## 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., 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 the latter case, the alleles may provide private benefits that are completely 27 independent from the public benefits of cooperation. In an asexual population 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 31 cooperator, by chance, wins the adaptive race. Hammarlund et al. (2015) recently demonstrated that in spatially-structured populations, the "Hankshaw effect" can give cooperators a substantial leg up on defectors in an adaptive race. When cooperation increases local population density, the likelihood of acquiring beneficial mutations is also increased. The cooperative trait can rise in abundance by hitchhiking along with these adaptations. 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) demonstrated that cooperation can be maintained
indefinitely 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 not simply play passive roles in environmental change and in evolution. Through their activities, their interactions
with others, and even their deaths, organisms constantly modify their envi-

environmental change alters selection, which, in turn, alters the distribution of phenotypes and their corresponding influence on the environment (Odling-Smee et al., 2003). Because of these feedback loops, populations may find themselves continually chasing beneficial mutations as their adaptive land-scape perpetually shifts beneath them.

Here, we explore whether the selective feedbacks that result from niche con-

ronment. These changes can produce evolutionary feedback loops in which

struction can prolong cooperation. We build upon the model presented by
Hammarlund et al. (2015) to allow populations to modify their local environments in ways that affect fitness. We first use this model to address whether
niche construction can extend the Hankshaw effect, allowing cooperation to
keep hitchhiking as populations continually adapt. We then focus on how
niche construction influences outcomes when isolated cooperator populations
encounter populations of defectors, either through migration or through mutations that inevitably produce defectors that share the same adaptations.

Finally, niche construction has frequently been shown to increase diversity (???). We explore whether this diversity helps or hinders the evolution of cooperation.

### 67 Methods

We develop an individual-based model in which populations of cooperators and defectors evolve and compete in a spatially-structured metapopulation (a collection of populations). Through mutations, individuals gain adaptations to their environment, which increase reproductive fitness, and allow those lineages to rise in abundance. Migration among neighboring populations allows more

We expand upon the model described by Hammarlund et al. (2015) to allow populations to modify their local environment. As this process occurs, environmental changes feed back to affect selection. We perform simulations using this model to explore how niche construction affects this adaptation process, and whether selective feedbacks allow cooperation to be maintained.

## 79 Model Description

successful lineages to spread.

### 80 Individual Genotypes and Adaptation

Each individual in a population has a genotype, which is an ordered list of L+1 integers, or *loci* (see Table 1 for model parameters and their values). Different

values at these loci represent different alleles. A binary allele at locus L+1 determines whether that individual is a cooperator (1), which carries fitness cost c, or a defector (0). Cooperation is independent from adaptation to the environment. The first L loci are adaptive loci, and are each occupied by 0 or an integer from the set  $\{1, 2, ..., A\}$ . Allele 0 represents a lack of adaptation, while a non-zero allele represents one of the A possible adaptations at that locus. Adaptations confer a fitness benefit  $\delta$ , regardless of which non-zero allele is present. We choose  $\delta > c$ , which allows a minimally adapted cooperator to recoup the cost of cooperation and gain a fitness advantage. The benefits that these adaptations engender are purely endogenous, and are not affected by the other individuals or the state of the environment.

#### 94 Niche Construction and Selective Feedbacks

- Individual fitness is also affected by the current state of the local environment.
- <sub>96</sub> Here, we represent the "niche" implicitly based on the allelic states present in
- 97 the population. As allelic states change, populations alter their environment
- 98 in different ways, creating a unique niche.

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- We use a form of density dependent selection to favors individuals that better match their niche. Specifically, the selective value of adaptive allele a at locus l increases with the number of individuals in the population that have allele a+1 at locus l+1. As a consequence, genotypes with sequentially increasing
- $_{104}$  as "circular": the selective value of an allele at locus L is affected by the allelic

allelic states will tend to evolve. We treat both adaptive loci and allelic states

composition of the population at locus 1. Similarly, the selective value of allele A at any locus increases with the number of individuals carrying allele 1 at the next locus. This circularity is represented by the function  $\beta(x, X)$ , which gives the integer that follows an arbitrary value x in the set  $\{1, 2, ..., X\}$ :

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

Here,  $\operatorname{mod}_X(x)$  is the integer remainder when dividing x by X. Thus, the selective value of adaptive allele a at locus l increases with the number of individuals that have allele  $\beta(a,A)$  at locus  $\beta(l,L)$ . The slope of this increase is  $\epsilon$ , which specifies the intensity of niche construction.

