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1 **Non-linear frequency-dependent selection promotes long-term**
2 **coexistence between bacteria species**

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16 Supplementary material: Detailed methods, appendix on the model, Sup Table S1, Fig S1, S2
17 available on HAL

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26 **Abstract**

27 Negative frequency-dependent selection (NFDS) is an important mechanism for species
28 coexistence and for the maintenance of genetic polymorphism. Long-term coexistence
29 nevertheless requires NFDS interactions to be resilient to further evolution of the interacting
30 species or genotypes. For closely related genotypes, NFDS interactions have been shown to be
31 preserved through successive rounds of evolution in coexisting lineages. On the contrary, the
32 evolution of NFDS interactions between distantly related species has received less attention. Here,
33 we tracked the coevolution of *Escherichia coli* and *Citrobacter freundii* that initially differ in their
34 ecological characteristics. We showed that these two bacterial species engaged in an NFDS
35 interaction particularly resilient to further evolution: despite a very strong asymmetric rate of
36 adaptation, their coexistence was maintained owing to an NFDS pattern where fitness increases
37 steeply as the frequency decreases towards zero. Using a model, we showed how and why such
38 NFDS pattern can emerge. These findings provide a robust explanation for the long-term
39 maintenance of species at very low frequencies.

40 **Keywords**

41 Experimental coevolution, *Escherichia coli*, *Citrobacter freundii*, Nalidixic acid, polymorphism,
42 negative frequency dependent selection, NFDS

43 **Introduction**

44 Astounding biodiversity can be observed at different scales: from locus to species, from newly
45 emerging polymorphism to trans-specific polymorphisms maintained for millions of years (e.g. in
46 (Devier et al. 2009; van Diepen et al. 2013). Understanding the maintenance of such diversity
47 among species (coexistence in communities) and within species (genetic polymorphism) is a long-
48 standing question in ecology and evolution. Various mechanisms can explain stable coexistence
49 by a balance between selection and other forces, such as migration or mutation (listed in Débarre
50 and Lenormand 2011). However, one of the most effective ways to maintain long-term
51 coexistence or polymorphism is when selection itself operates in a frequency-dependent manner,
52 favoring rare types. With such negative frequency-dependent selection (NFDS), by definition, a
53 small frequency perturbation below (resp. above) the equilibrium frequency leads to positive
54 (resp. negative) selection, bringing the system back to its equilibrium point and ensuring stability
55 (Lewontin 1958; Haldane and Jayakar 1963; Ayala and Campbell 1974; Bell 2008; Felsenstein
56 2017). Apart from selection on alleles caused by overdominance in diploids, NFDS can emerge
57 from a diversity of underlying ecological mechanisms (see Table S1). Demonstrating NFDS in field
58 or in laboratory experiments requires measuring how relative fitness varies when the competing
59 types are manipulated to be at different frequencies (we term “NFDS pattern” the relationship
60 between the selection coefficient and the frequency). Such findings are usually taken as a strong
61 argument in favor of long-term coexistence, as long as environmental conditions remain
62 unchanged (e.g. in Turner et al. 1996; Gigord et al. 2001; Weeks and Hoffmann 2008; Takahashi
63 and Kawata 2013; Healey et al. 2016). However, NFDS patterns may evolve, which may
64 compromise coexistence, even under constant environmental conditions. For instance, when two
65 types are maintained by NFDS, each can still independently adapt to the surrounding
66 environmental conditions and to the presence of the other type, hence putatively disrupting NFDS
67 interactions.

68 The evolution of NFDS patterns and long-term coexistence have been intensively studied
69 particularly for interactions within species. The long-term persistence of polymorphic alleles in
70 sexual species has been repeatedly demonstrated and can last hundreds of millions of years
71 (Takahata and Nei 1990; Devier et al. 2009; Karasov et al. 2014; Těšický and Vinkler 2015). Many
72 studies using experimental evolution on bacteria have demonstrated that emerging
73 polymorphisms can arise, evolve and reach metastable equilibria (Turner et al. 1996; Rainey and
74 Travisano 1998; Rozen and Lenski 2000; Friesen et al. 2004; Rozen et al. 2007; Blount et al. 2008,
75 2012; Plucain et al. 2014; Maddamsetti et al. 2015; Healey et al. 2016; Good et al. 2017). In those
76 cases, coexistence can be disrupted as beneficial mutations accumulate in one competing lineage
77 (Maddamsetti et al. 2015) or be maintained in the very long-term owing to successive rounds of

78 evolution in the two diverging ecotypes (Le Gac et al. 2012). The evolution of NFDS among
79 divergent species has received, comparatively, much less attention. There are however reasons to
80 expect a different regime for long-term coexistence in this case.

81 At first sight, the mechanism of negative frequency dependence applies equally well to alleles at a
82 locus within a sexual species as it does to different clones in an asexual species or to different
83 species in a community. And indeed, the basic theoretical models are virtually indistinguishable
84 in these cases (Levin 1988; Mazancourt and Dieckmann 2004). However, there might be an
85 important difference between these situations in terms of long-term coexistence. In a sexual
86 species, alleles maintained polymorphic at a locus by NFDS are likely to share the same genetic
87 background, except perhaps for the portion of the genome in the close vicinity of that locus. Hence
88 the evolution of the NFDS pattern is expected to be limited since it will mostly occur by the
89 occurrence of new alleles at that locus (e.g. for self-incompatibility, Castric and Vekemans 2004,
90 or mating type loci, Billiard et al. 2011), as illustrated in Figure 1a. Polymorphic clones *de novo*
91 emerging in an asexual species also share similar backgrounds, but, in that case, coexistence is
92 expected to be less stable compared to the previous case, as NFDS patterns may evolve due to
93 mutations and frequency variation at a large number of loci, in entire genomes evolving
94 independently. However, the initial similarity of the two types will be unlikely to cause fast NFDS
95 destabilization (Figure 1b). Divergent species engaging new NFDS interactions are expected to be
96 the least prone to long-term coexistence since such genetically and phenotypically different types
97 may immediately exhibit differences in their rate of adaptation to their common environment,
98 which could quickly destabilize their coexistence (Figure 1c). Overall, because of intercept
99 variation of the NFDS pattern, we would expect polymorphism maintained by NFDS to persist
100 longer among alleles in a sexual species than among species, and longer between recently-
101 diverged than anciently-diverged species.

