Negative Niche Construction Favors the

Evolution of Cooperation

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$_{\scriptscriptstyle 11}$ Abstract

By benefitting others at a cost to themselves, cooperators face an ever present threat from defectors, or individuals that avail themselves of the cooperative benefit without contributing. A longstanding challenge to evolutionary biology is to understand the mechanisms that support the many instances of cooperation that nevertheless exist. Hammarlund et al. recently demonstrated that cooperation can persist by hitchhiking along with beneficial non-social adapta-17 tions. Importantly, cooperators play an active role in this process. In spatiallystructured environments, clustered cooperator populations reach greater densities, which creates more mutational opportunities to gain beneficial non-social adaptations. Cooperation rises in abundance by association with these adaptations. However once adaptive opportunities have been exhausted, the free ride ends as cooperators are displaced by equally-adapted defectors. Using an agent-based model, we demonstrate that the selective feedback that is created as populations alter their environments can maintain cooperation indefinitely. We show that cooperator success depends specifically on negative niche construction. Here, negative niche construction acts as a perpetual source of 27 adaptive opportunities. As populations adapt, they further alter their environment in ways that reveal additional opportunities for adaptation. Despite being independent of niche construction in our model, cooperation feeds this cycle. We show that by reaching larger densities, populations of cooperators are better able to adapt both to changing environments and to the constant threat posed by defectors. We relate these findings to previous studies from

- $_{34}$ the niche construction literature and discuss how this model could be extended
- $_{35}$ to provide a greater understanding of how cooperation evolves in the complex
- environments in which it is found.

37 Introduction

Cooperative behaviors are common across all branches of the tree of life. Insects divide labor within their colonies, plants and soil bacteria exchange essential nutrients, birds care for others' young, and the trillions of cells in the human body coordinate to provide vital functions. Each instance of cooperation presents an evolutionary challenge: How can individuals that sacrifice their own well-being to help others avoid subversion by those that do not? Over time, we would expect these defectors to rise in abundance at the expense of others, eventually driving cooperators—and perhaps the entire population—to extinction. Several factors can prevent this tragedy of the commons (Hamilton, 1964; Nowak, 2006; West et al., 2007b). One such factor involves non-random social interaction, in which cooperators benefit more from the cooperative act than defectors. This can occur when cooperators are clustered together in spatially-structured populations (Fletcher and Doebeli, 2009; Nadell et al., 2010; Kuzdzal-Fick et al., 2011) or when cooperators use communication (Brown and Johnstone, 2001; Darch et al., 2012) or other cues (Sinervo et al., 2006; Gardner and West, 2010; Veelders et al., 2010) to cooperate conditionally with kin. Cooperation can also be bolstered by pleiotropic connections to personal benefits (Foster et al., 2004; Dandekar et al., 2012) or through association with alleles encoding self-benefitting traits (Asfahl et al., 2015). In these cases, the alleles may provide private benefits that are completely independent from the public benefits of cooperation. In asexual populations of cooperators

and defectors, this sets the stage for an "adaptive race" in which both types vie for the first highly beneficial adaptation (Waite and Shou, 2012; Morgan et al., 2012). The tragedy of the commons can be deferred if a cooperator, by chance, wins the adaptive race.

Hammarlund et al. (2015) recently showed that in spatially-structured populations, the "Hankshaw effect" can give cooperators a substantial leg up on defectors in an adaptive race. This advantage is reminiscent of Sissy Hankshaw, a fictional character in Tom Robbins' Even Cowgirls Get the Blues, whose oversized thumbs—which were otherwise an impairment—made her a prolific hitchhiker. Similarly, cooperation is costly, but it increases local population density. As a result, cooperators are more likely to acquire beneficial mutations. By hitchhiking along with these adaptations, cooperation can rise in abundance. Nevertheless, this advantage is fleeting. As soon as the oppor-72 tunities for adaptation are exhausted, cooperators are once again at a selective 73 disadvantage against equally-adapted defectors that arise via mutation. However, Hammarlund et al. (2015) also demonstrated that cooperation can be maintained when frequent environmental changes produce a steady stream of new adaptive opportunities. Although organisms typically find themselves in dynamic environments, the nature and frequency of these changes might not ensure long-term cooperator survival.

However, organisms do more than simply experience changing environments passively. Through their activities, their interactions with others, and even their deaths, organisms constantly modify their environment. This *niche construction* process can produce evolutionary feedback loops in which environ-

mental change alters selection, which, in turn, alters the distribution of types and their corresponding influence on the environment (Odling-Smee et al., 2003). The nature of this feedback can have dramatic evolutionary consequences. One critical distinction is whether the constructing type is favored in the resulting environment. Under positive niche construction, selection favors the constructor, and evolution stagnates as this type fixes. Under negative niche construction, selection favors a type other than the constructor, which creates an opportunity for novel adaptation. If the resulting adapted type also engages in negative niche construction, cycles of construction and adaptation can ensue, such that populations find themselves continually chasing beneficial mutations as their adaptive landscape perpetually shifts.

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Here, we show that the selective feedbacks that result from niche construction
can maintain cooperation indefinitely. We find that it is specifically negative
niche construction that is responsible for this result because of the endless
opportunities for adaptation that it produces. These results indicate that
cooperators can ensure their survival when they play an active role in their
own evolution.

₁ Methods

Building upon Hammarlund et al. (2015), we describe an individual-based model in which cooperators and defectors evolve and compete in a population of subpopulations (i.e., a metapopulation). Through mutations, individuals gain adaptations to their environment, which increase reproductive fitness, and allow those lineages to rise in abundance. More successful lineages spread to neighboring subpopulations by migration.

In this expanded model, subpopulations additionally modify their local environment. As this process occurs, environmental changes feed back to affect selection. We explore how niche construction affects the evolution of cooperation; specifically, how cooperative behavior can hitchhike along with adaptations to modified environments. Box 1 provides additional information about the model.