Consider a genotype g with the allelic state at locus l given by  $a_{g,l}$ ; the fitness of an individual with this genotype is defined as:

$$W_g = z + \delta \sum_{l=1}^{L} I(a_{g,l}) + \epsilon \sum_{l=1}^{L} n(\beta(a_{g,l}, A), \beta(l, L)) - ca_{g,L+1}$$
 (2)

where z is a baseline fitness 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 endogenously by adaptation ( $\delta$ ) and exogenously by its niche ( $\epsilon$ ).

Because mutations occur randomly (see below), each population will evolve different consecutive sequences. These different sequences represent the unique niches constructed by populations.

### Population Growth and the Benefit of Cooperation

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

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

During growth, individuals compete for inclusion in the resulting population.

Each individual's probability of success is determined by its fitness. The composition of a population with size P and cooperator proportion p after growth is multinomial with parameters and S(p) and  $\{\pi_1, \pi_2, \dots, \pi_P\}$ , where:

$$\pi_i = \frac{W_{\gamma(i)}}{\sum_{j=1}^P W_{\gamma(j)}} \tag{5}$$

Here,  $W_{\gamma(i)}$  is the fitness of an individual i with genotype  $\gamma(i)$  (see Equation 2).

The value  $\pi_i$  represents an individual's reproductive fitness relative to others in the population.

#### 135 Mutation

For simplicity, we apply mutations after population growth. Mutations occur independently at each locus and cause an allelic state change. At each adaptive locus, mutations occur at rate  $\mu_a$ . These mutations replace the existing allele with a random selection from the set  $\{0\} \cup \{1, 2, ..., A\}$ . Note that this allows for the possibility of an allele replacing itself, thus slightly reducing the effective mutation rate. 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.

### 144 Migration

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

### Metapopulation Initialization and Simulation

Metapopulations are initiated in a state that follows an environmental change, which leaves most patches empty. First, populations are seeded at all patches with cooperator proportion  $p_0$  and grown to density  $S(p_0)$ . An environmental

challenge is then introduced, which subjects the population to a bottleneck. 156 For each individual, the probability of survival is  $\mu_t$ , which represents the 157 likelihood that tolerance arises via mutation. Because individuals have not 158 yet adapted to this new environment, the allelic state of each individual's 159 genotype is 0 at each adaptive locus. Following initialization, simulations 160 are run for T cycles, where each discrete cycle consists of population growth, 161 mutation, and migration. At the end of each cycle, populations are thinned to 162 allow for growth in the next cycle. Each individual persists with probability 163 d, regardless of allelic state.

### Source Code and Software Environment

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 follow the evolution of cooperation in a metapopulation consisting of populations connected by spatially-limited migration. Individuals compete in these populations by gaining a limited number of adaptations that confer fit-

ness benefits. While cooperation does not directly affect the selective value of these adaptations, cooperation can have indirect effects on the adaptive pro-177 cess. Specifically, cooperation increases population density. As a result, larger 178 populations of cooperators experience more mutational opportunities to gain 179 adaptations. Cooperation can hitchhike along with these adaptations, which 180 compensate for the cost of cooperation. During this process, populations alter 181 their local environments, which, in turn, influences selection. Here, we explore 182 how niche construction affects the evolution of cooperation in the simulation environment defined by the parameter values listed in Table 1. 184

### 185 Cooperation Persists with Niche Construction

Without any opportunity for adaptation (L=0), cooperators are swiftly elim-186 inated in competition with defectors (Figure 1A). Despite an initial lift due 187 to increased productivity, the cost of cooperation becomes disadvantageous as 188 migration mixes the initially isolated populations. When there are opportuni-189 ties for adaptation (L=5) but no niche construction  $(\epsilon=0)$ , cooperators are 190 maintained transiently (Figure 1B). Here, larger cooperator populations can 191 more quickly adapt to their environment. As previously described by Hammar-192 lund et al. (2015), however, cooperation is subsequently lost once populations 193 become fully adapted to their environment. Once this has occurred, adapted 194 defectors that arise via mutation at the cooperation locus have a selective 195 advantage and drive cooperators from the population. However, when niche 196 construction creates selective feedbacks, cooperation persists in 13 of 18 repli-197

cate populations (Figure 2A).

