensate those lost through displacement. Similarly, Lafferty et al. [21] revealed snails in a coastal salt marsh were infected with up to 17 host-specific parasites, thus any population displacement by an invasive snail would be likely to result in substantial losses in food web nodes and links, and decreased robustness through increased secondary extinctions.

Implications of parasite 'spillback' and 'spillover' for food web structure

The shifts in food web topology resulting from co-introduced parasites will be strongly influenced by three processes: (i) enemy release; (ii) parasite introduction and spillover; and (iii) parasite acquisition and spillback (Table 1) [6]. The role of enemy release in determining the actual number of parasites co-introduced into the food web has already been outlined [5]. Following their introduction, these parasites might now 'spillover' to native species, i.e. they 'host-switch' to native species [6,7] and so would represent a new consumer in the ecosystem that increases the number of food web links (Fig. 1). The most substantial shifts in food web topology are likely to result from those spilled-over parasites with complex lifecycles that are trophically transmitted and have intermediate hosts, as their lifestages will form a series of new nodes and links across multiple trophic levels (Fig. 1) [48].

Parasite acquisition occurs when introduced free-living species become infected by native parasites; in the topological food web, new links are thus formed that might increase connectance and nestedness. The process might also have implications for the quantitative food web, as acquisition might result in parasite 'spillback' to the native species and disrupt trophic interactions [6]. This is dependent on whether the introduced host is competent [6,49]. If it is, then its population can act as a 'reservoir' in which the parasite persists and reproduces, and from which its infective stages disperse and result in increased parasite prevalence in native hosts [6]. Conversely, if the introduced host species is not competent then it can act as an infection 'sink' that dilutes infection levels in native hosts [6,7]. Some introduced hosts actually incur higher infection levels than native hosts, as observed in introduced European starlings in the USA that acted as a reservoir for equine encephalitis virus [49]. Across these reservoir and sink scenarios, considerable alterations in the trophic interactions are thus likely between the native and introduced hosts and parasites as infection

modifies the phenotypic traits of infected individuals, causing shifts and reorganisation in the quantitative food web [50].

Shifts in food web structure are affected by parasite lifecycles and host characteristics

The parasite lifecycle, host phenotypic modifications and confounding infections by native parasites are additional factors that determine how introduced parasites might alter food web structure, particularly with regard to quantitative changes (Table 1). Modifications to the host phenotype will vary depending on the hosts and the parasite concerned, but generally include altered foraging behaviour, feeding rates, competitive relationships, and shifts in life history traits such as altered growth rates and reproductive traits [13]. These modifications to host phenotype might impact trophic interactions through shifts in intra- and inter-specific competition and trophic relationships, and potentially result in considerable alterations to energy flow through the food web.

Direct and complex parasite lifecycles

In the topological food web, an introduced parasite with a direct lifecycle and a single definitive host might result in only one new node and link, i.e. there would be little reorganisation. By contrast, the spillover to native hosts of an introduced parasite with a complex lifecycle involving trophic transmission could result in a number of new nodes and multiple new links that causes substantial reorganisation, with this supported by some empirical evidence from introduced parasites (Box 1) and supplementary examples from native parasites [12,20,24]. In the quantitative food web, parasite lifecycles are important regarding how the host phenotypes might be altered and shift trophic relationships. As native parasites with complex lifecycles often increase their likelihood of transmission by manipulating host anti-predator behaviours that increase the likelihood of predation and so

completion of the parasite lifecycle [13], then similar scenarios could be likely for spilledover parasites in native hosts and acquired native parasites in introduced hosts. Notwithstanding, Lagrue *et al.* [51] found that while the acanthocephalan parasite *Pomphorhynchus laevis* manipulated the drifting behaviour of its native intermediate hosts (e.g. *Gammarus pulex*), this was not apparent in an introduced intermediate host (*Gammarus roeseli*).

Influence of host characteristics on food web consequences

Where an introduced parasite spills over into a native species, the potential shifts in food web structure depend on a range of factors relating to the host species, including their functional role(s), affected life stage, modified phenotypic trait(s), genetic diversity and parasite resistance and tolerance [32,52-54]. Resistance and tolerance of native hosts to the parasite might be low due to poor immune responses and anti-predator behaviours resulting from their lack of shared evolutionary history [55]. A spilled-over parasite that meets low resistance and tolerance in native hosts can lead to high mortality rates, i.e. there will be an epizootic that could have substantial implications for food web structure (Box 2). Note, however, that the level of parasite resistance and tolerance in naïve hosts will depend on a wide range of environmental and biological factors, including host genetic diversity [5,8]. Moreover, rapid evolutionary responses over two or three generations have been recorded in host populations following disease emergence that have provided enhanced immune responses to infection and so minimised the pathology [8] and consequently the impacts for food web structure.