102 The shape and slope of the NFDS pattern may also evolve. In the three situations mentioned above
103 and illustrated in Figure 1a,b,c, the evolution of a steeper NFDS pattern will promote long-term
104 coexistence as with a larger slope, larger changes in intercept are required to selectively eliminate
105 one type (Figure 1d). Ultimately, it is the relative evolution of the intercept and slope of the NFDS
106 pattern that will dictate long-term coexistence. While the variation in intercept may be expected
107 to differ for the different cases mentioned above (alleles, recently-diverged, anciently-diverged
108 species), there are no clear predictions for the slope. As for most situations of specialization, the
109 relative availability of the different types of mutations (on intercept or slope) depends mostly on
110 the shape of the underlying trade-offs that are difficult to predict (Lenormand 2012). Some of this
111 variation might depend on the age of the NFDS relationship. Ancient coexistence situations are
112 likely to involve interaction-stabilizing traits and relatively steeper NFDS slopes (as they have

113 been maintained for a long time) than more recent NFDS interactions. However, the current traits
114 of coexisting species might have been shaped by multiple interactions with other functionally-
115 related species in the past, even if the observed NFDS interaction is recent and involves new
116 partners. In other words, the degree to which divergent species are expected to maintain long-
117 term coexistence in an NFDS interaction is difficult to predict. For these different reasons,
118 expectations based on the wealth of knowledge accumulated for NFDS among alleles within sexual
119 species and among emerging asexual lineages is unlikely to be sufficient to draw global,
120 interspecific conclusions about the role of NFDS for long-term coexistence.

121 To address this question, we study the long-term coexistence between two bacteria species,
122 *Escherichia coli* (hereafter *E. coli* or *E*) and *Citrobacter freundii* (hereafter *C. freundii* or *C*) grown
123 on a medium with two carbon sources, glucose and citrate. *E* is particularly efficient at using
124 glucose, but cannot take up citrate under aerobic conditions, while *C* can use both carbon sources.
125 As we show, *E* and *C* can coexist by NFDS. We investigate the evolution of their NFDS patterns in
126 multiple replicates in a ca. 900 generations experimental evolution setting. In particular, we
127 investigate whether different abiotic conditions (different concentration of an antibiotic, nalidixic
128 acid) change long-term coexistence, as would be expected if different environments represent
129 different adaptive challenges to the two coevolving species.

130 **Material and methods**

131 **Strains and experimental evolution**

132 In our experiment, NFDS occurs between one Lenski's long term experiment derived *E. coli* strain
133 (Lenski and Travisano 1994) (*E*) constitutively expressing yellow fluorescent proteins (YFP) that
134 has been previously evolved in the presence of nalidixic acid (Nal) (Gallet et al. 2012; Harmand et
135 al. 2017, 2018) and one *C. freundii* strain (*C*) at very low starting frequency resulting from
136 contamination of *E* glycerol stocks. Hence, *E* is well adapted to the experimental conditions in the
137 absence of *C* whereas these conditions are new for *C*. Bacterial populations were propagated
138 through daily 100-fold dilutions (Sup. Methods) into fresh Davis minimal medium containing 250
139 µg/mL of glucose and Nal at six different concentrations: 0, 3, 8, 20, 100 and 200 µg/mL (referred
140 below to as Nal0, Nal3, etc.). We followed the frequency of the coevolving species in 44
141 coevolutionary replicates (hereafter CRep, 8 per dose of Nal, and 4 in absence of Nal). The
142 frequencies of *E* and *C* were estimated for each CRep at ca. 0, 200, 550 and 870 generations using
143 flow cytometry. See supplementary methods for details about the coevolution protocol and
144 frequency measures. *E* and *C* were isolated based on phenotypic differences between the two
145 species. Their growth patterns on glucose, on different doses of citrate or on filtrate of medium

146 where the other species had grown were measured following optical density using standard
147 methods (Sup. Methods).

148 **NFDS patterns and fitness measures**

149 Two CRep from Nal20 (CRep20A and CRep20B) were selected to investigate detailed variations
150 in the NFDS patterns during coevolution. Six independent clones of both *C* and *E* type were
151 isolated from those CRep at time 0, 214 and 870 generations and mixed to constitute *E* and *C* lines
152 representative of each time point. Variations of the NFDS patterns were investigated by
153 performing five different series of competitions: 1) *E* against *C* at time zero (*E0* against *C0*), 2) *E*
154 against *C* at 214 generations (*E2* against *C2*), 3) *E* against *C* at 870 generations (*E8* against *C8*), 4)
155 *E0* against *C8*, and 5) *E8* against *C0*. The first three series should indicate whether the interaction
156 changed during coevolution, while the last two should indicate how each species contributed to
157 that change. Separate overnight cultures of the *E* and *C* lines were mixed with a large range of
158 initial frequencies and mixes were incubated for 24h in the same conditions as the experimental
159 evolution. Selection coefficients per generation were calculated in a standard way, as in Harmand
160 *et al.* (2018), eq. (1), using frequency variation over 24h (supp. methods).

161 **Results**

162 **NFDS patterns evolved rapidly and consistently across abiotic conditions**

163 The two species were still coexisting in 39 CRep out of 44 at generations 200, 550 and 870 (Fig.
164 2). The frequency of *E* decreased very consistently across replicates. This decrease was also
165 consistently (much) stronger for increasing Nal concentrations (Fig. 2). A clear NFDS pattern was
166 present at generations 0 and 870 for CRep20A and CRep20B (Fig. 3). Remarkably, these NFDS
167 patterns were highly nonlinear, with a steep fitness change near frequency = 0 or 1. In the five
168 remaining CRep (one at Nal0, two at Nal3, one at Nal100 and one at Nal200), *C* was never detected,
169 and probably did not establish from the start (supp. methods).

170 Fig. 3 shows that the NFDS patterns changed dramatically along coevolution in CRep20A and
171 CRep20B replicates investigated in detail. Their shape was not strongly changed between initial
172 and final time, but was shifted significantly downwards (which is analogous to a change in
173 intercept as illustrated in Fig. 1). Consistent with this shift, *E* frequency declined in these
174 replicates. In other replicates, where we did not investigate the NFDS pattern, *E* frequency also
175 declined regularly. This decline is consistently more pronounced at increasing Nal concentrations

176 (Fig. 2). CRep20A and CRep20B do not particularly stand out: they are just two replicates among
177 many others showing a very similar trend in frequency variation.

178 **Evolution of NFDS pattern is mostly driven by the adaptation of *C. freundii***

179 We performed competitions at generations 0, 214 and 870 as well as time-shifted competitions in
180 order to assign specific patterns of variation in NFDS patterns to *E* or *C* evolution. Competitions
181 between *E0* and *C8* should indicate whether the evolution of *C* was responsible for the change in
182 NFDS pattern, and the reciprocal competition should measure how much change of the pattern
183 was due to the evolution of *E*. Fig. 3 shows that *E8-C0* pattern is nearly identical to the *E0-C0*
184 pattern, and that *E0-C8* patterns are nearly identical to the *E8-C8* pattern. This indicates that *E* did
185 not evolve much and therefore that nearly all the evolutionary changes modifying the NFDS
186 pattern occurred in *C*. The growth curves of isolated *C* and *E* lines confirm that *C* was adapting
187 rapidly during coevolution. In particular, these growth curves show a reduction in *C* lag time for
188 citrate consumption (CRep20A and B, Fig. 4) and improved growth on glucose (CRep20B, Fig. 4).
189 In contrast, *E* showed no obvious adaptation to the abiotic conditions. Hence, the variation of the
190 NFDS is almost entirely due to the evolution of *C*.