# Fitness Increases Alone do not Support Persisting Cooperation

In the model, both adaptation and niche construction contribute to an individual's fitness. To determine whether cooperation is maintained solely because 202 to the larger selective values, we performed simulations in which the selective 203 contributions of niche construction were removed ( $\epsilon = 0$ ), and we instead in-204 creased the fitness benefits conferred by adaptation ( $\delta = 0.6$ ). In doing so, we 205 conservatively represent the selective effects of niche construction, as fitness 206 benefits of this magnitude would only be given for sequential allelic states that 207 are fixed in full populations. We find that simply increasing selective values 208 does not allow cooperators to persist (Figure 2B).

# Negative Niche Construction is Critical to Cooperator Persistence

Negative niche construction can occur in our model due to the selection for sequentially-increasing allelic states and the circular arrangement of these alleles. When this occurs, adaptations at one locus reduce the selective effects at another locus, and thus negatively affect fitness. This occurs when when the genome length (L) is not evenly divided by the number of adaptive alleles (A), which makes it impossible to evolve sequentially increasing allelic states. When negative niche construction is removed (L = 5, A = 5), cooperators are again driven to extinction after an initial lift in abundance (Figure 2C).

### 220 Selective Feedbacks Limit Defector Invasion

The adaptation resulting from selective feedbacks can limit invasion by de-221 fectors, which arise either through immigration from neighboring patches or 222 through mutation from a cooperator ancestor. The latter pose a particularly 223 challenging threat, as they are equally adapted, yet do not incur the cost of cooperation. When homologous defectors (i.e., defectors with identical stress 225 loci) are introduced as a single population in the center of an 11x11 metapopulation of cooperators, they quickly spread (Figure 3A). However, when res-227 ident cooperators can adapt and respond to defector invasion, the situation 228 improves dramatically, allowing cooperation to evade defector invasion in 91 229 of 160 replicate populations (Figure 3B). Figure 4 depicts one such instance 230 where cooperators gained an adaptation that stopped and eliminated invading 231 defectors. We further highlight this process in Figure 3C, where an adapted 232 cooperator genotype can rapidly invade a population of defectors. 233

### Diversity Hampers the Evolution of Cooperation

TODO: defector can invade a diverse population of cooperators, while adaptation to a matching defector can't spread to stop invasion.

### 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-239 nard Smith and Haigh, 1974). In a process termed the "Hankshaw effect", 240 Hammarlund et al. (2015) recently demonstrated that cooperative behaviors 241 can prolong their existence by actively increasing their likelihood of hitchhiking 242 with a beneficial trait. While this process does favor cooperation in the short 243 term, it eventually reaches a dead end; when the opportunities for adaptation 244 are exhausted, and cooperators can no longer hitchhike, they face extinction. 245 In this work, we have considered whether niche construction can maintain 246 cooperation indefinitely. When niche construction occurs, cooperation can indeed persist (Figure 2A). In 248 our model, niche construction introduces additional selective effects that could 249 influence the evolutionary process, leading to a more pronounced Hankshaw 250 effect. However, simply raising the selective benefits provided by adaptations 251 does not prolong cooperation (Figure 2B), and indicates that niche construc-252 tion plays an important role. 253 We find that cooperator success is due to negative niche construction. Without adaptive opportunities, adaptation eventually slows. Once this occurs, cooperators face the threat of invasion by defectors that arise de novo through mutation. Since these defectors are equally adapted but do not bear the cost of cooperation, they are favored by selection, and quickly drive cooperators to extinction. By reducing fitness, negative niche construction creates adaptive

opportunities. These opportunities can allow cooperators to resist invasion by defectors, even when defectors are equally adapted (Figure 3B). Here we observe another facet of the Hankshaw effect: because populations of cooperators are larger, they are better able to respond to the adaptive opportunities that result from negative niche construction.