The indigenous parasites of the native hosts can also represent confounding infections that add complexity in determining which parasites - native or introduced - are actually adapting the host phenotype [13]. This is compounded by native parasite infections also leading to

reciprocal effects between the host and parasite [56]. These effects include 'vicious circles' of infection whereby hosts become more vulnerable to subsequent infections [57], leading to further complexity in deciphering which parasites are modifying the host phenotype.

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Concluding remarks and research perspectives

Consequences for the topological food web of an introduced free-living species and its parasites are associated with the addition of new nodes and links across different trophic levels that should increase connectivity and complexity. For the quantitative food web, shifts in trophic interactions and patterns of energy flow might be caused by cascading effects arising from the modified phenotypic traits of the native hosts (e.g. through parasite spillover, Fig. 1, Table 1) and introduced hosts (e.g. through parasite acquisition). Empirical evidence for these food web shifts supports the opinion I have expressed here that there is considerable potential for introduced parasites to substantially alter native food web structure (Boxes 1-3). There are, however, a series of factors, such as enemy release and biotic resistance, which strongly influence how these alterations will be manifest (Table 1). Nevertheless, it should be noted that these alterations are resulting from an aspect of human-mediated global change that is usually irreversible and often associated with exploitation of ecosystem services and their management (Box 1) [58]. Thus, from a management perspective, this emphasises the requirement for risk-based regulations and policies to be implemented on the global movements of free-living species that minimises opportunities for inadvertent parasite cointroduction [1,2].

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Research perspectives require the further testing of hypotheses relating to the potential shifts in food web structure that arise from introduced parasites of different functional groups and with contrasting lifecycles, and in relation to the factors outlined in Table 1. Approaches

can integrate the qualitative food web approaches that have dominated native parasite food web research with stable isotope analyses that can help reveal the more quantitative perspectives, including parasite-host trophic relationships [27], the effect of parasite loading on trophic niche [55] and the modifications to energy flow between trophic levels (Box 1). Evolutionary perspectives can be informed by the potential development of rapid and inheritable acquired immune responses of naïve hosts to introduced parasites [8]. The outputs of this research will then be important in refining the underlying theory that bridges the fields of parasitology, invasion ecology and evolution (Table 1). Aspects of these research perspectives are likely to be already progressing, albeit indirectly, through the use of introduced parasites in classical biological control programmes (CBC) that represent large-scale field experiments on introduced parasites in food webs (Box 3). That the parasite's natural host is already present and invasive in the food web increases their probability of establishment and so their utility to this field.

In closing, I emphasise that the increasing rate of species introductions is accelerating the rate of non-native parasite introductions and their consequences in the receiving ecosystems depends on their host specificity. Should parasite spillover occur then the native components of the food web are likely to be altered. If the parasite is host-specific then only impact the invaded component of the food web would be affected, potentially diminishing its effect. Indeed, in these circumstances, the parasite may protect the native aspects of the food web through diminishing the interaction strength of the invader. Irrespective, the increasing presence of non-native species and their parasitic fauna in food webs might be resulting in their irreversible structural re-organisations and represents a further consequence of global change.

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Table 1. Factors affecting the structural consequences of introduced parasites in food webs and examples of their potential outcomes

Factor	Implication	Potential outcome for food web structure
Lifecycle of the	(i) Parasites with complex lifecycles require definitive intermediate	No change in food web structure.
introduced parasite	and final hosts absent in the new range and fail to establish	
	(ii) Parasites with complex lifecycles might spillover at their	Shifts in the topological food web through new links and the
	different lifestages to native hosts at different trophic levels	quantitative food web via host phenotypic modifications.
	(iii) Parasites with direct lifecycles infect only the introduced	Minor changes in food web topology.
	species	
Enemy release	The introduced free-living species hosts a reduced number of	Minor changes in food web topology.
	parasites than in their native range.	
Biotic resistance	The introduced host and their parasites fail to establish as they are	No change in food web structure.
	out-competed, predated and/or parasitized by native species.	
Parasite spillover	Co-introduced parasites are transmitted to native species through	Shifts in the topological food web through new links and the
	host-switching.	quantitative food web via host phenotypic modifications.