114 Model Description

Individual Genotypes and Adaptation

Each individual has a haploid genome with L+1 loci (see Table 1 for model 116 parameters and their values). Different alleles at each locus are represented 117 by different integers. A binary allele at the cooperation locus (locus zero) 118 determines whether that individual is a cooperator (1), which carries fitness 119 cost c, or a defector (0). The remaining L loci are adaptive loci, and are each 120 occupied by 0 or a value from the set $\{1, 2, \dots, A\}$. Allele 0 represents a lack of 121 adaptation, while a non-zero allele represents one of the A possible adaptations 122 at that locus. 123

These non-zero alleles signify two types of adaptations, both of which increase fitness. First, adaptations to the external environment confer a fitness benefit δ . This selective value is the same regardless of which non-zero allele is present and is not affected by other individuals. We assume $\delta > c$, which allows a

minimally adapted cooperator to recoup the cost of cooperation and gain a fitness advantage.

Niche Construction and Selective Feedbacks

Individual fitness is also affected by aspects of the local environment that are 131 affected by organisms. We implicitly represent this constructed "niche" based 132 on the specific allelic states present in the subpopulation. As allelic states 133 change, the subpopulation alters its environment, creating a unique niche. As 134 described below, the specific alleles that are present at each locus matter. 135 In our model, the feedback from niche construction takes the form of den-136 sity dependent selection, and individuals evolve to better match their niche. 137 Specifically, the selective value of non-zero allele a at adaptive locus l—and 138 consequently the fitness of an individual carrying that allele—increases with 139 the number of individuals in the subpopulation that have allele a-1 at locus 140 l-1. For example, if L=5 and A=6, and allele 4 has fixed at locus 2, then 141 a genotype with allele 5 at locus 3 is favored. And as allele 5 fixes at locus 142 3, the niche that this population constructs will favor allele 6 at locus 4 (see 143 Box 1). As a consequence, genotypes with sequentially increasing allelic states 144 will tend to evolve. We treat both adaptive loci and their non-zero allelic 145 states as "circular": the selective value of an allele at locus 1 is affected by the allelic composition of the subpopulation at locus L. Similarly, the selective 147 value of allele 1 at any locus increases with the number of individuals carrying 148 allele A at the previous locus. This circularity is represented by the function

 $\beta(x,X)$, which gives the integer that is below an arbitrary value x in the set $\{1,2,\ldots,X\}$:

$$\beta(x,X) = \operatorname{mod}_X(x-2+X) + 1 \tag{1}$$

Here, $\operatorname{mod}_X(x)$ is the integer remainder when dividing x by X. Using this function, the selective value of allele a at adaptive locus l is increased by ϵ for each individual in the subpopulation that has allele $\beta(a,A)$ at locus $\beta(l,L)$. Thus, ϵ specifies the intensity of selection due to niche construction.

Consider a genotype g with allelic state $a_{g,l}$ at locus l; the fitness of an indi-

vidual with this genotype is defined as:

$$W_{g} = z - \underbrace{ca_{g,0}}_{\text{cost of cooperation}} + \underbrace{\delta \sum_{l=1}^{L} I(a_{g,l})}_{\text{adaptation to}} + \underbrace{\epsilon \sum_{l=1}^{L} n(\beta(a_{g,l}, A), \beta(l, L))}_{\text{adaptation to external environment}}$$
(2)

where z is a baseline fitness, n(a,l) is the number of individuals in the subpopulation with allele a at locus l, and I(a) indicates whether a given allele is non-zero:

$$I(a) = \begin{cases} 1 & \text{if } a \in \{1, 2, \dots, A\} \\ 0 & \text{otherwise} \end{cases}$$
 (3)

Thus, an individual's fitness is determined both by adaptations to the external environment and by adaptations to its constructed environment. Box 1 illustrates the process of adaptation to the constructed environment. While cooperation is costly, we assume its effects are independent of the external and constructed components of the environment.

Subpopulation Growth and the Benefit of Cooperation

Cooperation enables a subpopulation to reach a greater density. This benefit affects all individuals equally and accumulates linearly with the proportion of cooperators in the subpopulation. If p is the proportion of cooperators present at the beginning of a growth cycle, then that subpopulation reaches the following size:

$$S(p) = S_{min} + p(S_{max} - S_{min}) \tag{4}$$

During subpopulation growth, individuals compete through differential reproduction. Each individual's probability of success is determined by its fitness. The composition of a subpopulation with size P and cooperator proportion pafter growth is multinomial with parameters S(p) and $\{\pi_1, \pi_2, \dots, \pi_P\}$, where π_i represents individual i's reproductive fitness relative to others in the subpopulation (using Equation 2).

178 Mutation

For simplicity, we apply mutations after subpopulation growth. Mutations occur independently at each locus and cause an allelic state change. At the

binary cooperation locus, mutations occur at rate μ_c . These mutations flip the allelic state, causing cooperators to become defectors and vice versa. Mutations occur at rate μ_a at each adaptive locus. These mutations replace the existing allele with a random selection from the set $\{0\} \cup \{1, 2, ..., A\}$. Because mutations are stochastic, the allelic sequences that evolve depend on which allele arises first and at which locus.

187 Migration

Populations are composed of N^2 patches arranged as an $N \times N$ lattice, where each patch can support a subpopulation. After mutation, individuals emigrate to an adjacent patch with probability m. During each migration event, a single destination patch is randomly chosen from each source patch's Moore neighborhood, which is composed of the nearest 8 patches on the lattice. Because the population lattice has boundaries, patches located on the periphery have smaller neighborhoods.

Population Initialization and Simulation

Following Hammarlund et al. (2015), we begin simulations with sparse populations. Subpopulations are first seeded at all patches with size $S(p_0)$ and cooperator proportion p_0 . The population is then thinned. Each individual survives this bottleneck with probability σ . Starting from this initial state, simulations then proceed for T cycles, where each discrete cycle consists of subpopulation growth, mutation, migration, and dilution. Dilution reduces each subpopulation to support growth in the next cycle. Each individual remains with probability d, regardless of its genotype.

