# Negative Niche Construction Favors the

# Evolution of Cooperation

3

4 Abstract

5 TODO

#### 6 Introduction

- <sup>7</sup> Cooperative behaviors are common across all branches of the tree of life. In-
- 8 sects divide labor within their colonies, plants and soil bacteria exchange es-
- 9 sential nutrients, birds care for others' young, and the trillions of cells in the
- human body coordinate to provide vital functions. Each instance of cooper-
- 11 ation 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
- 15 extinction.

Several factors can prevent this tragedy of the commons (Hamilton, 1964; Nowak, 2006; West et al., 2007). 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 27 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 opportunities for adaptation are exhausted, cooperators are once again at a selective 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.

Importantly, 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 environmental 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 or some other type is most adapted 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. In this latter case, populations find themselves continually chasing beneficial mutations as their adaptive landscape perpetually shifts.

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.

#### 69 Methods

Building upon Hammarlund et al. (2015), we develop 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 this process hitchhiking

along with adaptations, and whether cooperation can be maintained by selec-

80 tive feedbacks.

#### 81 Model Description

#### 82 Individual Genotypes and Adaptation

Each individual has a haploid genome with L+1 loci (see Table 1 for model parameters and their values). Different alleles at each locus are represented by 84 different integers. A binary allele at the first locus (here, locus zero) determines 85 whether that individual is a cooperator (1), which carries fitness cost c, or a 86 defector (0). Cooperation is independent from adaptation to the environment. 87 The remaining L loci are adaptive loci, and are each occupied by 0 or a value 88 from the set  $\{1, 2, \ldots, A\}$ . Allele 0 represents a lack of adaptation, while a 89 non-zero allele represents one of the A possible adaptations at that locus. 90 These non-zero alleles signify two types of adaptations, both of which increase 91 fitness. First, adaptations to the external environment confer a fitness benefit 92  $\delta$ . This selective value is the same regardless of which non-zero allele is present and is not affected by other individuals or the local niche. We assume  $\delta > c$ , which allows a minimally adapted cooperator to recoup the cost of cooperation 95 and gain a fitness advantage.

#### 97 Niche Construction and Selective Feedbacks

- Individual fitness is also affected by the current state of the local environment.
- We represent the "niche" implicitly based on the specific allelic states present
- in the subpopulation. As allelic states change, subpopulations alter aspects of
- their environment, creating a unique niche. As described below, the specific

alleles that are present at each locus matter.

Niche construction takes the form of density dependent selection, and individ-103 uals evolve to better match their niche by an additional form of adaptation. 104 The niche is defined by the distribution of alleles at each locus. Non-zero 105 alleles that are more common will improve fitness by a larger selective value 106 (beyond  $\delta$ ). Specifically, the selective value of adaptive allele a at locus l, and 107 consequently the fitness of an individual carrying that allele, increases with 108 the number of individuals in the subpopulation that have allele a-1 at locus 109 l-1. As a consequence, genotypes with sequentially increasing allelic states 110 will tend to evolve. We treat both adaptive loci and allelic states as "circular": 111 the selective value of an allele at locus 1 is affected by the allelic composition 112 of the subpopulation at locus L. Similarly, the selective value of allele 1 at 113 any locus increases with the number of individuals carrying allele A at the 114 previous locus. This circularity is represented by the function  $\beta(x, X)$ , which 115 gives the integer that is below an arbitrary value x in the set  $\{1, 2, \dots, X\}$ : 116

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

Here,  $\operatorname{mod}_X(x)$  is the integer remainder when dividing x by X. Using this function, the selective value of adaptive allele a at locus l is increased by  $\epsilon$  for each individual in the subpopulation that has allele  $\beta(a,A)$  at locus  $\beta(l,L)$ . Thus,  $\epsilon$  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 individual with this genotype is defined as:

$$W_{g} = z - \underbrace{ca_{g,0}}_{\text{cost of cooperation}} + \underbrace{\delta \sum_{l=1}^{L} I(a_{g,l})}_{\text{adaptation to external environment}} + \underbrace{\epsilon \sum_{l=1}^{L} n(\beta(a_{g,l}, A), \beta(l, L))}_{\text{adaptation to constructed environment}}$$
(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 adaptive 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 ( $\delta$ ) and by adaptations to its constructed environment ( $\epsilon$ ). Figure 128 1 illustrates the effects of these two components.