191 The shapes of the NFDS patterns at initial and final time are very similar (Fig. 3). The main
192 difference is an overall shift downwards, as would be expected if *C* accumulated many more
193 unconditionally beneficial mutations over this period. Here we refer as unconditionally beneficial
194 to mutations that confer the same fitness advantage at all frequencies. However, this
195 interpretation is not correct. At intermediate time, the NFDS patterns show a very different shape
196 (see NFDS patterns at generation 214 in Fig. 5). They are only shifted downwards for large starting
197 frequencies of *E*, not for small frequencies. The modification of the NFDS pattern therefore
198 occurred in a stepwise fashion. *C* first acquired mutations that presented an advantage when *C*
199 was rare. In a second step, other beneficial mutations occurred that were beneficial in a higher
200 range of *C* frequencies. The same pattern holds very consistently for CRep20A and CRep20B.

201 **Ecological context of the frequency-dependent selection**

202 *C* presents a diauxic growth (i.e. a growth curve with two exponential phases on two different
203 resources), but *E* does not (Fig 4). *C* overall carrying capacity increases linearly with citrate
204 concentration, confirming that *C* consumes citrate. Citrate concentration has almost no effect on
205 the first growth phase of *C*, indicating that *C* consumes glucose first (Fig S2). *E* maximal carrying
206 capacity occurs at 100% citrate (i.e. the concentration corresponding to that of the experimental
207 evolution medium). At other citrate concentrations, this carrying capacity is smaller, but the
208 difference is modest in all cases. This small variation does indicate that citrate plays a role in *E*

209 metabolism, but not as a resource (e.g. pH buffering). The modest effect of citrate concentration
210 on *E* carrying capacity did not change between generation 0 and 870 (all these results are shown
211 in Fig S2). At the start of the coevolution, *C* growth achieved on glucose is always lower than *E*
212 growth, indicating that *C* has a lower efficiency on glucose. To achieve a larger overall carrying
213 capacity than *E*, *C* requires some citrate (at least ~one quarter of the concentration in the DM250,
214 Fig S1a). At generation 870, these comparisons cannot be made easily as the diauxic shift for *C*
215 cannot be well identified (see for instance Fig 4). In most *C* lines, the lag time of switching from
216 glucose to citrate resource becomes hardly detectable. Finally, *E* did not grow in the filtrated
217 medium produced after a growth cycle of *C* (while *C* could use citrate in the reverse situation,
218 Suppl. results.). This indicates that they do not excrete byproducts that are left undigested. This
219 observation does not rule out the possibility that some metabolites are excreted and then
220 reabsorbed and consumed later (e.g. acetate can be produced and temporarily excreted).

221 **Modelling non-linear NFDS patterns**

222 We investigated theoretically the NFDS patterns that can be obtained in competition scenarios of
223 two strains with two resources and diauxy. We detail this model in Appendix 1 and Fig S1 presents
224 it graphically. Figure 6a-c illustrates the NFDS patterns expected under one-niche, two-niches and
225 three-niches models. In the one niche model, the NFDS patterns emerge from a differential
226 specialization on growth rate and conversion efficiency for two types exploiting the same
227 resource. The two niches model corresponds to the Levene's model (1953) with two types
228 competing on two resources. Finally, in the three niches model, there are two resources/niches,
229 but a third one is created by an anticipated metabolic switch of one type to the second resource
230 before the first resource is depleted (i.e. the lag phase on the second resource starts earlier by a
231 quantity Δt). The other type then has a 'private' third niche corresponding to the leftover of the
232 first resource.

233 Interestingly, the NFDS patterns of this third model are generally consistent with the overall "S-
234 shape" of the NFDS patterns observed with strong curvature at extreme frequencies (Fig. 3). The
235 model can further indicate ways to mimic the observed NFDS pattern evolution through time (Fig
236 6d). Initially, increasing *C* citrate conversion efficiency can shift down the NFDS pattern at high
237 frequency of *E* (as observed at generation 214 for CRep20A and CRep20B). Then, *C* may evolve to
238 consume more glucose, switching to citrate closer to the time of glucose depletion (smaller Δt).
239 This change favors *C* when frequent while maintaining a small 'private niche' for *E*, and an upward
240 NFDS curvature at very low frequencies of *E* (as observed at generation 870 for CRep20A and
241 CRep20B), hence preserving *E* and *C* coexistence. The model also shows that *C* growth rate and
242 lag time on citrate do not change selection coefficients against *E*, once all resources are exhausted.

243 Indeed these parameters only influence how fast citrate is consumed, which is irrelevant to *E* that
244 does not consume citrate.

245 **Discussion**

246 **Interactions between co-evolving species evolve quickly, but can be maintained** 247 **through nonlinear NFDS patterns**

248 In this study, we investigated the long-term NFDS coexistence of two species. Following the
249 scenario developed in the introduction, we hypothesized that the two species may not coexist in
250 the long-term, especially if they present asymmetrical rates of adaptation (Fig 1c), as expected for
251 divergent species. Consistent with this scenario, one species (*C. freundii*) evolved at a considerably
252 faster rate than the other (*E. coli*), and became dominant in most of the coevolving cultures with
253 antibiotics (Fig. 2). The shift in the NFDS pattern is rapid so that the frequency of the two species
254 do not even have time to equilibrate at the frequency where their fitness is equal: by the time this
255 frequency is reached, the NFDS has further shifted down so that *E* is always more frequent than
256 would be expected by the current NFDS pattern. This is observed for both CRep20A and CRep20B
257 at intermediate and final times (Fig. 3, 5). The rate of change of NFDS patterns is here somewhat
258 faster (at the scale of 100s of generations) than those observed in the case of emerging
259 polymorphism within an asexual species (see e.g. the changes occurring at ~500 generation scale
260 in Rozen and Lenski 2000). However, in those cases where a polymorphism emerges within
261 asexual species, such an asymmetry quickly favoring one genotype is unexpected since the
262 coevolving genotypes are initially identical, save for a handful of mutations unlikely to drastically
263 favor one at the expense of the other. In our case, *E* and *C* are genetically largely divergent, and
264 did not have the same history of adaptation. *E* was initially relatively well adapted to the
265 environment, due to 10,000 generations of evolution in similar conditions in DM25, and then
266 ~500 generations in DM250-Nal (Harmand et al. 2018). Because adaptation most often shows a
267 pattern of diminishing returns (Lenski and Travisano 1994; Elena and Lenski 2003), *E* was most
268 probably not prone to important and rapid further adaptation to the abiotic conditions (including
269 to the Nal antibiotic, Harmand et al. 2018). In contrast, *C* had probably not been previously
270 exposed to serial batch culture in minimal medium (although the history of this strain cannot be
271 established), and hence was prone to faster and larger adaptive changes in those new conditions.
272 Such asymmetry in adaptive responses are likely to be pervasive among interacting species in
273 many natural situations, for example among populations that were temporarily isolated or
274 brought into contact secondarily or during invasion events. In those cases, long-term species co-
275 existence can be maintained through NFDS interactions only if those interactions are resilient to

276 further adaptation of these species, meaning that observation of NFDS interactions at one given
277 evolutionary time does not guarantee long-term coexistence. In our case, and despite a strong
278 asymmetry in rates of adaptation of the two interacting species, they were still coexisting after
279 870 generations (in the 39 replicates where *C* initially established). This striking outcome was due
280 to the particular S-shape of the NFDS patterns with extensive curvature near the fixation points
281 (Fig. 3) protecting *E* from extinction by strongly favoring it when rare. Such “S-shaped” NFDS
282 pattern can considerably extend long-term coexistence of interacting species, as an internal
283 equilibrium can still be preserved even with large shifts in intercepts. This is an important
284 departure from the simplified scenario usually envisioned and illustrated in Fig. 1, where linear
285 NFDS patterns do not strongly protect competing types against extinction if one is evolving
286 quicker. Hence, the shape of initial NFDS patterns between interacting types can be qualitatively
287 different among divergent species than among emerging polymorphisms, with a strong
288 consequence on their long-term coexistence.