Simulation Source Code and Software Dependencies

The simulation software and configurations for the experiments reported are available online. Simulations used Python 3.4, NumPy 1.9.1, Pandas 0.15.2 (McKinney, 2010), and NetworkX 1.9.1 (Hagberg *et al.*, 2008). Data analyses were performed with R 3.1.3 (R Core Team, 2015). Reported confidence intervals were estimated by bootstrapping with 1000 resamples.

Results

Using the model described in the previous section, we perform simulations that 211 follow the evolution of cooperation in a population of subpopulations that are 212 connected by spatially-limited migration. Individuals increase their compet-213 itiveness by gaining adaptations. While cooperation does not directly affect 214 the fitness benefits that these adaptations confer, cooperation has indirect 215 effects on the adaptive process. Specifically, cooperation increases subpopulation density. As a result, larger subpopulations of cooperators experience more 217 mutational opportunities. Cooperation can rise in abundance by hitchhiking along with beneficial mutations, which compensate for the cost of cooperation. 219 Importantly, subpopulations alter their local environments, which feeds back

 $^{^{1}}$ These materials will be made public prior to publication.

to influence selection. Here, we explore how such niche construction affects the evolution of cooperation.

223 Cooperation Persists with Niche Construction

Without any opportunity for adaptation (L=0), cooperators are swiftly eliminated from populations (Figure 1A). Despite an initial lift in cooperator abun-225 dance due to increased productivity, the cost of cooperation becomes disad-226 vantageous as migration mixes the initially isolated subpopulations. When 227 populations can adapt to the external environment $(L > 0 \text{ and } \delta > 0)$, but 228 niche construction is absent ($\epsilon = 0$), cooperators are maintained only transiently (Figure 1B). Here, larger cooperator subpopulations can more quickly 230 adapt to their external environment. However, as previously described by 231 Hammarlund et al. (2015), cooperation is subsequently lost once populations 232 become fully adapted. This occurs when isogenic defectors (i.e., defectors with 233 identical adaptive loci) arise via mutation and displace cooperators due to their 234 selective advantage. However, when niche construction feeds back to influence 235 selection ($\epsilon > 0$), cooperation persists in the majority of the replicate popula-236 tions (Figure 1C). We see in Figure 2A that despite oscillations, cooperation 237 is maintained at high levels in these populations. 238

Fitness Increases Alone do not Support Persisting Cooperation

In the model, adaptations to both the external environment and the constructed environment contribute to an individual's fitness. To determine 242 whether cooperation is maintained solely due to the larger selective values 243 that result from the contributions of niche construction, we performed simula-244 tions in which these contributions were removed ($\epsilon = 0$). We instead increased 245 the fitness benefits conferred by adaptation to the external, non-constructed 246 environment ($\delta = 0.6$). In doing so, we conservatively estimate the selective 247 effects of niche construction by supplementing the selective benefits of adapta-248 tions to the external environment by the maximum possible selective benefit 249 that results from niche construction. Nevertheless, we find that simply increas-250 ing selective values does not enable cooperators to persist (Figure 2B). Niche 251 construction, therefore, plays a decisive role here.

Negative Niche Construction is Critical to Cooperator Persistence

In our model, an adaptation to a constructed aspect of the environment initiates a new instance of niche construction, leading to sequentially increasing allelic states across the adaptive loci. Under certain conditions, this construction always makes the constructor sub-optimal for the niche it creates (see Box 1). This negative niche construction occurs when the number of adaptive alleles (A) does not divide evenly into the number of adaptive loci (L). In such a case, any sequence of integers on the circular genome will always contain a break in the sequence; that is, one locus will will have an allele that is not one less than the allele at the next locus. Given this unavoidable mismatch, any type that has fixed will always construct a niche that favors selection for a new type. When negative niche construction is removed (by setting L=5, A=5), cooperators are again driven extinct after an initial lift in abundance (Figure 2C). These results indicate that the type of niche construction matters. Specifically, negative niche construction is crucial for maintaining cooperation.

²⁶⁹ Selective Feedbacks Limit Defector Invasion

The adaptation resulting from selective feedbacks can limit invasion by defectors, which arise either through immigration from neighboring patches or 271 through mutation at the cooperation locus. The latter challenge is particularly 272 threatening, as these isogenic defectors are equally adapted, yet do not incur 273 the cost of cooperation. As demonstrated in Figure 3A, isogenic defectors 274 rapidly spread when introduced at a single patch in the center of a population 275 of cooperators if mutations do not occur. However, when cooperators can gain 276 adaptations via mutation, cooperators resist defector invasion in over half of 277 the replicate populations (Figure 3B). Figure 4 depicts one such instance. In 278 that population, isogenic defectors are seeded at a single patch in an otherwise 279 all-cooperator population. These defectors quickly begin to spread. However, 280 a neighboring cooperator population gains an adaptation, which increases its 281

fitness above that of the defector. This type spreads more quickly, stopping the spread of defectors and eventually driving them to extinction. Because this 283 adaption occurs in a cooperator population, cooperation is able to hitchhike 284 to safety. Importantly, this new cooperator is favored because of the niche 285 that its ancestral type created. Here, cooperators can find safety in numbers— 286 because their larger populations have more mutational opportunities, they are 287 more likely to gain adaptations that rescue them from invasion. Further, these 288 larger cooperators exert a greater influence on their niche, which increases selection for an adapted type. This allows that type to appear and to spread more quickly in the population. Figure 3C shows how quickly an adapted coop-291 erator type can invade a population of defectors. Importantly, this cooperator 292 type is adapted to the niche constructed by the defector.