TODO: diversity results TODO: references about diversity

In our model, cooperation and niche construction are orthogonal, which al-266 lows us to focus on hitchhiking. However, the form of cooperation used in 267 this model could itself be seen as a niche constructing behavior. Explicitly 268 modeling this cooperative behavior, which is akin to the production of pub-269 lic goods, would likely yield additional insights into the relationship between 270 cooperation and niche construction. For example, previous work has shown 271 that niche construction can favor deleterious alleles (Laland et al., 1996, 1999). 272 Cooperation, especially in competition against equally-adapted defectors, can be considered deleterious, so introducing selective feedbacks from cooperation 274 could further bolster cooperation. Van Dyken and Wade (2012) showed that when two cooperative behaviors co-evolve and niche construction feedbacks benefit the other type, niche construction can increasingly favor these traits, 277 which were otherwise disfavored when alone. Arguably, this can be viewed 278 as another instance of hitchhiking: the maladaptive form of cooperation is 279 maintained by association with the adaptive form. However, negative niche 280 construction then reverses these roles and perpetuates the cycle. 281

By their very nature, public goods benefit populations by making their envi-

ronment more hospitable (West et al., 2007a). For example, bacteria produce a host of extracellular products that scavage soluble iron (Griffin et al., 2004), 284 digest large proteins (Diggle et al., 2007; Darch et al., 2012), and reduce the 285 risk of predation (Cosson et al., 2002). While many studies have explored 286 how the environment affects the evolution of cooperative behaviors such as 287 the production of these public goods, relatively few have examined how the 288 resulting selective feedbacks influence evolution as public goods modify the 289 environment. In these instances, the timescale at which the environment is likely to be decoupled from the timescale at which reproduction occurs. These 291 differences can have profound effects. For example, a multitude of factors in-292 cluding protein durability (Brown and Taddei, 2007; Kümmerli and Brown, 293 2010), diffusion (Allison, 2005; Driscoll and Pepper, 2010), and resource avail-294 ability (Zhang and Rainey, 2013; Ghoul et al., 2014) influence both the rate 295 and the degree to which public goods alter the environment. Lehmann (2007) 296 demonstrated that cooperative, niche constructing behaviors can be favored 297 when they affect selection for future generations. When this occurs, conflict 298 among contemporary kin is reduced. The evolutionary inertia that this cre-299 ates, however, may ultimately work against cooperators. When public good 300 accumulates in the environment, cooperators must decrease production to re-301 main competitive (Kümmerli and Brown, 2010; Dumas and Kümmerli, 2012). 302 This favors cooperation that occurs facultatively, perhaps by sensing the abi-303 otic (Bernier et al., 2011; Koestler and Waters, 2014) or biotic environment 304 (Brown and Johnstone, 2001; Darch et al., 2012). 305

In many instances of cooperation, the environment is itself a biological entity,

which can produce additional evolutionary feedbacks. As the host population changes, so too does selection on their symbiont populations. Here, evolution-308 ary outcomes depend greatly on the degree of shared interest between the host 309 and symbiont. For example, the cooperative production of virulence factors by 310 the human pathogen P. aeruqinosa in lung infections is harmful to those with 311 cystic fibrosis (Harrison, 2007). Conversely, cooperative light production by A. fischeri is vital for the survival of its host, the Hawaiian bobtail squid (Ruby, 313 1996). It was recently argued that incorporating the effects of niche construction is critical for improving our understanding of viral evolution (Hamblin et 315 al., 2014) and evolution in co-infecting parasites (Hafer and Milinski, 2015). 316 Incorporating host dynamics, co-evolution, and the feedbacks that they pro-317 duce into models is likely to be equally important for gaining an understanding 318 of how cooperative behaviors evolve in these host-symbiont settings.