289 **The evolution of NFDS patterns through time**

290 Asymmetric rates of adaptation to the abiotic conditions between *E* and *C* are clearly not a
291 sufficient explanation for all the results. For CRep20A and B, the NFDS patterns were mainly
292 shifted downwards between the initial and final time points. This pattern would be expected if
293 unconditionally beneficial mutations accumulated in *C*. However, NFDS patterns at an
294 intermediate time (generation 214, Fig 5) showed that, in both cases, the shape of the NFDS
295 changed sequentially, first shifted downward for the range of frequencies where *C* is rare (i.e. on
296 the right of *x*-axis in Fig 5), and then, later, shifted downward for the range of frequencies where
297 *C* is frequent (i.e. on the left of *x*-axis in Fig. 5). This degree of malleability of NFDS patterns is
298 surprising, but not unexpected. Initially, *C* is rare. It will therefore mostly fix mutations that confer
299 an advantage at this frequency (irrespective of their effect when *C* is frequent). Then the
300 equilibrium frequency of *C*, dictated by the NFDS pattern, increases, which triggers the fixation of
301 other mutations that confer an advantage at higher *C* frequency, as is observed. Consequently, the
302 final NFDS pattern appears as shifted down at all frequencies, giving the impression that it was
303 modified by the accumulation of beneficial but frequency-independent mutations. On the
304 contrary, the evolution of the NFDS pattern appears to be highly dynamic. It indicates that many
305 mutations are available but that some confer different selective advantages at different
306 frequencies, providing ample flexibility in the possible evolutionary deformation in the NFDS
307 pattern. Because these mutations influence the shape of the NFDS pattern, and have a frequency-
308 dependent effect, they probably modify the interaction between the two species, rather than being
309 unconditionally beneficial to the laboratory condition. This is a second important departure from
310 the simplified scenario usually envisioned (presented in the introduction and in Fig. 1). Many

311 mutations with a frequency-dependent effect seem to be available for adaptation, which allows
312 for considerable malleability in the evolution of the shape of NFDS pattern through time, beyond
313 mere changes in intercepts or slopes.

314 **The origin of non-linear NFDS patterns**

315 Different biological mechanisms can lead to NFDS (Table S1), and they can be difficult to tease
316 apart as they can lead to identical NFDS patterns. In our case study, many possibilities can be ruled
317 out (e.g. effect of parasites or predators) and some possibilities are worth discussing. First, NFDS
318 can emerge from environmental heterogeneity (mainly represented here by different resources)
319 provided that relevant fitness trade-offs exist among the different niches, as in Levene's model
320 (Levene 1953; Ravigné et al. 2004). This is a good candidate mechanism in this study since the
321 medium includes two carbon sources, glucose and citrate, on which *E* and *C* are known to be
322 specialized, respectively. In particular, *E* does not usually consume citrate under aerobic
323 conditions (Dimroth 2013), but *C* does. Second, additional niches can be created by the strains
324 themselves, as e.g. with cross-feeding interactions (Rosenzweig et al. 1994; Treves et al. 1998;
325 Doebeli 2002; Plucain et al. 2014) or detoxification of the environment (Dugatkin et al. 2005;
326 Kelsic et al. 2015). Third, the coexistence can rely on different strategies of exploitation of the
327 same resource via a trade-off between the uptake efficiency and the energetic conversion of a
328 resource. Finally, these mechanisms can be combined. For example, a strain which consumes the
329 resource rapidly but with low efficiency (third mechanism) may be prone to excrete byproducts,
330 which provides an opportunity for cross-feeding interactions (second mechanism).

331 The NFDS patterns alone do not provide sufficient information to decipher among these
332 mechanisms. Nevertheless, by modelling competition between one diauxic and one non-diauxic
333 species (Fig. S1), we pinpoint likely hypotheses at the basis of the establishment and evolution of
334 the interaction between the two species. First, we showed that an overall "S-shape" of the NFDS
335 patterns can be obtained in a two-resource model only if the diauxic species switches to the
336 second resource before the first one is entirely depleted. The remaining glucose constitutes a
337 private niche to the non-diauxic species that becomes strongly favored when rare. Such a private
338 niche cannot be revealed by growth curves of *E* and *C* in isolation (Fig 4). These curves showed
339 that *C* initially exhibits a clear diauxic growth on two resources and that only *C* evolved, notably
340 by decreasing the lag on citrate (λ_{C2} smaller). This may erroneously suggest that the evolution of
341 the NFDS pattern mainly rely on this smaller λ_{C2} . However, as shown by our model, such change
342 is neutral with respect to competition with *E*. It probably only evolved due to within-*C* competition
343 for citrate consumption. Within-*C* competition may also be key for the evolution of *C* anticipated
344 switch since it may be worthwhile for a diauxic species to anticipate the switch to a very abundant

345 second resource before the first one is entirely depleted. Ultimately, the model showed that it is
346 the time at which *C* initiates its switch to citrate rather than the time it takes to switch that is
347 central for the evolution of NFDS pattern between *E* and *C*. Our modelling clarifies these points.
348 Yet, it is likely that NFDS interactions can be generally more complex than the baseline situation
349 depicted in the model. For instance, in our case study, it is very possible that other intermediate
350 carbon sources (e.g. acetate, succinate) are temporarily excreted and exploited by *E* and *C*. The
351 model also provides keys to pointing the most promising candidate mechanisms that can yield
352 frequency-dependent changes that cannot be summarized by mere changes in intercept as
353 presented in Fig. 1. In our case, such mechanism creates a situation where adaptation in one
354 species indirectly opens a niche for another, in absence of cross-feeding. Further experiments are
355 however required to identify the exact mechanism at work, notably by investigating the
356 competition of the two species on glucose alone, provided that pH buffering in absence of citrate
357 can be controlled for.