Discussion

Despite their negative effects, deleterious traits can rise in abundance due to genetic linkage with other traits that are strongly favored by selection (Maynard Smith and Haigh, 1974). In a process termed the "Hankshaw effect", Hammarlund et al. (2015) recently demonstrated that cooperation can actively prolong its existence by increasing its likelihood of hitchhiking with a beneficial trait. In that work and in ours, subpopulations of cooperators grow to a higher density than those of defectors. Because of this, these larger cooperator subpopulations experience more mutations and are therefore more likely to gain adaptations. While this process does favor cooperation in the

short term, it eventually reaches a dead end; when the opportunities for adaptation are exhausted, and cooperators can no longer hitchhike, they face extinction. Here, we have considered whether niche construction might serve to
perpetually generate new adaptive opportunities and thus favor cooperation
indefinitely.

When niche construction occurs, cooperation can indeed persist (Figures 1C and 2A). In our model, niche construction introduces additional selective effects that influence the evolutionary process, leading to a more pronounced Hankshaw effect. However, these fitness benefits alone do not maintain cooperation (Figure 2B). Niche construction and the selective feedbacks that it produces play a crucial role.

We find that it is specifically negative niche construction that maintains coop-315 eration (Figure 2C). As cooperator and defector types gain adaptations, they 316 alter their environment in ways that favor other types. Thus, negative niche 317 construction serves as a perpetual source of adaptation. Here we observe an-318 other facet of the Hankshaw effect: because populations of cooperators are larger, they are better able to respond to the adaptive opportunities that follow from negative niche construction. By gaining adaptations more quickly, 321 cooperators resist invasion by defectors (Figure 3B). Although defectors ini-322 tially have an advantage by saving on the cost of cooperation, subpopulations 323 of cooperators can quickly gain an advantage because they are larger. Even in 324 the presence of an equally-adapted defector type, cooperator subpopulations 325 are more likely to produce the mutant most adapted to the current niche, 326 which can then displace the slower evolving defectors. These recurring cycles

of defector invasion and cooperator adaptation underlie the oscillations in cooperator proportion seen in Figure 2A. When cooperators do not gain these adaptations, they are driven to extinction by the defector. This is something that we see occur stochastically in Figures 2A and 3B.

332 Cooperation as Niche Construction

In our model, niche construction and adaptation are independent of cooperation, which allows us to focus on hitchhiking. However, by increasing the size of the subpopulation, this form of cooperation can itself be seen as a kind of 335 niche construction. Cooperative benefits often take similar forms in natural systems. For example, bacteria produce a host of extracellular products that 337 scavenge soluble iron (Griffin et al., 2004), digest large proteins (Diggle et al., 338 2007; Darch et al., 2012), and reduce the risk of predation (Cosson et al., 2002), 339 among many others (West et al., 2007a). As in our model, these forms of co-340 operation are likely to increase local population density. While many studies 341 have focused on how the environment affects the evolution of these cooperative 342 traits, relatively few have addressed how the environmental changes created 343 by these products feed back to influence evolution. Perhaps most similar to this study, Van Dyken and Wade (2012) demon-345 strated that when two negative niche constructing, cooperative behaviors coevolve, selection can increasingly favor these traits, which are disfavored when alone. In that model, "reciprocal niche construction" occurred when the negative feedback resulting from one strategy positively influenced selection for

the other, creating a perpetual cycle that maintained both forms of cooperation. Arguably, this can be seen as an instance of hitchhiking: the currently-351 maladaptive form of cooperation is maintained by association with the adap-352 tive form. 353 When dispersal is limited, competition among kin can undermine cooperation. 354 To separate kin competition from kin selection, Lehmann (2007) developed 355 a model in which a cooperative, niche-constructing behavior only benefitted 356 future generations. Kin competition was thereby reduced, and cooperation 357 instead benefitted descendants. This work highlights an important aspect of 358 niche construction—often, the rate of selective feedback from niche construc-359

Evolution at Multiple Timescales

tion is different from the rate at which populations grow.

In our work, the niche is modeled implicitly by the composition of the population. Any changes in the population, therefore, produce immediate effects on 363 the constructed environment and the resulting selective feedbacks. However, 364 timescales in our model could be de-coupled in two ways. First, cooperators 365 modify their niche by enabling their subpopulation to reach larger density 366 (Equation 4). These increased subpopulation sizes play a critical role by ef-367 fectively increasing the rate of evolution in these subpopulations. Because of 368 the importance of this process, it would be very informative to explore how sensitive our results are to changes in how quickly population sizes increase and for how long they are upheld. Similarly, changes in the rate at which

a niche changes in response to population changes could substantially alter our results. Not only would such changes in timescale affect the selective values of alleles as the population changed, they could also influence whether or not populations were able to evolve adapted types and if so, how well those adapted types can propagate through the population to address the threat of a defector.

Other studies, while not focused on cooperation, have similarly shown that the 378 timescales at which niche construction feedbacks occur can strongly influence 379 evolutionary outcomes (Laland et al., 1996, 1999). This perspective is likely to 380 be crucial for understanding the evolution of cooperative behaviors like the pro-381 duction of public goods. In these instances, environmental changes are likely 382 to occur on different timescales than growth, which can have profound effects. 383 For example, a multitude of factors including protein durability (Brown and 384 Taddei, 2007; Kümmerli and Brown, 2010), diffusion (Allison, 2005; Driscoll 385 and Pepper, 2010), and resource availability (Zhang and Rainey, 2013; Ghoul 386 et al., 2014) influence both the rate and the degree to which public goods alter 387 the environment. While Lehmann (2007) showed that cooperation was favored 388 when selective feedbacks act over longer timescales, niche construction may in 389 fact hinder cooperation when selection is more quickly altered. For example, 390 when public goods accumulate in the environment, cooperators must decrease production to remain competitive (Kümmerli and Brown, 2010; Dumas and 392 Kümmerli, 2012). This favors cooperation that occurs facultatively, perhaps 393 by sensing the abiotic (Bernier et al., 2011; Koestler and Waters, 2014) or biotic environment (Brown and Johnstone, 2001; Darch et al., 2012). To allow our model to more fully address how traits such as these evolve, we could de-couple the niche from the composition of the population and represent the niche explicitly.