	Parasite spillback	Native parasites infect the introduced free-living species that:	Shifts in the topological food web through new links and the
		(i) become a competent host that acts as infection reservoirs for	quantitative food web via host phenotypic modifications in native
		native hosts; or	and/or introduced hosts.
		(ii) become an incompetent host acting as an infection sink.	
-	Naïve hosts rapidly	Host populations suffering an epizootic due to initial low parasite	Decreased host populations that substantially reorganise food web
	develop inheritable	resistance and tolerance acquire inheritable immunity and anti-	topology and disrupt trophic relationships are temporary.
	anti-parasite traits	parasite behaviours in two to three generations	
-	Co- infections of	Hosts with modified phenotypic traits are infected with a number of	Shifts in the quantitative food web might relate to infections by
	native and	native and introduced parasites.	native and/or introduced parasites.
	introduced parasites		

- 412 Glossary
- 413 **Competent host:** A host species that is capable of supporting and transmitting the parasite.
- 414 **Connectance:** A food web metric expressing the proportion of possible links in a food web
- 415 that are realized.
- 416 Complex lifecycle: A parasite lifecycle that requires more than one host species for
- completion through use of intermediate and final hosts.
- 418 **Direct lifecycle:** A parasite lifecycle involving a single host species.
- 419 Food web topology: A qualitative representation of feeding interactions that illustrates who
- 420 eats who in the food web, but with no information on the strength of those feeding
- 421 interactions.
- 422 Macro-parasite: Defined here as relatively large parasites such as cestode tapeworms; can
- be seen with the naked eye.
- 424 Micro-parasite: Defined here as parasites requiring microscopy to view, such as viruses,
- 425 bacteria and fungi.
- Nestedness: An expression of the extent to which species with a low number of food web
- links have a sub-set of the links of other species, and is determined by characteristics such as
- 428 network size and connectivity.
- 429 Parasite spillover: The cross-species transmission of a non-native parasite from its
- introduced free-living host to a native free-living host.
- 431 **Parasite spillback:** The cross-species transmission of a non-native parasite from its acquired
- ative free-living host via spillover back to its introduced free-living host.
- Paratenic host: A host that maintains the lifecycle of the parasite but is not needs for its
- 434 development.
- 435 Quantitative food web: a representation of feeding interactions within an ecosystem that
- illustrates both the presence and strength of feeding interactions within that ecosystem.

Qualitative food web: a representation of the feeding interactions within an ecosystem that
illustrates who eats whom but provides no information about the strength of the feeding
interactions.
Robustness: Refers to the probability of secondary extinctions arising from species'
removal; often defined as the proportion of species that must be removed to result in 50 % of
species going extinct as they lack resource species. It is a topological measure of stability.
Stability: The likelihood of interacting species within the food web being persistent and
usually expressed as a continuous metric that measures this likelihood. A stable food web
tends to be one with minimal secondary extinctions following species removal.

Box 1. Alterations to food-web structure by introduced parasites

Direct empirical evidence for shifts in food web topology arising from the introduction of free living species with their parasites is provided by invasive fishes in the pelagic food web of Lake Takvatn, Norway [4]. Introductions into this sub-Arctic lake of Arctic charr *Salvelinus alpinus* and three-spined stickleback *Gasterosteus aculatus* and their cointroduced parasites strongly altered pelagic food web structure through increasing: (i) species richness from 39 to 50 species (the two fishes plus nine parasites); (ii) the number of nodes and trophic links in the topological food web; (iii) food-chain length; and (iv) the total number of trophic levels in the food web [4]. Food web complexity also increased, revealed through increased linkage density, degree distribution, vulnerability to natural enemies, omnivory and nestedness, all of which might have consequences for network functioning and stability [4]. The study concluded that when parasites are co-introduced with their free-living hosts, substantial alterations in the structure of the qualitative food web can result, especially when the parasites are tropically transmitted with complex lifecycles that form new links across multiple trophic levels. It thus underpins the importance of accounting for both native and introduced hosts and parasites in food-web studies.