#### Population 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:

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

35 During 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 p after growth is multinomial with parameters S(p) and  $\{\pi_1, \pi_2, \dots, \pi_P\}$ , where  $\pi_i$  represents individual i's reproductive fitness relative to others in the subpopulation.

#### 140 Mutation

For simplicity, we apply mutations after growth. Mutations occur independently at each locus and cause an allelic state change. At the binary cooperation locus, mutations occur at rate  $\mu_c$ . These mutations flip the allelic state, causing cooperators to become defectors and vice versa. Mutations occur at rate  $\mu_a$  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.

#### 149 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 to create empty patches. Each individual survives this bottleneck with probability  $\sigma$ . Starting from this initial state, simulations then proceed for T cycles, where each discrete cycle consists of subpopulation growth, mutation, migration, and dilution. Dilution thins the population to support growth in the next cycle. Each individual remains with probability d, regardless of allelic state.

#### 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.

#### 172 Results

Using the model described in the previous section, we perform simulations that follow the evolution of cooperation in a population consisting of subpopulations that are connected by spatially-limited migration. Individuals increase their

<sup>&</sup>lt;sup>1</sup>These materials will be made public prior to publication.

competitiveness by gaining a limited number of adaptations. While coopera-176 tion does not directly affect the fitness benefits that these adaptations confer, 177 cooperation has indirect effects on the adaptive process. Specifically, cooper-178 ation increases subpopulation density. As a result, larger subpopulations of 179 cooperators experience more mutational opportunities to gain adaptations. Co-180 operation can rise in abundance by hitchhiking along with these adaptations, 181 which compensate for the cost of cooperation. During this process, subpopu-182 lations alter their local environments, which feeds back to influence selection. 183 Here, we explore how niche construction affects the evolution of cooperation. 184

#### Cooperation Persists with Niche Construction

Without any opportunity for adaptation (L=0), cooperators are swiftly elim-186 inated in competition with defectors (Figure 2A). Despite an initial lift in 187 cooperator abundance due to increased productivity, the cost of cooperation 188 becomes disadvantageous as migration mixes the initially isolated subpopula-189 tions. When populations can adapt to the external environment (L=5), but 190 niche construction is absent ( $\epsilon = 0$ ), cooperators are maintained only tran-191 siently (Figure 2B). Here, larger cooperator subpopulations can more quickly 192 adapt to their external environment as before. However, as previously de-193 scribed by Hammarlund et al. (2015), cooperation is subsequently lost once 194 populations become fully adapted to their environment. This occurs when iso-195 genic defectors (i.e., defectors with identical adaptive loci) arise via mutation 196 and displace cooperation due to their selective advantage. However, when 197

niche construction feeds back to influence selection, cooperation persists in over 2/3 of the replicate populations (Figure 2C). We see in Figure 3A that despite oscillations in the proportion of cooperators, cooperation is maintained at high levels in these populations.

# 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 205 whether cooperation is maintained solely due to the larger selective values 206 that result from the contributions of niche construction  $(\epsilon)$ , we performed 207 simulations in which these contributions were removed ( $\epsilon = 0$ ), and we in-208 stead increased the fitness benefits conferred by adaptation to the external, 209 non-constructed environment ( $\delta = 0.6$ ). In doing so, we conservatively esti-210 mate the selective effects of niche construction by supplementing the selective 211 benefits of adaptations to the external environment by the maximum possible 212 selective benefit that results from niche construction. Nevertheless, we find 213 that simply increasing selective values does not enable cooperators to persist 214 (Figure 3B). Niche construction, therefore, plays a decisive role here.