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## Figures 527

## Figure 1

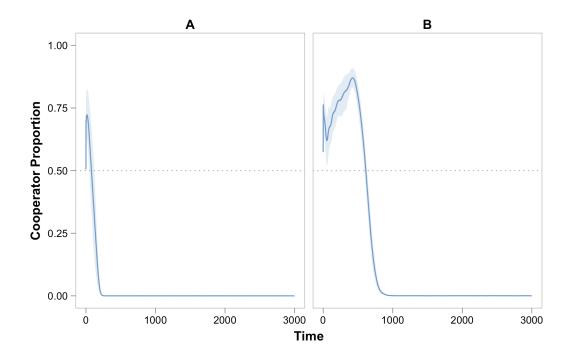


Figure 1: Adaptation, Hitchhiking, and the Evolution of Cooperation. The proportion of cooperators present in the population is shown for the duration of simulations. Curves show the average among replicate populations, and shaded areas indicate 95% confidence intervals. Unless otherwise noted, parameter values are listed in Table 1. (A) Without any opportunity to adapt (L, the number of adaptive loci, is zero), cooperation is quickly lost. (B) When adaptation can occur (L=5), but populations do not alter their environment  $(\epsilon, \text{the intensity of niche construction, is zero)}$ , cooperation hitchhikes along with adaptions, allowing cooperators to temporarily rise in abundance before eventually going extinct.

### Figure 2

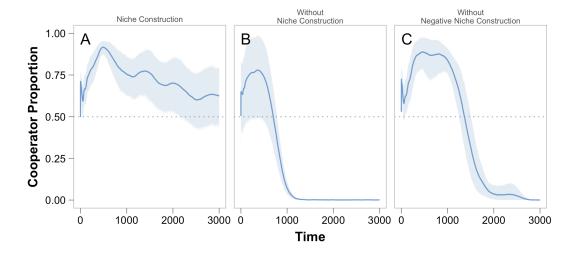


Figure 2: Niche Construction and the Evolution of Cooperation. The proportion of cooperators present in the population is shown for the duration of simulations. Curves show the average among replicate populations, and shaded areas indicate 95% confidence intervals. (A) Niche construction enables cooperation to be maintained indefinitely. In the majority of populations, cooperation remained as the dominant strategy. (B) When niche construction is removed and the fitness benefit of adaptation is increased as compensation ( $\epsilon = 0, \delta = 0.6$ ), adapted defectors arise and drive cooperators to extinction. (C) Without negative niche construction, cooperation is not maintained (A = 5). Here again, cooperators are at a selective disadvantage against equally-adapted defectors.

### Figure 3

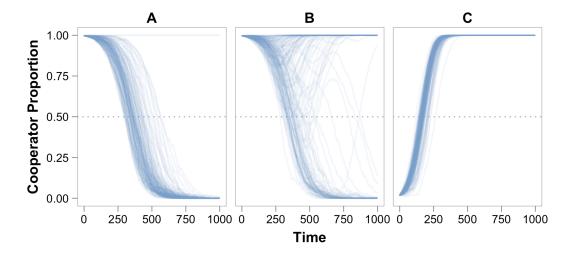


Figure 3: Niche Construction and Invasion. Curves trace the proportion of cooperators present in the population for the duration of 160 replicate simulations. These experiments examine whether a cooperator or defector strategy can invade when initiated at a single population in the center of the metapopulation lattice ( $N^2 = 121$ ). Unless otherwise noted, we disable mutations  $(\mu_a = 0, \mu_c = 0)$  to focus on the dynamics of invasion. This limitation is removed in the results shown in Figure SX. (A) When cooperators and defectors are matched (i.e., genotypes [1,2,3,4,5]) and adaptation can not occur, defectors quickly drive cooperators to extinction due to the cost of cooperation. Defectors were stochastically eliminated in 2 replicate populations. (B) However, the adaptive opportunities produced by negative niche construction can allow cooperators to resist invasion by initially-matching defectors. Here, cooperation persisted in the majority of populations ( $\mu_a = 0.00005$ , the base mutation rate). (C) Here we demonstrate that these adaptations can enable an adapted cooperator (genotype [1,2,3,4,6]) to displace a population of defectors when defectors can not arise or adapt via mutation.

### Figure 4

TODO: snapshots of cooperators adapting to thwart defector invasion

# Figure 5

<sup>334</sup> TODO: A: defector invading diverse C popuation, B: Adapted cooperators can

 $_{335}$  not spread to resist defector invasion.