Cooperation and Niche Construction in Host-Symbiont Co-Evolution

As the niche becomes more independent from the population, it develops its 401 own state and dynamics. A logical next step, then, could be to treat the 402 environment as a biological entity itself, which could introduce additional evo-403 lutionary feedbacks. Such a model could be used to explore the evolution of 404 cooperation in host-symbiont systems, where cooperation among symbionts 405 affects host fitness. As the host population changes, either in response to sym-406 biont cooperation or other factors, so too does selection on their symbiont pop-407 ulations. Here, evolutionary outcomes depend greatly on the degree of shared 408 interest between the host and symbiont. Future models could explicitly cap-409 ture the environment as a biological entity to explore the rich co-evolutionary 410 dynamics that these systems might offer. 411 For example, the cooperative production of virulence factors by the human 412 pathogen P. aeruginosa is harmful to hosts with cystic fibrosis (Harrison, 2007). 413 Following what we have shown in this work, these antagonistic, negative niche 414 constructing behaviors might actually work to maintain these infections. If 415 these populations do indeed perpetually benefit from adaptations that are 416 created by niche construction as we have shown, a case could perhaps be

made for developing treatments that target the selective feedback loop that provides adaptive opportunities in these spatial environments. While the idea 419 of removing negative selective feedbacks and supporting stability may seem 420 counterintuitive, if it leaves the infecting population more susceptible, then 421 perhaps pairing such a treatment with ones in which mutants are introduced 422 (see e.g., Rumbaugh et al. (2009)), could significantly improve host fitness. Expanding models such as ours to address the additional dynamics present in host-symbiont systems such as these could be quite productive. It was recently argued that incorporating the effects of niche construction is 426 critical for improving our understanding of viral evolution (Hamblin et al., 427 2014) and evolution in co-infecting parasites (Hafer and Milinski, 2015). In-428 corporating host dynamics, co-evolution, and the feedbacks that they produce 429 is likely to be equally important for gaining a greater understanding of how 430 cooperative behaviors evolve in these host-symbiont settings.

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Box 1: Description of niche construction in our

$_{\scriptscriptstyle{ ext{ t 441}}}$ $oxdot{oxmodel}$

442 See Figure B1

- (A) Individuals. The genome of each individual consists of a single coop-443 eration locus and L adaptive loci (here, L=5). At the cooperation locus (labeled 0), this individual has allele 1, making it a cooperator. The adaptive 445 loci (labeled 1-5) are arranged as a circular chromosome, where each locus has 446 an integer allele between 0 and A, inclusive. In the description that follows, 447 we focus exclusively on these adaptive loci. Genotypes are given by their al-448 lelic states starting with locus 1. For instance, the genotype shown here is 449 [2,0,5,2,1]. Because of their circular structure, allele 2 at the first locus follows 450 allele 1 at the fifth locus.
- (B) Niche Construction. Consider a subpopulation fixed for genotype 452 [1,2,0,0,0]. This subpopulation constructs environment $E_{[1,2,0,0,0]}$. Every non-453 zero allele influences selection at the next locus, favoring sequential allelic 454 states. In this constructed environment, allele 3 at locus 3 would be favored. 455 If genotype [1,2,3,0,0] arises via mutation, it is expected to fix. However, 456 genotype [1,2,3,0,0] affects the environment differently. As [1,2,3,0,0] rises in 457 abundance, the constructed environment changes to $E_{[1,2,3,0,0]}$, which favors 458 [1,2,3,4,0]. 459
- 460 (C) Niche Construction and Adaptation. The evolutionary transition 461 shown in Part B is indicated in the dashed box. Here, we depict entire sub-

populations fixed for a genotype using a single instance of that genotype. Simi-462 larly, an arrow represents niche construction and adaptation to the constructed 463 environment. We start with a case in which there are five alleles (A = 5). Sub-464 populations begin with the non-adapted genotype [0,0,0,0,0], shown on the far 465 left. A non-zero allele is introduced via mutation, which represents an adapta-466 tion to external aspects of the environment. Here, allele 1 arises and fixes at 467 locus 1. The remainder of this figure focuses on adaptation to the constructed 468 aspects of the environment. This genotype has a "mismatch" (shown by the 469 red sector), because $E_{[1,0,0,0,0]}$ favors [1,2,0,0,0]. Assuming allele 2 arises at the 470 second locus, it will be selected, creating a "match" at the first and second 471 loci (green sector). Now there is a mismatch between the second and third 472 loci in the resulting environment, which a new round of mutation and selection 473 corrects, and so on. The green sector grows as the red sector shifts clockwise. 474 When the population reaches [1,2,3,4,5], it constructs $E_{[1,2,3,4,5]}$. Here, since 475 allele 1 follows allele 5, there is no longer a mismatch, so no further adaptation 476 occurs. 477

478 (**D**) Negative Niche Construction. A different case emerges when the 479 number of alleles does not evenly divide into the number of loci. Here, we 480 change the number of alleles to six (A = 6). As shown on the far left, we 481 begin with a subpopulation fixed for genotype [1,2,3,4,5]. This genotype has a 482 mismatch, because the niche constructed by allele 5 favors allele 6 (not 1) at the 483 next locus (locus 1). A mutant with genotype [6,2,3,4,5] has a fitness advantage 484 and can fix in $E_{[1,2,3,4,5]}$. However, as this type constructs $E_{[6,2,3,4,5]}$, a new 485 mismatch appears. In this instance of negative niche construction, adapting

- $_{486}$ to correct one mismatch generates a new mismatch. This system can never
- $_{\tt 487}$ $\,$ escape its mismatches—the red sector just shifts clockwise around the genome
- 488 perpetually.