That this study highlighted that trophically transmitted parasites with complex lifecycles will cause the most substantial shifts in food web structure suggests that globally invasive parasites with complex lifecycles, such as the Asian tapeworm *Bothriocephalus acheiolognathi*, might have already resulted in major reorganisations in food web structure. This trophically-transmitted freshwater parasite has been introduced around the world through the aquaculture industry [59]. In their invasive range, species within six copepod genera have been identified as intermediate hosts and at least 200 fish species as final hosts [59]. Transmission can also be through piscivory by fish and birds (postcyclic transmission)

[59]. Thus, as already measured in [4], their widespread introduction might have resulted in substantial shifts in food webs structure and complexity. Quantitative food web impacts might also be likely through their host impacts, as they cause substantial phenotypic modifications, albeit with a variable severity according to the fish host and infected lifestage [55]. In particular, infected *Cyprinus carpio* were discovered to be feeding at lower trophic levels than uninfected individuals, changing the symmetry of their competitive interactions [55] and potentially impacting trophic diversity and disrupting patterns of energy flow.

Box 2. Emerging infectious diseases and epizootics

Epizootics are often associated with emerging infectious diseases (EIDs) [8,9]. These include parasites which have recently increased in incidence, host species and/or geographic extent [8,9,54,58]. Their effects can be severe, mediating community dynamics, shrinking host ranges and potentially extirpating local populations [8,9,58]. Consequently, they could disrupt food web topology, potentially decrease robustness via secondary extinctions and cause shifts in trophic relationships [9,44,45]. Examples of introduced pathogens in UK food webs responsible for substantial declines of native species are crayfish plague Aphanomyces astaci impacting native crayfish Austopotamobius pallipes following spillover from introduced signal crayfish Pacifastacus leniusculus [60] and the parapoxvirus that spilledover from invasive grey squirrels Sciurus carolinensis to native reds Sciurus vulgaris [61]. The disease emergence associated with Gyrodactylus salaris in Norway dramatically reduced populations of Atlantic salmon Salmo salar in 45 rivers [62] and triggered large-scale eradication efforts [62]. This highlights that EIDs in economically important species are often highly managed [9] and although outbreaks can have substantial food web implications, structural changes might be temporary if their management enables population recovery (Table 1).

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Epizootics can also have profound effects on native ecological-engineering species, such as grazing animals that can have consequences for plant communities and food web structure. For example, EIDs impacting grazing animals can have substantial implications on the prevailing vegetation cover, as revealed by the cattle disease Rinderpest [63]. Savannah ecosystems comprise open grasslands, woodlands and closed thickets of broad-leaved shrubs; in the Serengeti National Park, Tanzania, the small tree *Euclea divinorum* facilitates establishment of the closed thickets [63]. Under current park management this, however, is

not occurring, with the existing thickets estimated as having established between 1890 and 1920 when their growth was enabled by a Rinderpest epizootic that extirpated the park's ungulates and severely reduced the cattle population [63]. Rinderpest has subsequently been successfully managed at a regional and global level, preventing further outbreaks and inhibiting further thicket establishment [64].

Control of engineering invasive species has utilised introduced pathogens to facilitate ecosystem restoration that might have invoked cascading food web effects. The *Myxoma* virus has been used widely to control invasive rabbit numbers and enabled severely grazed vegetation to recover in impacted food webs [65]. In the case of the sub-Antarctic Macquarie Island, *Myxoma* decreased rabbit numbers and enabled recovery of tall tussock grassland; when rabbit numbers subsequently recovered to former levels then a uniform pattern of degraded vegetation and increased bare ground returned [65].

Box 3. Classic biocontrol: field experiments on introduced parasites in food webs

As classic biological control (CBC) is the deliberate introduction of the natural enemies of an introduced species into their invaded region then aspects of their study have high utility in informing how non-native parasites may alter the structure of native food webs [66]. Relevant ecological theories and hypotheses to CBC include enemy release and parasite spillover. Indeed, ERH provides CBC with its underlying principles as it assumes the invasion success of the target species was the loss of its native enemies so that their introduction into the new range will control it [39,66,67].

CBC has been argued by practitioners as being the most environmentally desirable control method as it has limited ecological consequences beyond the target species. However, the traditional CBC approach, the release of generalist parasites to control the invader, suggests that substantial impacts have been incurred in native non-target species as a result of spillover, including local extinctions, [66,67]. For example, Hawkins and Marino [68] found that of 313 introduced parasitoids released in North America for CBC, 51 were present on non-target native insects. In field experiments in the USA on the parasitoid fly *Compsilura concinnata*, Boettner *et al.* [69] found high levels of parasitism (up to 100 %) in two native silk moths that could have been responsible for local extinctions. Henneman and Memmott [67] studied a remote and relatively pristine Hawaiian swamp within a region where at least 122 releases of parasitic wasps and flies have been released to control Lepidopteran agricultural pests [67]. They recovered 216 parasitoids from 58 native moth species of which 83 % were introduced through CBC and a further 14 % were from accidental introductions; only 3 % were native species [67]. Thus, introduced parasites from CBC had profoundly reorganised the trophic relationships in this quantitative food web.