358 **NFDS evolution across environments**

359 The evolution of *E* frequency in our experiment was strongly influenced by abiotic environmental
360 conditions (here the concentration of Nal). The patterns of variation are highly regular with
361 respect to the gradient of Nal concentration. The importance of environmental conditions in the
362 emergence or maintenance of biotic interactions has already been pointed out in other
363 experiments (e.g. in Hansen and Hubbell 1980; Healey et al. 2016). These observations suggest
364 that the environmental context can have a large influence on the long-term maintenance of NFDS
365 interactions. This would be easily interpretable if such environmental variation was related to the
366 mechanism of coexistence, for instance the proportion of the different available resources
367 (glucose or citrate). But this is not the case here: the environmental variable playing such a strong
368 role is the concentration of the antibiotic, which seems entirely unrelated to the mechanism of
369 coexistence and orthogonal to the issue of resource utilization. A possible explanation might be
370 that Nal represents an asymmetrical challenge for *C* and *E*. For instance, contrary to *E*, *C* may be
371 mostly unaffected by the presence of Nal at any concentration (e.g. because of reduced uptake or
372 a Nal-proof gyrase target). Some tests (not shown) indicated that *C* growth rate was not affected,
373 but *E* growth rate reduced with increasing Nal concentration up to 200 µg/mL. Another possibility
374 might be that, fortuitously, Nal resistance in *E* favored *C*. In particular, some (loss-of-function)
375 mutations on enzymes of the Krebs cycle, have been shown to be selected in *E* to resist Nal.
376 Presumably, metabolites that accumulate upstream of the blocks caused by the mutations
377 increase the expression levels of generalist efflux pumps, thereby removing Nal from the organism
378 (Helling and Kukora 1971; Lakshmi and Helling 1976; Helling et al. 2002). This resistance
379 mechanism, however, reduces metabolic efficiency and leads to the excretion of intermediate

380 metabolite that could be exploited by *C*. In all cases, adding a direct advantage to *C* with increasing
381 Nal concentration is sufficient to explain why *E* equilibrium frequency is lower at increasing Nal
382 doses (scenario modelled and shown in Fig. 6e). Irrespective of the exact underlying mechanism,
383 our results show that environmental conditions that are *a priori* unrelated to the mechanism of
384 coexistence can largely impact the evolution of NFDS patterns.

385 **The maintenance of rare species**

386 The persistence of rare species in communities is often difficult to understand, as they should be
387 very vulnerable to stochastic perturbations or to slight adaptation of their competitor. The
388 occurrence of a sharply increasing selective advantage at very low frequencies, as demonstrated
389 in our case, could explain these observations. This increased coexistence timespan may provide
390 sufficient time for further niche specialization, and eventually stabilization of interactions among
391 coevolving competitors. Here we showed that this non-linearity may occur because of incomplete
392 exploitation of resources by the more generalist species. There is indeed a strong selection
393 pressure within that generalist population (here *C* that consumes both glucose and citrate) to
394 switch resources when the first resource starts to be too rare to be worth exploiting. This
395 inevitably opens a private niche for a more specialized species that can persist in the long term at
396 very low frequency. This process of indirect niche construction may provide a general mechanism
397 to explain persistence of rare species in ecological systems.

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405 **References**

- 406 Allen, J. A. 1988. Frequency-dependent selection by predators. *Philos. Trans. R. Soc. Lond. B. Biol.*
407 *Sci.* 319:485–503.
- 408 Ayala, F. J., and C. A. Campbell. 1974. Frequency-dependent selection. *Annu. Rev. Ecol. Syst.* 115–
409 138.
- 410 Barnard, C. J., and R. M. Sibly. 1981. Producers and scroungers: A general model and its application
411 to captive flocks of house sparrows. *Anim. Behav.* 29:543–550.
- 412 Bell, G. 2008. *Selection: the mechanism of evolution*, second edition. Oxford University Press.
- 413 Billiard, S., M. Lopez-Villavicencio, B. Devier, M. Hood, C. Fairhead, and T. Giraud. 2011. Having sex,
414 yes, but with whom? Inferences from fungi on the evolution of anisogamy and mating types.
415 *Biol. Rev.* 86:421–442.
- 416 Blount, Z. D., J. E. Barrick, C. J. Davidson, and R. E. Lenski. 2012. Genomic analysis of a key
417 innovation in an experimental *Escherichia coli* population. *Nature* 489:513–8.
- 418 Blount, Z. D., C. Z. Borland, and R. E. Lenski. 2008. Historical contingency and the evolution of a
419 key innovation in an experimental population of *Escherichia coli*. *Proc. Natl. Acad. Sci. U. S. A.*
420 105:7899–906.
- 421 Bonsall, M. B. 2006. Longevity and ageing: appraising the evolutionary consequences of growing
422 old. *Philos. Trans. R. Soc. Lond. B. Biol. Sci.* 361:119–135.
- 423 Borghans, J. A. M., J. B. Beltman, and R. J. De Boer. 2004. MHC polymorphism under host-pathogen
424 coevolution. *Immunogenetics* 55:732–739.
- 425 Bürger, R. 2010. Evolution and polymorphism in the multilocus Levene model with no or weak
426 epistasis. *Theor. Popul. Biol.* 78:123–138.
- 427 Castric, V., and X. Vekemans. 2004. Plant self-incompatibility in natural populations: a critical
428 assessment of recent theoretical and empirical advances. *Mol. Ecol.* 13:2873–2889.
- 429 Clarke, B. 1962. Balanced polymorphism and the diversity of sympatric species. Pp. 47–70 *in* D.
430 Nichols, ed. *Taxonomy and Geography*. Systematics Association, Oxford.
- 431 Cordero, O. X., and M. F. Polz. 2014. Explaining microbial genomic diversity in light of evolutionary
432 ecology. *Nat Rev Microbiol* 12:263–273.
- 433 Débarre, F., and T. Lenormand. 2011. Distance-limited dispersal promotes coexistence at habitat
434 boundaries: reconsidering the competitive exclusion principle. *Ecol. Lett.* 14:260–6.
- 435 Devier, B., G. Aguilera, M. E. Hood, and T. Giraud. 2009. Ancient trans-specific polymorphism at
436 pheromone receptor genes in basidiomycetes. *Genetics* 181:209–223.
- 437 Dimroth, P. 2013. Molecular basis for bacterial growth on citrate or malonate. *EcoSalPlus* 1–36.
- 438 Doebeli, M. 2002. A model for the evolutionary dynamics of cross-feeding polymorphisms in
439 microorganisms. *Popul Ecol* 44:59–70.
- 440 Dugatkin, L. A., M. Perlin, J. S. Lucas, and R. Atlas. 2005. Group-beneficial traits, frequency-
441 dependent selection and genotypic diversity: an antibiotic resistance paradigm. *Proc. Biol.*
442 *Sci.* 272:79–83.
- 443 Elena, S. F., and R. E. Lenski. 2003. Evolution experiments with microorganisms: the dynamics and
444 genetic bases of adaptation. *Nat. Rev. Genet.* 4:457–69.
- 445 Felsenstein, J. 2017. *Theoretical evolutionary genetics*. J. Felsenstein, University of Washington,
446 Seattle.
- 447 Fincke, O. M. 2004. Polymorphic signals of harassed female odonates and the males that learn