# Negative Niche Construction is Critical to Cooperator

Persistence

Adaptations can negatively effect fitness in our model due to selection for sequentially-increasing allelic states and the circular arrangement of these al-219 leles (see Figure 1). This negative niche construction occurs when the number 220 of adaptive alleles (A) does not divide evenly into the number of adaptive loci 221 (L). In such a case, any sequence of integers on the circular genome will always 222 contain a break in the sequence; that is, one locus with an allele that is not 223 one less than the allele at the next locus. Given this unavoidable mismatch, 224 any type that has fixed will always construct a niche that favors selection for 225 a new type. When negative niche construction is removed (by setting L=5, 226 A=5), cooperators are again driven extinct after an initial lift in abundance 227 (Figure 3C). These results indicate that the type of niche construction matters. 228 Specifically, negative niche construction is crucial for maintaining cooperation.

#### 230 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 through mutation from a cooperator ancestor. The latter challenge is particularly threatening, as these isogenic defectors are equally adapted, yet do not incur the cost of cooperation. As demonstrated in Figure 4A, isogenic defectors rapidly spread when introduced at a single patch in the center of a population of cooperators if mutations do not occur. However, when resident cooperators can gain adaptations via mutation, cooperators resist defector invasion in
over half of the replicate populations (Figure 4B). Figure 5 depicts one such
instance where cooperation survived. In that population, defectors quickly
began to spread. However, an adaptation arose in a neighboring cooperator
population. This type spread more quickly, halting defectors and eventually
driving them to extinction. Because this adaption occurred in a cooperator
population, cooperation was able to hitchhike to safety. Figure 4C shows how
quickly an adapted cooperator type can invade a population of defectors.

#### Negative Niche Construction Must Follow a Path

We have seen that negative niche construction plays a critical role in maintaining cooperation by creating adaptive "escape routes" for cooperators. But 248 in some cases, cooperator populations were not able to gain these adaptations 249 quickly enough, which led to extinction (Figure 3A). To determine whether 250 stronger negative feedbacks from niche construction would increase the rate at 251 which cooperator populations gained the adaptations needed to escape defec-252 tor invasion, we performed simulations in which the allelic state at each locus 253 was always mismatched. This was accomplished in the model by removing 254 selection for sequential allelic states. Instead, the selective value of an allele 255 at each locus increased with the number of individuals in the population that 256 had the next allelic state at that same locus. For example, selection in this 257 modified model would favor a type with [2, 5, 1, 4, 4] in a niche constructed 258 by [1,4,6,3,3] (L=5, A=6). However, this strongly negative niche construction does not better enable cooperators to stave off defection. In fact, cooperation is quickly lost under these conditions (Figure 6A). We then performed simulations to determine whether it is the rate of adaptation in response to negative niche construction that is important, not the strength of its feedback. When the mutation rate at adaptive loci is raised 100-fold ( $\mu_a = 0.001$ ), cooperation is maintained at higher levels and in more replicate populations (Figure 6B).

#### Discussion

Despite their negative effects, deleterious traits can rise in abundance due to genetic linkage with other traits that are strongly favored by selection (May-269 nard Smith and Haigh, 1974). In a process termed the "Hankshaw effect", 270 Hammarlund et al. (2015) recently demonstrated that cooperation can ac-271 tively prolong its existence by increasing its likelihood of hitchhiking with a 272 beneficial trait. While this process does favor cooperation in the short term, 273 it eventually reaches a dead end; when the opportunities for adaptation are 274 exhausted and cooperators can no longer hitchhike, they face extinction. In this work, we have considered whether niche construction might serve to perpetually generate new adaptive opportunities and thus favor cooperation in-277 definitely. When niche construction occurs, cooperation can indeed persist (Figures 2C) and 3A). In our model, niche construction introduces additional selective ef-280 fects that could influence the evolutionary process, leading to a more pronounced Hankshaw effect. However, simply raising the fitness benefits conferred by adaptations does not prolong cooperation (Figure 3B), which indicates that niche construction and the selective feedbacks that it produces play a crucial role.