Figures Figures

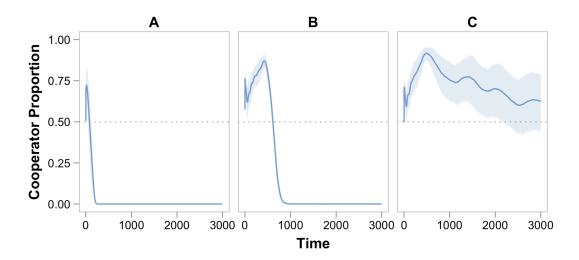


Figure 1: Adaptation, Hitchhiking, and the Evolution of Cooperation. Curves show the average cooperator proportion among replicate populations for the duration of simulations, and shaded areas indicate 95% confidence intervals. Unless otherwise noted, parameter values are listed in Table 1. (A) Without any opportunity to adapt (L=0), cooperation is quickly lost. (B) When adaptation can occur (L=5), but niche construction does not affect selection $(\epsilon=0)$, cooperators rise in abundance by hitchhiking along with adaptions to the external environment. Nevertheless, this effect is transient, and cooperators eventually become extinct. (C) Niche construction enables cooperation to be maintained indefinitely. In the majority of populations, cooperation remained the dominant strategy. The trajectories of individual populations are shown in Figure 2A.

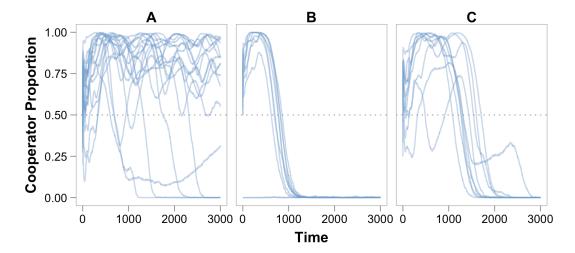


Figure 2: Niche Construction and the Evolution of Cooperation. The proportion of cooperators present in each replicate population is shown for the duration of simulations. (A) Despite some oscillations, cooperation dominates in 13 of 18 populations when niche construction affects selection. (B) When the selective effects of niche construction (ϵ) are removed, and the selective benefit of adaptation to the external environment (δ) is increased to compensate, cooperators are driven to extinction by isogenic defectors that arise by mutation ($\epsilon = 0$, $\delta = 0.6$). Note that cooperation was not present after initialization in one replicate population. (C) Cooperators are also driven to extinction without negative niche construction (A = 0).

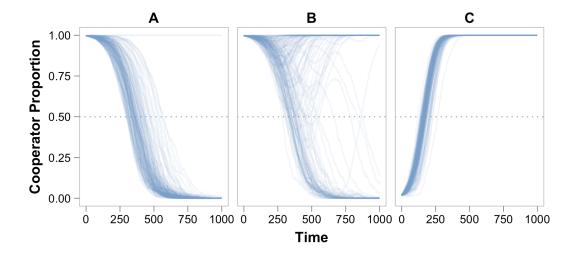


Figure 3: Niche Construction and Invasion. Curves trace the proportion of cooperators present in each replicate population for the duration of simulations (T = 1000). In each simulation, a rare type was initiated at a single patch in the center of the population lattice ($N^2 = 121$). Unless otherwise noted, mutations are disabled in these ecological simulations to highlight the dynamics of invasion ($\mu_a = 0, \mu_c = 0$). (A) When cooperators and defectors are isogenic (i.e., both types have stress alleles [1,2,3,4,5]), rare defectors quickly invade and drive cooperators to extinction due to the cost of cooperation. Defectors were stochastically eliminated in 2 replicate populations. (B) However, negative niche construction creates adaptive opportunities that enable cooperators to resist invasion by isogenic defectors. Here, cooperation remained the dominant in 91 of 160 populations ($\mu_a = 0.00005$). Results from simulations where mutations also occurred at the cooperation locus are shown in Figure S1. (C) In fact, an adapted cooperator type (stress alleles [6,2,3,4,5], see Box 1) can swiftly displace defectors when isogenic defectors cannot arise or adapt via mutation.

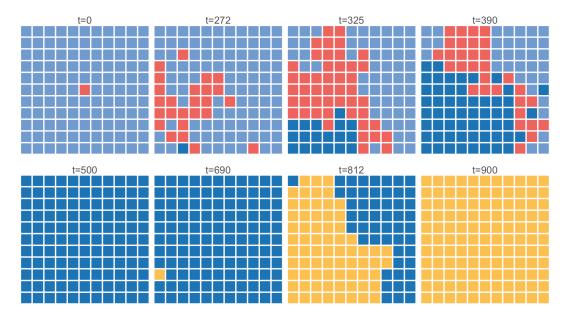


Figure 4: Cooperator Adaptation Prevents Defector Invasion. Here we depict the distribution of dominant types among subpopulations over time for one representative simulation in which isogenic defectors arise. To highlight the effects of adaptation, mutations did not occur at the cooperation locus ($\mu_c = 0$). At time t = 0 (upper left panel), a single isogenic defector population (red) is placed among cooperator populations (light blue). Because these defectors do not bear the costs of cooperation, they spread (t = 272). However, cooperators in one population gain an adaptation that gives them a fitness advantage over defectors (second panel, dark blue, lower left). At t = 325, defectors continue to invade cooperator populations. However, the adapted cooperator type spreads more quickly due to its fitness advantage, invading both defector populations and ancestral cooperator populations (t = 390), until it eventually fixes in the population (t = 500). At t = 690, a new cooperator type emerges that is favored due to negative niche construction (orange). This new type spreads rapidly (t = 812) until reaching fixation (t = 900). At this point, it becomes susceptible to invasion by the next "adapted" cooperator type, and the cycle continues.