- 448 them support a novel frequency-dependent model. *Anim. Behav.* 67:833–845.
- 449 Friesen, M. L., G. Saxer, M. Travisano, and M. Doebeli. 2004. Experimental evidence for sympatric
450 ecological diversification due to frequency-dependent competition in *Escherichia coli*.
451 *Evolution* 58:245.
- 452 Gallet, R., T. F. Cooper, S. F. Elena, and T. Lenormand. 2012. Measuring selection coefficients below
453 10^{-3} : Method, Questions, and Prospects. *Genetics* 190.
- 454 Gigord, L. D., M. R. Macnair, and A. Smithson. 2001. Negative frequency-dependent selection
455 maintains a dramatic flower color polymorphism in the rewardless orchid *Dactylorhiza*
456 *sambucina* (L.) Soo. *Proc. Natl. Acad. Sci. U. S. A.* 98:6253–6255.
- 457 Goldford, J. E., N. Lu, D. Bajić, S. Estrela, M. Tikhonov, A. Sanchez-Gorostiaga, D. Segrè, P. Mehta,
458 and A. Sanchez. 2018. Emergent simplicity in microbial community assembly. *Science*
459 361:469–474.
- 460 Good, B. H., M. J. McDonald, J. E. Barrick, R. E. Lenski, and M. M. Desai. 2017. The dynamics of
461 molecular evolution over 60,000 generations. *Nature* 551:45–50.
- 462 Gross, M. R. 1996. Alternative reproductive strategies and tactics: diversity within sexes. *Trends*
463 *Ecol. Evol.* 11:92–98.
- 464 Haldane, J. B. S., and S. D. Jayakar. 1963. Polymorphism due to selection of varying direction. *J.*
465 *Genet.* 58:237–242.
- 466 Hansen, S. R., and S. P. Hubbell. 1980. Single-nutrient microbial competition: qualitative
467 agreement between experimental and theoretically forecast outcomes. *Science* 207:1491–
468 1493.
- 469 Harmand, N., R. Gallet, R. Jabbour-Zahab, G. Martin, and T. Lenormand. 2017. Fisher’s geometrical
470 model and the mutational patterns of antibiotic resistance across dose gradients. *Evolution*
471 71:23–37.
- 472 Harmand, N., R. Gallet, G. Martin, and T. Lenormand. 2018. Evolution of bacteria specialization
473 along an antibiotic dose gradient. *Evol. Lett.* 2:221–232.
- 474 Healey, D., K. Axelrod, and J. Gore. 2016. Negative frequency-dependent interactions can underlie
475 phenotypic heterogeneity in a clonal microbial population. *Mol. Syst. Biol.* 12:877.
- 476 Hedrick, P. W. 1978. Genetic variation in a heterogeneous environment. V. Spatial heterogeneity
477 in finite populations. *Genetics* 89:389–401.
- 478 Heino, M., J. A. J. Metz, and V. Kaitala. 1997. Evolution of mixed maturation strategies in
479 semelparous life histories: the crucial role of dimensionality of feedback environment.
480 *Philos. Trans. R. Soc. B Biol. Sci.* 352:1647–1655.
- 481 Helling, R. B., B. K. Janes, H. Kimball, T. Tran, M. Bundesmann, P. Check, D. Phelan, and C. Miller.
482 2002. Toxic waste disposal in *Escherichia coli*. *J. Bacteriol.* 184:3699–703.
- 483 Helling, R. B., and J. S. Kukora. 1971. Nalidixic acid-resistant mutants of *Escherichia coli* deficient
484 in isocitrate dehydrogenase. *J. Bacteriol.* 105:1224–1226.
- 485 Helling, R., C. Vargas, and J. Adams. 1987. Evolution of *Escherichia coli* during growth in a constant
486 environment. *Genetics* 116:349–358.
- 487 Joron, M., and J. L. B. Mallet. 1998. Diversity in mimicry: Paradox or paradigm? *Trends Ecol. Evol.*
488 13:461–466.
- 489 Karasov, T. L., J. M. Kniskerb, L. Gao, B. DeYoung, J. Ding, U. Dubiella, R. O. Lastra, S. Nallu, F. Roux,
490 R. W. Innes, L. G. Barrett, R. R. Hudson, and J. Bergelson. 2014. The long-term maintenance of
491 a resistance polymorphism through diffuse interactions. *Nature* 512:436–440.
- 492 Kelsic, E. D., J. Zhao, K. Vetsigian, and R. Kishony. 2015. Counteraction of antibiotic production and

- 493 degradation stabilizes microbial communities. *Nature* 521:516–519.
- 494 Lakshmi, T. M., and R. B. Helling. 1976. Selection for citrate synthase deficiency in *icd* mutants of
495 *Escherichia coli*. *J. Bacteriol.* 127:76–83.
- 496 Le Gac, M., J. Plucain, T. Hindre, R. E. Lenski, and D. Schneider. 2012. Ecological and evolutionary
497 dynamics of coexisting lineages during a long-term experiment with *Escherichia coli*. *Proc.*
498 *Natl. Acad. Sci.* 109:9487–9492.
- 499 Lenormand, T. 2012. From local adaptation to speciation: specialization and reinforcement. *Int. J.*
500 *Ecol.* 2012:e508458.
- 501 Lenski, R. E., and M. Travisano. 1994. Dynamics of adaptation and diversification: a 10,000-
502 generation experiment with bacterial populations. *Proc Natl Acad Sci U S A* 91:6808–6814.
- 503 Levene, H. 1953. Genetic equilibrium when more than one ecological niche is available. *Am. Nat.*
504 87:331–333.
- 505 Levin, B. 1972. Coexistence of Two Asexual Strains on a Single Resource. *Science* 175:1272–1274.
- 506 Levin, B. R. 1988. Frequency-dependent selection in bacterial populations. *Philos. Trans. R. Soc.*
507 *Lond. B. Biol. Sci.* 319:459–472.
- 508 Lewontin, R. C. 1958. A General Method for Investigating the Equilibrium of Gene Frequency in a
509 Population. *Genetics* 43:419–34.
- 510 Maddamsetti, R., R. E. Lenski, and J. E. Barrick. 2015. Adaptation, clonal interference, and
511 frequency-dependent interactions in a long-term evolution experiment with *Escherichia coli*.
512 *Genetics* 200:619–631.
- 513 Manhart, M., B. V. Adkar, and E. I. Shakhnovich. 2018. Trade-offs between microbial growth phases
514 lead to frequency-dependent and non-transitive selection. *Proc. R. Soc. B Biol. Sci.*
515 285:20172459.
- 516 Mazancourt, C. de, and U. Dieckmann. 2004. Trade-Off Geometries and Frequency-Dependent
517 Selection. *Am. Nat.* 164:765–778.
- 518 O’Connell, L. M., and M. O. Johnston. 1998. Male and Female Pollination Success in a Deceptive
519 Orchid, a Selection Study. *Ecology* 79:1246–1260.
- 520 Oaten, A., and W. W. Murdoch. 1975. Functional response and stability in predator-prey systems.
521 *Am. Nat.* 109:289–298.
- 522 Olendorf, R., F. H. Rodd, D. Punzalan, A. E. Houde, C. Hurt, D. N. Reznick, and K. a Hughes. 2006.
523 Frequency-dependent survival in natural guppy populations. *Nature* 441:633–636.
- 524 Penn, D. J., and W. K. Potts. 1999. The evolution of mating preferences and major
525 histocompatibility complex genes. *Evolution* 153:145–164.
- 526 Plucain, J., T. Hindré, M. Le Gac, O. Tenaillon, S. Cruveiller, C. Médigue, N. Leiby, W. R. Harcombe, C.
527 J. Marx, R. E. Lenski, and D. Schneider. 2014. Epistasis and allele specificity in the emergence
528 of a stable polymorphism in *Escherichia coli*. *Science* 343:1366–1369.
- 529 Rainey, P. B., and M. Travisano. 1998. Adaptive radiation in a heterogeneous environment. *Nature*
530 394:69–72.
- 531 Ravigné, V., I. Olivieri, and U. Dieckmann. 2004. Implications of habitat choice for protected
532 polymorphisms. *Evol. Ecol. Res.* 6:125–145.
- 533 Reusch, T. B. H., M. A. Häberli, P. B. Aeschlimann, and M. Milinski. 2001. Female sticklebacks count
534 alleles in a strategy of sexual selection explaining MHC polymorphism. *Nature* 414:300–302.
- 535 Rosenzweig, R. F., R. R. Sharp, D. S. Treves, and J. Adams. 1994. Microbial evolution in a simple
536 unstructured environment: Genetic differentiation in *Escherichia coli*. *Genetics* 137:903–