Further, we find that it is specifically negative niche construction that main-286 tains cooperation (Figure 3C). Here we observe another facet of the Hankshaw 287 effect: because populations of cooperators are larger, they are better able to 288 respond to the adaptive opportunities that result from negative niche construc-289 tion. Without these adaptive opportunities, adaptation eventually grinds to 290 a halt. Once this occurs, cooperators face the threat of invasion by isogenic 291 defectors. Since these defectors are equally adapted but do not bear the cost 292 of cooperation, they quickly drive cooperators to extinction. Importantly, be-293 cause every type constructs an environment in which a different type is more fit. 294 negative niche construction creates continual adaptive opportunities. These 295 opportunities can allow cooperators to resist invasion by defectors, even when 296 defectors are equally adapted (Figure 4B). It is these recurring cycles of inva-297 sion and adaptation that underlie the oscillations in cooperator populations 298 that we see in Figure 3A. When stochastic mutations do not engender these 299 adaptations, defectors invade, and the cycle is broken. 300

While negative niche construction is necessary, we find that making the selective effects of niche construction more negative is disruptive to cooperation (Figure 6A). Instead, it is the rate at which cooperators gain adaptations that allow them to escape invasion. When the mutation rate at adaptive loci was increased, cooperation was maintained in more populations, despite the

increased opportunity for gaining deleterious mutations (Figure 6B). Taken together, these results indicate that cooperators are better able to escape defector invasion when adaptation has a clear path.

When the selective effects of niche construction are more negative, populations become more diverse. This diversity could potentially mean that when 310 a defector arises, the type that outcompetes it may already be present in the 311 population. But diversity is a double edged sword. Because a neighboring 312 patch in a diverse population is also more likely to have constructed a differ-313 ent niche. This greatly limits how quickly an adapted type can spread, since 314 the immigrant type evolved in a different niche and is therefore maladapted to 315 the new environment. Instead, when a cooperator population is homogeneous, 316 the evolutionary trajectories of its subpopulations are more aligned. As we see 317 in Figure 5, when an adapted type emerges in a more homogeneous population, 318 that type can quickly spread throughout the population, thwarting invasion 319 by isogenic defector types. 320

#### Niche Construction and the Evolution of Cooperation

In our model, cooperation is orthogonal to niche construction, which allows us to focus on hitchhiking. However, by increasing the size of the local patch, this form of cooperation could itself be seen as form of niche construction. Previous studies have more directly explored how niche construction and cooperation interact. Lehmann (2007) showed that cooperation can be favored when niche construction acted to decouple kin competition from kin selection in spatially-

structured populations. Perhaps most similar our work, Van Dyken and Wade (2012) demonstrated that when two negative niche constructing cooperative 329 behaviors co-evolve, selection can increasingly favor these traits, which were 330 disfavored when alone. In that model, "reciprocal niche construction" occurred 331 when the negative feedback created by one strategy positively influenced se-332 lection on the other, creating a perpetual cycle that maintained both forms of cooperation. Arguably, this can be viewed as an instance of hitchhiking: the 334 currently-maladaptive form of cooperation is maintained by association with the adaptive form. Outside of the context of cooperation, Laland, Odling-Smee, and Feldman have shown that niche construction can allow deleterious 337 alleles to be maintained (1996, 1999). Indeed, cooperation, especially in com-338 petition against equally-adapted defectors, can be considered deleterious.

#### 340 Evolution at Multiple Timescales

In the model described by Lehmann (2007), the selective feedbacks produced by the cooperative, niche-constructing behavior only affected future generations. Kin competition thereby was reduced, and cooperation instead benefitted descendants. Other studies, while not focused on cooperation, have similarly shown that the timescales at which niche construction feedbacks occur can strongly influence evolutionary outcomes (Laland *et al.*, 1996, 1999). This perspective is likely to be crucial for understanding the evolution of cooperative behaviors like the production of public goods.