Box 1 Figures

Figure B1

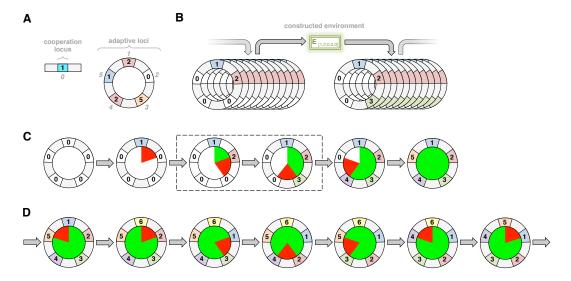


Figure B1: Figure for Box 1

Supplemental Figures

Figure S1

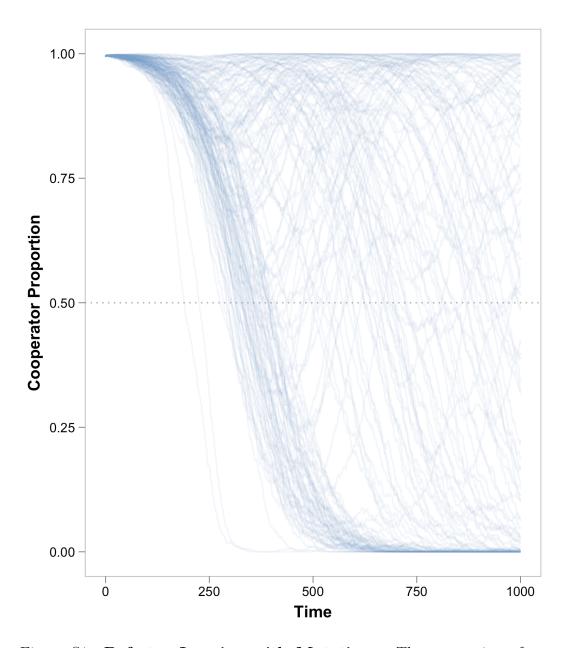


Figure S1: **Defector Invasion with Mutations.** The proportion of cooperators present in each replicate population is shown for the duration of simulations (T = 1000). When mutations occur both at the adaptive loci and the cooperation locus ($\mu_a = \mu c = 0.00005$), cooperation remains dominant in 58 of 160 replicate populations.

Tables

Table 1: Model parameters and their value

Parameter	Description	Base Value
\overline{L}	Number of adaptive loci	5
c	Fitness cost of cooperation	0.1
A	Number of alleles	6
δ	Fitness benefit, adaptation to external environment	0.3
ϵ	Fitness benefit, adaptation to constructed environment	0.00015
z	Baseline fitness	1
S_{min}	Minimum subpopulation size	800
S_{max}	Maximum subpopulation size	2000
μ_a	Mutation rate (adaptation)	10^{-5}
μ_c	Mutation rate (cooperation)	10^{-5}
N^2	Number of patches	625
m	Migration rate	0.05
p_0	Initial cooperator proportion	0.5
σ	Survival rate at population initialization	10^{-5}
T	Number of simulation cycles	3000
d	Subpopulation dilution factor	0.1

References

- Allison, S.D. 2005. Cheaters, diffusion and nutrients constrain decomposition
- by microbial enzymes in spatially structured environments. Ecology Letters,
- 502 **8**: 626–635.
- Asfahl, K.L., Walsh, J., Gilbert, K. and Schuster, M. 2015. Non-social adap-
- tation defers a tragedy of the commons in Pseudomonas aeruginosa quorum
- sensing. The ISME Journal, doi: 10.1038/ismej.2014.259.
- Bernier, S.P., Ha, D.-G., Khan, W., Merritt, J.H.M. and O'Toole, G.A. 2011.
- Modulation of Pseudomonas aeruginosa surface-associated group behaviors by
- individual amino acids through c-di-GMP signaling. Research in Microbiology,
- 509 **162**: 680–688.
- 510 Brown, S.P. and Johnstone, R.A. 2001. Cooperation in the dark: Signalling
- and collective action in quorum-sensing bacteria. Proceedings of the Royal
- 512 Society of London B: Biological Sciences, 268: 961–965.
- Brown, S.P. and Taddei, F. 2007. The durability of public goods changes the
- dynamics and nature of social dilemmas. PLoS ONE, 2: e593.
- ⁵¹⁵ Cosson, P., Zulianello, L., Join-Lambert, O., Faurisson, F., Gebbie, L. and
- Benghezal, M. et al. 2002. Pseudomonas aeruginosa virulence analyzed in a
- Dictyostelium discoideum host system. Journal of Bacteriology, 184: 3027-
- 518 3033.
- Dandekar, A.A., Chugani, S. and Greenberg, E.P. 2012. Bacterial quorum
- sensing and metabolic incentives to cooperate. Science, 338: 264–266.

- Darch, S.E., West, S.A., Winzer, K. and Diggle, S.P. 2012. Density-dependent
- 522 fitness benefits in quorum-sensing bacterial populations. Proceedings of the
- National Academy of Sciences, 109: 8259–8263.
- Diggle, S.P., Griffin, A.S., Campbell, G.S. and West, S.A. 2007. Cooperation
- and conflict in quorum-sensing bacterial populations. *Nature*, **450**: 411–414.
- Driscoll, W.W. and Pepper, J.W. 2010. Theory for the evolution of diffusible
- external goods. Evolution, 64: 2682–2687.
- Dumas, Z. and Kümmerli, R. 2012. Cost of cooperation rules selection for
- cheats in bacterial metapopulations. Journal of Evolutionary Biology, 25:
- ₅₃₀ 473–484.
- Fletcher, J.A. and Doebeli, M. 2009. A simple and general explanation for the
- evolution of altruism. Proceedings of the Royal Society B: Biological Sciences,
- 533 **276**: 13–19.
- Foster, K., Shaulsky, G., Strassmann, J., Queller, D. and Thompson, C. 2004.
- Pleiotropy as a mechanism to stabilize cooperation. *Nature*, **431**: 693–696.
- Gardner, A. and West, S.A. 2010. Greenbeards. Evolution, 64: 25–38.
- Ghoul, M., West, S.A., Diggle, S.P. and Griffin, A.S. 2014. An experimental
- test of whether cheating is context dependent. Journal of Evolutionary Biology,
- **27**: 551–556.
- Griffin, A.S., West, S.A. and Buckling, A. 2004. Cooperation and competition
- in pathogenic bacteria. Nature, 430: 1024–1027.
- Hafer, N. and Milinski, M. 2015. When parasites disagree: Evidence for