537 917.

538 Rozen, D. E., and R. E. Lenski. 2000. Long-Term experimental evolution in *Escherichia coli*. VIII.
539 Dynamics of a balanced polymorphism. *Am. Nat.* 155:24–35.

540 Rozen, D. E., L. McGee, B. R. Levin, and K. P. Klugman. 2007. Fitness costs of fluoroquinolone
541 resistance in *Streptococcus pneumoniae*. *Antimicrob. Agents Chemother.* 51:412–416.

542 Rozen, D. E., N. Philippe, J. Arjan de Visser, R. E. Lenski, and D. Schneider. 2009. Death and
543 cannibalism in a seasonal environment facilitate bacterial coexistence. *Ecol. Lett.* 12:34–44.

544 Sinervo, B., and C. M. Lively. 1996. The rock-paper-scissors game and the evolution of alternative
545 male strategies. *Nature* 380:240–243.

546 Takahashi, Y., and M. Kawata. 2013. A comprehensive test for negative frequency-dependent
547 selection. *Popul. Ecol.* 55:499–509.

548 Takahata, N., and M. Nei. 1990. Allelic genealogy under overdominant and frequency-dependent
549 selection and polymorphism of major histocompatibility complex loci. *Genetics* 124:967–
550 978.

551 Těšický, M., and M. Vinkler. 2015. Trans-Species Polymorphism in Immune Genes: General Pattern
552 or MHC-Restricted Phenomenon? *J. Immunol. Res.* 2015:838035.

553 Thijs, H., J. R. Shann, and J. D. Weidenhamer. 1994. The effect of phytotoxins on competitive
554 outcome in a model system. *Ecology* 75:1959–1964.

555 Treves, D. S., S. Manning, and J. Adams. 1998. Repeated evolution of an acetate-crossfeeding
556 polymorphism in long-term populations of *Escherichia coli*. *Mol. Biol. Evol.* 15:789–97.

557 Turner, P. E., V. Souza, and R. E. Lenski. 1996. Tests of ecological mechanisms promoting the stable
558 coexistence of two bacterial genotypes. *Ecology* 77:2119–2129.

559 van Diepen, L. T. A., Å. Olson, K. Ihrmark, J. Stenlid, and T. Y. James. 2013. Extensive Trans-Specific
560 Polymorphism at the Mating Type Locus of the Root Decay Fungus *Heterobasidion*. *Mol. Biol.*
561 *Evol.* 30:2286–2301. Oxford University Press.

562 Wang, J., E. Atolia, B. Hua, Y. Savir, R. Escalante-Chong, and M. Springer. 2015. Natural variation in
563 preparation for nutrient depletion reveals a cost-benefit tradeoff. *PLoS Biol.* 13:1–31.

564 Weeks, A. R., and A. a Hoffmann. 2008. Frequency-dependent selection maintains clonal diversity
565 in an asexual organism. *Proc. Natl. Acad. Sci. U. S. A.* 105:17872–17877.

566

567 **Figure Legends**

568 **Figure 1.** Evolution of frequency-dependence and long-term coexistence. The figure illustrates
569 patterns of negative frequency-dependent selection between types (NFDS, left panels) and the
570 corresponding frequency variation in co-existing populations (right panels, the time scale is such
571 that mutations appear as instantaneously fixed). Basically, NFDS can be described as a negative
572 linear dependence between the selection coefficient of one type (y -axis) against its frequency (x -
573 axis) which defines the equilibrium frequency at which both types can eventually co-exist
574 indefinitely (i.e. where the NFDS line intersects the x -axis). However, this selective equilibrium
575 may be displaced by further mutations: Panels (a): In sexual species, a new allele at one
576 polymorphic locus is supposed to have a limited impact on the equilibrium frequency because
577 recombination will disconnect such allele at one locus from other mutations in the genome; Panels
578 (b): In de novo divergent asexual species where recombination is absent, both emerging types
579 share similar mutational opportunity to further adapt to their common environment at a whole
580 genome scale. Any beneficial mutation fixed in one lineage will shift the NFDS line either up or
581 down depending in which type such mutation arise (on fig. 1b, see e.g. beneficial mutations 2 and
582 3 arising in type A shifting up the NFDS line whereas beneficial mutations 1 and 4 arising in type
583 B shift it down). Overall, 2 types adapting at a similar rate will exhibit successive up and down
584 variations of the NFDS line, temporally changing their equilibrium frequency but not disrupting
585 their long-term coexistence; Panels (c): When NFDS **occurs** between types from 2 divergent
586 species with an asymmetric rate of adaptation to a common environment, the NFDS line will shift
587 up or down but to an extent causing the selective fixation of the type more prone to adapt to the
588 common environment (on fig. 1c see e.g. mutations 1, 3, 4 with large effect in B that are not
589 ultimately counterbalanced by mutation 2 with smaller effect in A). Panel (d): The previous panels
590 only consider the fixation of beneficial mutations uniformly affecting the fitness of one type
591 independently of its frequency. However, mutation contributing to the specialization of one type
592 to a dedicated ecological niche may alternatively affect the slope of the NFDS line instead of the
593 equilibrium frequency (see Fig. 1d). Such a mutation may occur in previous cases (a, b, c) with
594 increased slope improving long-term coexistence since mutation with a larger effect will be
595 required to switch the NFDS line to a point where one type will be selectively outcompeted.