For example, bacteria produce a host of extracellular products that scavenge

soluble iron (Griffin et al., 2004), digest large proteins (Diggle et al., 2007; Darch et al., 2012), and reduce the risk of predation (Cosson et al., 2002). 351 While many studies have focused on how the environment affects the evolution 352 of these cooperative traits, relatively few have addressed how the environmen-353 tal changes created by public goods feed back to influence evolution. In these 354 instances, environmental changes are likely to occur on different timescales than growth, which can have profound effects. For example, a multitude of 356 factors including protein durability (Brown and Taddei, 2007; Kümmerli and Brown, 2010), diffusion (Allison, 2005; Driscoll and Pepper, 2010), and re-358 source availability (Zhang and Rainey, 2013; Ghoul et al., 2014) influence 359 both the rate and the degree to which public goods alter the environment. 360 While Lehmann (2007) showed that cooperation was favored when selective 361 feedbacks act over longer timescales, niche construction may in fact hinder 362 cooperation when selection is more quickly altered. For example, when public 363 goods accumulate in the environment, cooperators must decrease production 364 to remain competitive (Kümmerli and Brown, 2010; Dumas and Kümmerli, 365 2012). This favors cooperation that occurs facultatively, perhaps by sensing 366 the abiotic (Bernier et al., 2011; Koestler and Waters, 2014) or biotic environ-367 ment (Brown and Johnstone, 2001; Darch et al., 2012).

# Cooperation and Niche Construction in Host-Symbiont Co-Evolution

In many instances where cooperation occurs, the environment is itself a biological entity. This can introduce additional evolutionary feedbacks. As the host 372 population changes, so too does selection on their symbiont populations. Here, 373 evolutionary outcomes depend greatly on the degree of shared interest between 374 the host and symbiont. For example, the cooperative production of virulence 375 factors by the human pathogen P. aeruginosa in lung infections is harmful to 376 hosts with cystic fibrosis (Harrison, 2007). Conversely, cooperative light pro-377 duction by A. fischeri is vital for the survival of its host, the Hawaiian bobtail 378 squid (Ruby, 1996). It was recently argued that incorporating the effects of 379 niche construction is critical for improving our understanding of viral evolu-380 tion (Hamblin et al., 2014) and evolution in co-infecting parasites (Hafer and 381 Milinski, 2015). Incorporating host dynamics, co-evolution, and the feedbacks 382 that they produce into models is likely to be equally important for gaining 383 an understanding of how cooperative behaviors evolve in these host-symbiont settings.

### ${f Acknowledgments}$

We are grateful to Peter Conlin, Sylvie Estrela, Carrie Glenney, and Martha Kornelius for helpful comments on the manuscript, and to Anuraag Pakanati for assistance with simulations. This material is based upon work supported by the National Science Foundation Postdoctoral Research Fellowship in Biology under Grant No. DBI-1309318 (to BDC) and under Cooperative Agreement No. DBI-0939454 (BEACON STC). Computational resources were provided by an award from Google Inc. (to BDC and BK).

- Figures Figures
- Figure 1

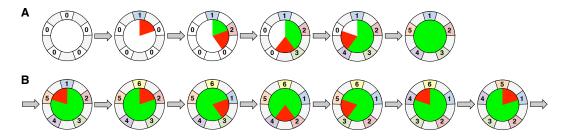


Figure 1: Adaptation to External and Constructed Environments. (A) We begin with the case with five adaptive loci (L=5) and five non-zero alleles (A = 5). All simulations are initialized with a non-adapted genotype with allele 0 at every locus (far left). Random mutation will introduce a nonzero allele, which will increase in frequency. In this example, allele 1 arises at the first locus (in the "12 o'clock" position). The rest of this schematic focuses on niche construction. Every non-zero allele at any locus influences selection at the next locus in the clockwise direction. There is a "mismatch" in this genotype (highlighted by the red sector), because the niche constructed by allele 1 at the first locus favors allele 2, not 0, at its immediate clockwise neighbor (the second locus). Once the appropriate allele arises, it will be selected. In this case, the genotype [1,2,0,0,0] receives an epsilon effect in addition to the extra delta. The "match" at the first and second locus is highlighted as a green sector. However, now there is a new mismatch between the second and third locus, which a new round of mutation and selection corrects, and so on. The green sector grows as the red sector ticks clockwise. Importantly, because A divides evenly into L, this genotype can evolve into a perfectly reinforcing sequence [1,2,3,4,5], which enjoys a maximal epsilon increment of fitness due to niche construction. (B) The case of negative niche construction is illustrated for the case of five loci (L=5) and six non-zero alleles (A=6). Here we start with a population fixed for the genotype on the far left [1,2,3,4,5]. There is a single mismatch in this genotype (highlighted by the red sector), because the niche constructed by allele 5 favors allele 6, not 1, at its immediate clockwise neighbor. If the fitter mutant [6,2,3,4,5] arises (see next genotype to the right), it will fix. (We note that the strength of selection will drop as its frequency increases). However, now there is a new mismatch in the genotype (highlighted again with a red sector). We see that correcting one mismatch generates a new mismatch. Thus, this system will never escape its mismatches—the red sector just clicks clockwise around the genome. Indeed, after six (or A) rounds of mismatch correction and generation, we have ended back where we started with the original genotype turned clockwise by one locus. Here, the adaptation to previous niche construction generates further niche construction that leads to novel adaptation.