These examples all suggest that the release of generalist, non-native parasites is likely to result in some degree of spillover to native species and alter food web structure. More contemporary applications of CBC has, however, started to test the use of specialist enemies that will not spillover and, if successful at controlling the target species, will diminish their interaction strength with native species [70]. By contrast to the release of generalist parasite, this aspect of CBC should consequently inform how host-specific introduced parasites might protect the native components of food webs from introduced free-living species [70].

Figure 1. (i) Graphical and (ii) matrices representing a simple five-node native food web [27]
before (a) and after (b) the introduction of a free-living non-native species, and before (c) and
after (d) the spillover of its parasite with a complex lifecycle to a native final host. Native
taxa are represented as basal (B), grazer (G1, G2) and predator (C1, C2) and the introduced
free-living non-native species as C_{INV} . In (c) and (d), the parasite introduced with C_{INV} is
represented as P; it has an adult stage (A) using (c) C_{INV} and (d) C_{INV} and C1 as a host, a free-
living larval stage (L1), and a parasitic larval stage (L2) that uses G2 as an intermediate host.
Transmission from intermediate host to final host requires the consumption of an infected
intermediate host.
In (i), the rectangular box at the top of (c) and (d) contains the three life stages of P, the
dashed ellipsoids indicate parasites occurring within hosts, and arrows represent feeding links
that also indicate the direction of energy flow (note the predator-parasite links are not shown
for brevity).
In (ii), the consumers are rows and resources are columns, and the shaded boxes indicate an
interaction. In (c) and (d) there are four quadrants (clockwise from the top left): predator-
prey, predator-parasite, parasite-parasite and parasite-host. In the initial free-living web (a),
20 % of the possible links (directed connectance) are present [27]; after the introduction of
the free-living species (b), this reduces to 19 %. The inclusion of the introduced parasite (c)
increases connectance to 26.5 % and parasite spillover to C1 increases it to 29 %.

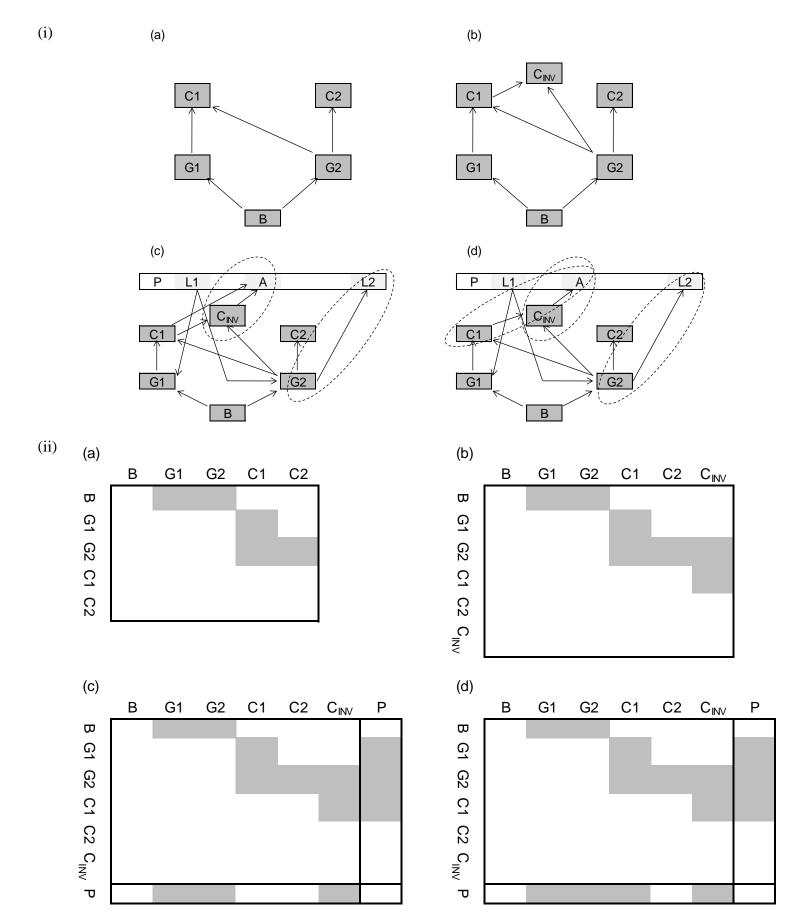


Figure 1.