596 **Figure 2.** Evolution of the proportion (y -axis in %) of *E. coli* against *C. freundii* throughout the
597 coevolution at different antibiotic concentrations (color code indicated in the figure for the
598 different Nal concentrations). In panel (a), mean values and standard deviations are calculated
599 per antibiotic dose (8 CRep per Nal dose, but 4 in Nal0). Values for each CRep are shown in panel
600 (b). Frequency estimates were obtained using samples of 100,000 cells (*E. coli* frequency was
601 computed as the proportion of YFP fluorescent cells).

602 **Figure 3:** Negative frequency-dependent selection patterns between *E. coli* and *C. freundii* at
603 initial (0) (grey dots) and final time (8) (black dots) of the coevolution from two sets of lines co-
604 evolved in DM250-Nal20 (CRep20A and B, top and bottom panels, respectively). Triangles
605 represent the crossed-time competitions. Each point is the selection coefficient of one
606 competition, some of which were done in duplicate (same *E* frequency). Green marks on the x-axis
607 represent the frequency of *E* measured in the mixes at initial (higher values) and final time (lower
608 values) of the coevolution.

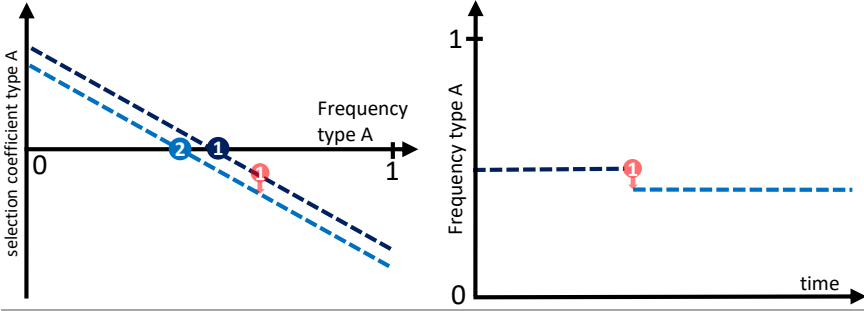
609 **Figure 4:** Growth curves in DM250-Nal20 of *E. coli* and *C. freundii* lines isolated from CRep20A
610 and CRep20B throughout the coevolution (grey gradient). Growth curves were repeated four
611 times, resulting in very similar curves, but for the clarity of the figure we selected one
612 representative set. The optical density (OD) was measured at regular intervals of 10 minutes and
613 each dot corresponds to one measure.

614 **Figure 5:** Transient deformation of the NFDS pattern at generation 200 (black squares) compared
615 to generation 0 (light grey dots) for CRep20A and CRep20B replicates. This deformation has
616 almost completely disappeared at generation 870 (not represented here for clarity, see Fig. 3).

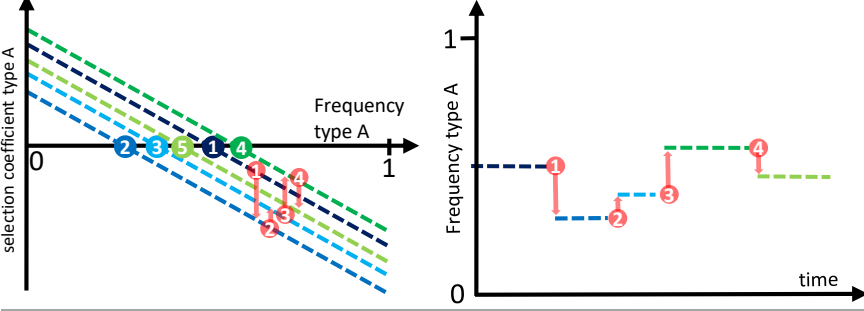
617 **Figure 6:** NFDS patterns obtained by modeling competition between *E* and *C* types. The x-axis
618 represents the frequency of *E* (from 0 to 1), and the y-axis the selection coefficient of *E* against its
619 competitor *C*. In (a), only one resource is considered and *C* initially has a growth advantage over
620 *E* (black line). The grey gradient represents a gradual increase of the efficiency of *E* in energetic
621 conversion that, if large enough, can result in NFDS with an internal frequency equilibrium. In (b)
622 and (c), two resources are considered. In the baseline situation (black lines) *E* and *C* are equally
623 able to exploit resource 1 (glucose), but *C* can switch to exploit resource 2 (citrate) after glucose
624 is depleted. In these conditions, *C* is favored at all frequencies. (b) If *E* evolves a shorter lag phase
625 (or equivalently higher growth rate) when exploiting resource 1, NFDS can emerge as indicated
626 by the grey lines crossing the x-axis (the larger the advantage of *E* on resource 1, the lighter the
627 line). (c) If *E* parameters remain constant but *C* evolves an anticipated switch to resource 2 (the
628 earlier the switch, the lighter the curve), a “S-shaped”-NFDS pattern emerges because *C* switches
629 to citrate before glucose is depleted, hence opening a ‘private’ niche for *E* on the remaining
630 glucose. This could equivalently be described as a three resources model (citrate, private glucose,
631 shared glucose). In (d) and (e), the model is used to identify parameters sustaining the observed
632 variations of NFDS patterns during the evolution of CRep. (d) Time dynamics in CRep20A (and
633 similarly CRep20B). At time T0, *E* and *C* coexist through NFDS with the S-shaped pattern signature
634 as described above. At time T2, the shift down of NDFS pattern mostly at high *E* frequency (step1
635 arrow, as in Fig. 5) can be obtained by increasing *C* conversion efficiency on citrate, but with an

636 earlier *C* switch to citrate (leaving more 'private' glucose to *E*). At time T8, the shift-down of NFDS
637 pattern mostly at low *E* frequency (step2 arrow, as in Fig. 3) can be obtained if *C* keeps its
638 conversion efficiency on citrate but evolves back to a later switch to citrate (i.e. leaving less
639 'private' glucose to *E* compared to T2). (e) Negative impact of Nal on *E* frequency at the
640 equilibrium. The shift-down of frequencies with increasing Nal concentrations (as in Fig. 2) can
641 be obtained if Nal imposes a greater reduction of growth rate on glucose (or greater increase of
642 lag phase) for *E* than for *C*. See sup. mat. for more details.

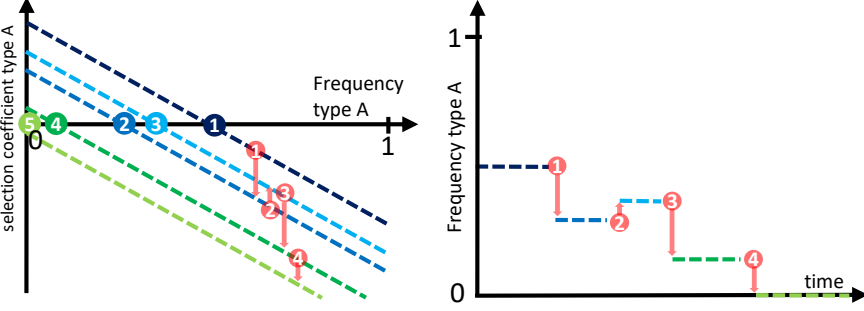
(a) Allele replacement sexual species



(b) Coevolving asexual lineages



(c) Coevolving divergent species



(d) Reinforcement of NFDS