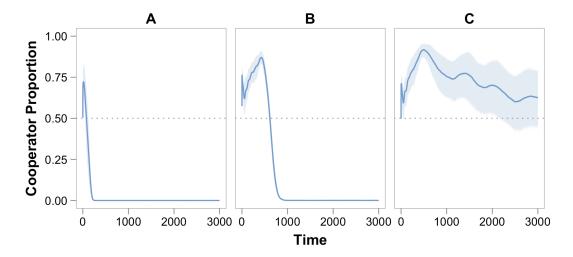


Figure 2: Adaptation, Hitchhiking, and the Evolution of Cooperation. Curves show the average cooperator proportion among replicate populations for the duration of simulations, and shaded areas indicate 95% confidence intervals. Unless otherwise noted, parameter values are listed in Table 1. (A) Without 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 3A.

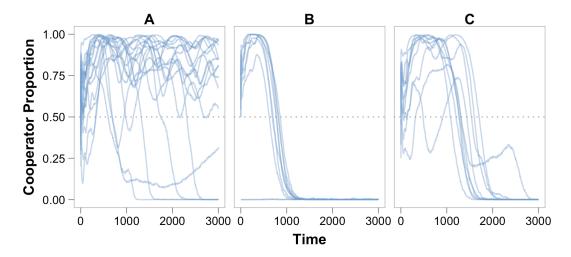


Figure 3: Niche Construction and the Evolution of Cooperation. The proportion of cooperators present in each replicate population is shown for the duration of simulations. (A) Despite some oscillations, cooperation dominates in 13 of 18 populations when niche construction affects selection. (B) When the selective effects of niche construction ( $\epsilon$ ) are removed, and the selective benefit of adaptation to the external environment ( $\delta$ ) 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).

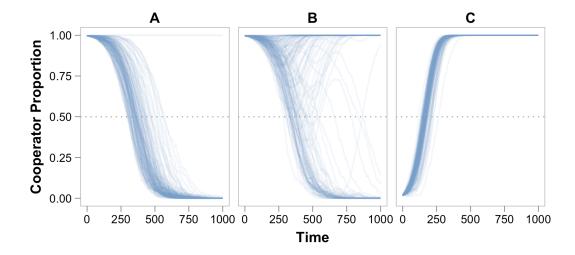


Figure 4: Niche Construction and Invasion. Curves trace the proportion of cooperators present in each replicate population for the duration of simulations (T = 1000). In each simulation, a rare type was initiated at a single patch in the center of the population lattice ( $N^2 = 121$ ). Unless otherwise 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 Figure 1) can swiftly displace defectors when isogenic defectors cannot arise or adapt via mutation.

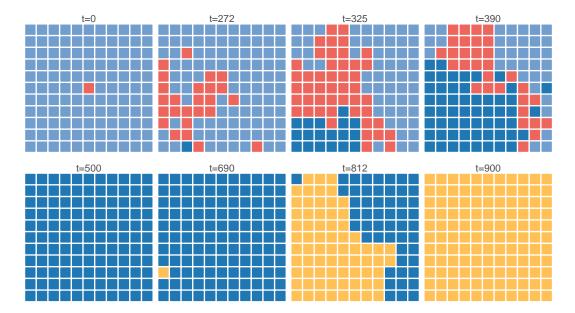


Figure 5: Cooperator Adaptation Prevents Defector Invasion. Here we depict the distribution of dominant types among subpopulations over time for one representative simulation in which isogenic defectors arise. To highlight the effects of adaptation, mutations did not occur at the cooperation locus ( $\mu_c = 0$ ). 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.

