

Article

Effects of Relatedness on the Evolution of Cooperation in Nonlinear Public Goods Games

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Abstract: Evolution of cooperation has traditionally been studied by assuming that individuals adopt either of two pure strategies, to cooperate or defect. Recent work have considered continuous cooperative investments, turning full cooperation and full defection into two opposing ends of a spectrum and sometimes allowing for the emergence of the traditionally-studied pure strategies through evolutionary diversification. These studies have typically assumed a well-mixed population in which individuals are encountered with equal probability. Here, we allow for the possibility of assortative interactions by assuming that, with specified probabilities, an individual interacts with one or more other individuals of the same strategy. A closely related assumption has previously been made in evolutionary game theory and has been interpreted in terms of relatedness. We systematically study the effect of relatedness and find, among other conclusions, that the scope for evolutionary branching is reduced by either higher average degree of, or higher uncertainty in, relatedness with interaction partners. We also determine how different types of non-linear dependencies of benefits and costs constrain the types of evolutionary outcomes that can occur. While our results overall corroborate the conclusions of earlier studies, that higher relatedness promotes the evolution of cooperation, our investigation gives a comprehensive picture of how relatedness affects the evolution of cooperation with continuous investments.

Keywords: adaptive dynamics; evolution; cooperation

1. Introduction

Cooperation, and in particular the willingness to cooperate with relatives, is regularly observed in many species. Observed forms of cooperation include egg trading, cooperative foraging, and predator inspection among fishes; defensive coalitions, cooperative hunting, food sharing, and alarm calls among birds; grooming behaviour, alarm signals, coalitions, alloparenting, and cooperative hunting among various mammals; bloodsharing among vampire bats; care-giving behaviour in dolphins; foraging, anti-predator behaviour, and hive thermoregulation in honeybee colonies; and “social contracts” among paper wasps [1]. Social insects such as ants and bees also cooperate frequently (e.g., [2,3]) and cooperation is observed even among microbes (e.g. [4]). Finally, human societies are striking examples of cooperation.

Since cooperation often involves a cost to the individual without conveying a commensurate immediate benefit, one would naively suspect that cooperation cannot evolve to any higher degree. This, however, is not the case, and several mechanisms capable of promoting and supporting cooperation has been proposed and explored [5,6]. An influential explanation of cooperation between related individuals is kin selection, which states that there can be positive selection for a gene conveying cooperative behavior if the beneficiaries of that behavior are likely to share the same gene. This is typically the case when interactions occurs between relatives, but the principle applies more generally and assortative interaction between individuals can arise for a range of other reasons such as spatial

36 structure or social dynamics. In certain settings, the principle of kin selection can be formulated
 37 quantitatively in a form that is now known as Hamilton's rule: a cooperative gene is selected for if and
 38 only if $rb > c$, where r is the average degree of relatedness, b is the average benefit to the recipients of
 39 the cooperative act, and c is the average cost to the individual actor [7].

40 Cooperation has often been studied in the setting of evolutionary game theory based on the
 41 prisoner's dilemma or its many-player generalization, the public-goods game (see e.g. [8,9]). Although
 42 the latter game in principle allows for continuous investment, it is often assumed that individuals adopt
 43 either of two pure strategies corresponding to cooperation and defection. The evolutionary dynamics
 44 are studied with the so-called replicator equations in which strategies with a higher payoff than
 45 average in the population increase in frequency. To allow for the possibility of assortative interactions
 46 in this framework, Grafen [10] introduced the assumption that a fraction of an individual's interaction
 47 are with individuals of the same type with the remaining interactions occurring with partners drawn
 48 at random from the population. This approach and related approaches has since been used to address
 49 a range of evolutionary questions [11–17].

50 Recently, several studies have moved beyond the traditional cooperative games by considering
 51 continuous cooperative investments as well as non-linearity of benefits and costs [4,18–23]. With this
 52 setting, additional evolutionary outcomes become possible, including evolutionary branching in the
 53 cooperative trait under consideration and the emergence of two or more different coexisting strategies.
 54 In a landmark study, Doebeli et al. [18] considered a setting in which individuals interact in pairs and
 55 each make a cooperative investment for their common good. Their joint benefit is given by a non-linear
 56 function of their joint investment and their costs are given by a non-linear function of their respective
 57 investments. Finally, their payoffs are determined as the difference of the benefit and their respective
 58 costs. By assuming that the benefit and cost function were given by quadrature polynomials, they
 59 classified all possible evolutionary outcomes. Building on this framework, Cornforth et al. [4] showed
 60 how interaction assortment, interpreted as relatedness, can be incorporated. Though developed
 61 independently, the idea is similar to Grafen [10]. Cornforth et al. investigated how assortment
 62 affects the evolution of cooperation for three different benefit functions and under the assumption of
 63 proportional costs.