- parasite-induced sabotage of host manipulation. Evolution, 69: 611–620.
- Hagberg, A.A., Schult, D.A. and Swart, P.J. 2008. Exploring network struc-
- 545 ture, dynamics, and function using NetworkX. In: Proceedings of the 7th
- Python in Science Conference (SciPy2008), pp. 11–15.
- Hamblin, S.R., White, P.A. and Tanaka, M.M. 2014. Viral niche construction
- alters hosts and ecosystems at multiple scales. Trends in Ecology & Evolution,
- **29**: 594–599.
- 550 Hamilton, W.D. 1964. The genetical evolution of social behaviour I & II.
- Journal of Theoretical Biology, 7: 1–52.
- Hammarlund, S.P., Connelly, B.D., Dickinson, K.J. and Kerr, B. 2015. The
- evolution of cooperation by the Hankshaw effect. bioRxiv, doi: 10.1101/016667.
- 554 Cold Spring Harbor Labs Journals.
- Harrison, F. 2007. Microbial ecology of the cystic fibrosis lung. *Microbiology*,
- 556 **153**: 917–923.
- 557 Koestler, B.J. and Waters, C.M. 2014. Bile acids and bicarbonate inversely
- regulate intracellular cyclic di-GMP in Vibrio cholerae. Infection and Immu-
- nity, **82**: 3002–3014.
- 560 Kuzdzal-Fick, J.J., Fox, S.A., Strassmann, J.E. and Queller, D.C. 2011. High
- relatedness is necessary and sufficient to maintain multicellularity in Dic-
- tyostelium. *Science*, **334**: 1548–1551.
- Kümmerli, R. and Brown, S.P. 2010. Molecular and regulatory properties of
- a public good shape the evolution of cooperation. Proceedings of the National

- 565 Academy of Sciences, **107**: 18921–18926.
- Laland, K.N., Odling-Smee, F.J. and Feldman, M.W. 1999. Evolutionary con-
- sequences of niche construction and their implications for ecology. *Proceedings*
- of the National Academy of Sciences, **96**: 10242–10247.
- Laland, K.N., Odling-Smee, F.J. and Feldman, M.W. 1996. The evolutionary
- consequences of niche construction: A theoretical investigation using two-locus
- theory. Journal of Evolutionary Biology, 9: 293–316.
- Lehmann, L. 2007. The evolution of trans-generational altruism: Kin selection
- meets niche construction. Journal of Evolutionary Biology, 20: 181–189.
- Maynard Smith, J. and Haigh, J. 1974. The hitch-hiking effect of a favourable
- gene. Genetics Research, 23: 23–35.
- McKinney, W. 2010. Data structures for statistical computing in Python. In:
- 577 Proceedings of the 9th Python in Science Conference (S. van der Walt and J.
- 578 Millman, eds), pp. 51–56.
- Morgan, A.D., Quigley, B.J.Z., Brown, S.P. and Buckling, A. 2012. Selection
- on non-social traits limits the invasion of social cheats. Ecology Letters, 15:
- 581 841-846.
- Nadell, C.D., Foster, K.R. and Xavier, J.B. 2010. Emergence of spatial struc-
- ture in cell groups and the evolution of cooperation. PLoS Computational
- Biology, 6: e1000716.
- Nowak, M.A. 2006. Five rules for the evolution of cooperation. Science, 314:
- ₅₈₆ 1560–1563.

- Odling-Smee, F.J., Laland, K.N. and Feldman, M.W. 2003. Niche construc-
- tion: The neglected process in evolution. Princeton University Press.
- R Core Team. 2015. R: A language and environment for statistical computing.
- Vienna, Austria: R Foundation for Statistical Computing.
- Rumbaugh, K.P., Diggle, S.P., Watters, C.M., Ross-Gillespie, A., Griffin, A.S.
- and West, S.A. 2009. Quorum sensing and the social evolution of bacterial
- virulence. Current Biology, 19: 341–345.
- Sinervo, B., Chaine, A., Clobert, J., Calsbeek, R., Hazard, L. and Lancaster,
- Let al. 2006. Self-recognition, color signals, and cycles of greenbeard mutu-
- ⁵⁹⁶ alism and altruism. Proceedings of the National Academy of Sciences, 103:
- ₅₉₇ 7372–7377.
- Van Dyken, J.D. and Wade, M.J. 2012. Origins of altruism diversity II: Run-
- away coevolution of altruistic strategies via "reciprocal niche construction".
- 600 Evolution, **66**: 2498–2513.
- Veelders, M., Brückner, S., Ott, D., Unverzagt, C., Mösch, H.-U. and Essen,
- 602 L.-O. 2010. Structural basis of flocculin-mediated social behavior in yeast.
- Proceedings of the National Academy of Sciences, 107: 22511–22516.
- Waite, A.J. and Shou, W. 2012. Adaptation to a new environment allows coop-
- erators to purge cheaters stochastically. Proceedings of the National Academy
- of Sciences, **109**: 19079–19086.
- West, S.A., Diggle, S.P., Buckling, A., Gardner, A. and Griffin, A.S. 2007a.
- The social lives of microbes. Annual Review of Ecology, Evolution, and Sys-
- 609 tematics, **38**: 53–77.

- 610 West, S.A., Griffin, A.S. and Gardner, A. 2007b. Evolutionary explanations
- $_{611}$ for cooperation. Current Biology, 17: R661–R672.
- ⁶¹² Zhang, X.-X. and Rainey, P.B. 2013. Exploring the sociobiology of pyoverdin-
- producing Pseudomonas. Evolution, 67: 3161–3174.