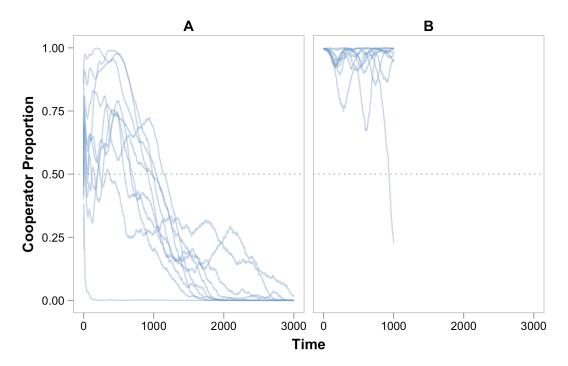


Figure 6: Negative Niche Construction and Adaptive Paths. The proportion of cooperators present in each replicate population is shown for the duration of simulations. (A) When the effects of negative niche construction are magnified, cooperators are eliminated from all replicate populations (n=10). (B) Instead, when the mutation rate at adaptive loci is increased 100-fold, cooperators remain dominant in TODO of TODO replicate populations ( $\mu_a = 0.001$ ).

Supplemental Figure 1

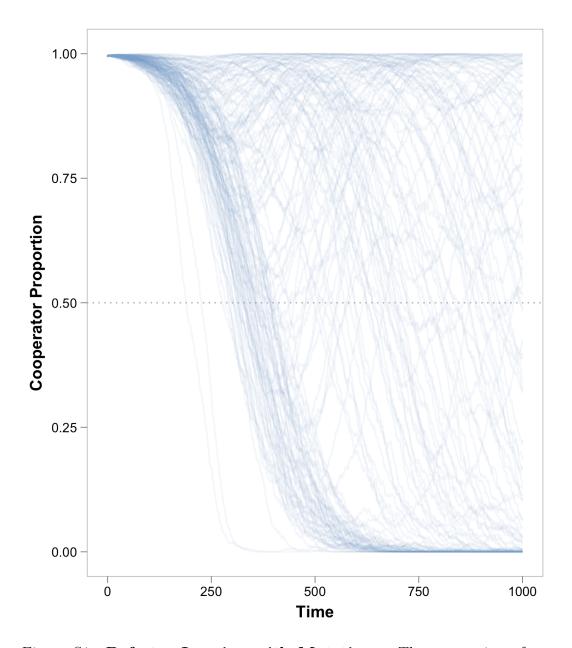


Figure S1: **Defector Invasion with Mutations.** The proportion of cooperators present in each replicate population is shown for the duration of simulations (T = 1000). When mutations occur both at the adaptive loci and the cooperation locus ( $\mu_a = \mu c = 0.00005$ ), cooperation remains dominant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

## <sup>402</sup> Supplemental Figure 2

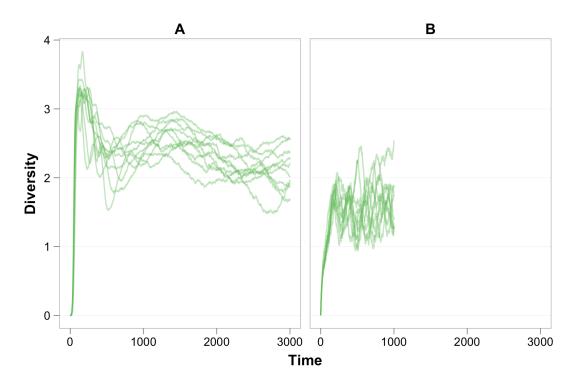


Figure S2: **TODO Diversity something.** TODO. Also explain Shannon (A) TODO. (n=TODO) (B) TODO. (n=TODO)