# 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, nonzero alleles	0.3
$\epsilon$	Fitness benefit, sequential alleles	0.00015
z	Baseline fitness	1
$S_{min}$	Minimum population size	800
$S_{max}$	Maximum population size	2000
$\mu_a$	Mutation rate (adaptation)	$10^{-5}$
$\mu_c$	Mutation rate (cooperation)	$10^{-5}$
$N^2$	Number of metapopulation sites	625
m	Migration rate	0.05
$p_0$	Initial cooperator proportion	0.5
$\mu_t$	Mutation rate (tolerance to new environment)	$10^{-5}$
d	Population dilution factor	0.1
T	Number of simulation cycles	1000

### References

- Allison, S.D. 2005. Cheaters, diffusion and nutrients constrain decomposition
- by microbial enzymes in spatially structured environments. Ecology Letters,
- 340 **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,
- **162**: 680–688.
- Brown, S.P. and Johnstone, R.A. 2001. Cooperation in the dark: Signalling
- and collective action in quorum-sensing bacteria. Proceedings of the Royal
- 350 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.
- <sup>353</sup> 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-
- 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
- 360 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
- 367 cheats in bacterial metapopulations. Journal of Evolutionary Biology, 25:
- 368 473–484.
- 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,
- 371 **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.
- 380 Hafer, N. and Milinski, M. 2015. When parasites disagree: Evidence

- for parasite-induced sabotage of host manipulation. Evolution, doi:
- 382 10.1111/evo.12612.
- Hagberg, A.A., Schult, D.A. and Swart, P.J. 2008. Exploring network struc-
- 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.
- Hamilton, W.D. 1964. The genetical evolution of social behaviour I & II.
- 390 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.
- <sup>393</sup> Cold Spring Harbor Labs Journals.
- Harrison, F. 2007. Microbial ecology of the cystic fibrosis lung. *Microbiology*,
- 395 **153**: 917–923.
- Koestler, B.J. and Waters, C.M. 2014. Bile acids and bicarbonate inversely
- regulate intracellular cyclic di-GMP in Vibrio cholerae. Infection and Immu-
- *піту*, **82**: 3002–3014.
- Kuzdzal-Fick, J.J., Fox, S.A., Strassmann, J.E. and Queller, D.C. 2011. High
- 400 relatedness is necessary and sufficient to maintain multicellularity in Dic-
- 401 tyostelium. Science, **334**: 1548–1551.
- Kümmerli, R. and Brown, S.P. 2010. Molecular and regulatory properties of

- 403 a public good shape the evolution of cooperation. Proceedings of the National
- 404 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
- 409 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:
- Proceedings of the 9th Python in Science Conference (S. van der Walt and J.
- 417 Millman, eds), pp. 51–56.
- Morgan, A.D., Quigley, B.J.Z., Brown, S.P. and Buckling, A. 2012. Selection
- 419 on non-social traits limits the invasion of social cheats. Ecology Letters, 15:
- 420 841-846.
- 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
- 423 Biology, **6**: e1000716.
- Nowak, M.A. 2006. Five rules for the evolution of cooperation. Science, 314:

- 425 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.
- 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,
- 434 L. et al. 2006. Self-recognition, color signals, and cycles of greenbeard mutu-
- alism and altruism. Proceedings of the National Academy of Sciences, 103:
- 436 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".
- 439 Evolution, **66**: 2498–2513.
- Veelders, M., Brückner, S., Ott, D., Unverzagt, C., Mösch, H.-U. and Essen,
- 441 L.-O. 2010. Structural basis of flocculin-mediated social behavior in yeast.
- 442 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.

- 447 The social lives of microbes. Annual Review of Ecology, Evolution, and Sys-
- 448 tematics, **38**: 53–77.
- West, S.A., Griffin, A.S. and Gardner, A. 2007b. Evolutionary explanations
- for cooperation. Current Biology, 17: R661–R672.
- <sup>451</sup> Zhang, X.-X. and Rainey, P.B. 2013. Exploring the sociobiology of pyoverdin-
- producing Pseudomonas. Evolution, 67: 3161–3174.