# Tables

Table 1: Model parameters and their value

Parameter	Description	Base Value
L	Number of adaptive loci	5
c	Fitness cost of cooperation	0.1
A	Number of alleles	6
$\delta$	Fitness benefit, nonzero alleles	0.3
$\epsilon$	Fitness benefit, sequential alleles	0.00015
z	Baseline fitness	1
$S_{min}$	Minimum subpopulation size	800
$S_{max}$	Maximum subpopulation size	2000
$\mu_a$	Mutation rate (adaptation)	$10^{-5}$
$\mu_c$	Mutation rate (cooperation)	$10^{-5}$
$N^2$	Number of patches	625
m	Migration rate	0.05
$p_0$	Initial cooperator proportion	0.5
$\sigma$	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,
- 407 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.
- 412 Modulation of Pseudomonas aeruginosa surface-associated group behaviors by
- individual amino acids through c-di-GMP signaling. Research in Microbiology,
- 414 **162**: 680–688.
- Brown, S.P. and Johnstone, R.A. 2001. Cooperation in the dark: Signalling
- 416 and collective action in quorum-sensing bacteria. Proceedings of the Royal
- 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.
- 420 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-
- 423 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
- 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:
- 435 473-484.
- 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,
- 438 **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.
- 442 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.
- 447 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-
- 450 ture, dynamics, and function using NetworkX. In: Proceedings of the 7th
- Python in Science Conference (SciPy2008), pp. 11–15.
- 452 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.
- 455 Hamilton, W.D. 1964. The genetical evolution of social behaviour I & II.
- 456 Journal of Theoretical Biology, 7: 1–52.
- 457 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.
- 459 Cold Spring Harbor Labs Journals.
- 460 Harrison, F. 2007. Microbial ecology of the cystic fibrosis lung. *Microbiology*,
- 461 **153**: 917–923.
- 462 Koestler, B.J. and Waters, C.M. 2014. Bile acids and bicarbonate inversely
- regulate intracellular cyclic di-GMP in Vibrio cholerae. Infection and Immu-
- 464 nity, **82**: 3002–3014.
- 465 Kuzdzal-Fick, J.J., Fox, S.A., Strassmann, J.E. and Queller, D.C. 2011. High
- 466 relatedness is necessary and sufficient to maintain multicellularity in Dic-
- 467 tyostelium. *Science*, **334**: 1548–1551.
- 468 Kümmerli, R. and Brown, S.P. 2010. Molecular and regulatory properties of
- 469 a public good shape the evolution of cooperation. Proceedings of the National

- 470 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.
- 477 Lehmann, L. 2007. The evolution of trans-generational altruism: Kin selection
- meets niche construction. Journal of Evolutionary Biology, 20: 181–189.
- 479 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:
- 482 Proceedings of the 9th Python in Science Conference (S. van der Walt and J.
- 483 Millman, eds), pp. 51–56.
- 484 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:
- 486 841-846.
- 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
- 489 Biology, **6**: e1000716.
- Nowak, M.A. 2006. Five rules for the evolution of cooperation. Science, 314:
- 491 1560-1563.

- Odling-Smee, F.J., Laland, K.N. and Feldman, M.W. 2003. Niche construc-
- 493 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.
- Ruby, E.G. 1996. Lessons from a cooperative, bacterial-animal association:
- The Vibrio fischeri–Euprymna scolopes light organ symbiosis. Annual Review
- of Microbiology, **50**: 591–624.
- Sinervo, B., Chaine, A., Clobert, J., Calsbeek, R., Hazard, L. and Lancaster,
- Let al. 2006. Self-recognition, color signals, and cycles of greenbeard mutu-
- alism and altruism. Proceedings of the National Academy of Sciences, 103:
- <sub>502</sub> 7372–7377.
- Van Dyken, J.D. and Wade, M.J. 2012. Origins of altruism diversity II: Run-
- <sup>504</sup> away coevolution of altruistic strategies via "reciprocal niche construction".
- 505 Evolution, **66**: 2498–2513.
- Veelders, M., Brückner, S., Ott, D., Unverzagt, C., Mösch, H.-U. and Essen,
- 507 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., Griffin, A.S. and Gardner, A. 2007. Evolutionary explanations for
- cooperation. Current Biology, 17: R661–R672.

- <sup>514</sup> Zhang, X.-X. and Rainey, P.B. 2013. Exploring the sociobiology of pyoverdin-
- producing Pseudomonas. Evolution, 67: 3161–3174.