64 Here, we extend the previous results by Cornforth et al. and systematically study the effects
 65 of relatedness on the evolution of cooperation in non-linear public-goods games with continuous
 66 investment. In Section 2, we present the general model. We consider both quadratic cost and benefit
 67 functions and general, increasing, cost and benefit functions. In Section 3, we analyze the dynamics
 68 of quadratic cost and benefit functions, and then derive results for general cost and benefit functions.
 69 Finally, in Section 4, we recapitulate and discuss our main results and identify important challenges
 70 for future research.

71 2. Model description

72 2.1. Demographical dynamics

73 We consider a model based upon [18] and [4]. We assume an infinite population in which the i th
 74 individual has trait value x_i representing cooperative investment, with $0 \leq x_i \leq 1$. At each generation,
 75 N random individuals with respective trait values x_1, \dots, x_N are chosen for an interaction, in which
 76 the payoff to the i th individual, with $1 \leq i \leq N$, is given by

$$77 \quad \frac{B(x_1 + \dots + x_N)}{N} - C(x_i), \quad (1)$$

78 where $B : [0, N] \rightarrow \mathbb{R}^+ \cup \{0\}$ and $C : [0, 1] \rightarrow \mathbb{R}^+ \cup \{0\}$ are two increasing functions such that
 79 $B(0) = C(0) = 0$. $B(x_1 + \dots + x_N)$ is the collective benefit of the cooperative investment of the entire
 80 group, and $C(x_i)$ is the cost of the individual's own investment. The collective benefit is assumed to be
 81 divided equally among the members of the group, yielding a per capita benefit of $B(x_1 + \dots + x_N)/N$.

Each individual is assumed to be identical by descent (i.b.d.) to a random number of individuals in the group including itself. We define $\text{Pr}(k)$ to be the probability for an individual to be i.b.d. to exactly k individuals in the group including itself. Just like in Cornforth et al. [4], the expected payoff to a rare mutant with trait value m in a monomorphic population of residents with trait value r becomes

$$P(m, r) = \sum_{k=1}^N \frac{\text{Pr}(k)B(km + (N - k)r)}{N} - C(m). \quad (2)$$

We assume that the population dynamics follows the replicator equation [8], i.e., that successful strategies increase in frequency at a rate proportional to the difference with the average population-level payoff.

We first carry out a comprehensive analysis of quadratic benefit and cost functions $B(x) = b_2x^2 + b_1x$, $C(x) = c_2x^2 + c_1x$ that typically look like in Figure 1. We restrict the range of possibilities by requiring that $B(x_1 + \dots + x_N)$ and $C(x_i)$ should be increasing, i.e., that $b_1 > 0$, $c_1 > 0$, $b_1 \geq -2Nb_2$, and $c_1 \geq -2c_2$. The signs of b_2 and c_2 will depend on whether the functions are accelerating or decelerating. As a second step, we derive results for general increasing benefit and cost functions.

2.2. Evolutionary dynamics

We base our analysis of the evolutionary dynamics on adaptive-dynamics techniques [24–26]. Readers not familiar with these methods may want to refer to an introductory text such as Brännström et al. [27]. In brief, we consider the initial growth rate of a rare strategy m in an environment dominated by a resident strategy r . This growth rate is called the invasion fitness and written $S_r(m)$. The sign of the invasion fitness determines whether the rare strategy can grow in numbers and invade. From the invasion fitness we obtain the selection gradient $S'_r(r)$. Strategies at which selection ceases, $S'_r(r) = 0$ are called evolutionary singular. They can be evolutionary attractors, in which case they are called convergence stable. Once established, the subsequent evolutionary dynamics depends on whether the singular strategy is evolutionarily stable or an evolutionary branching point. In the former case, evolution comes to a halt while in the latter case, the population diversifies and two coexisting strategies emerges.

From the replicator equation (see Appendix A.1) we obtain the following expression for the invasion fitness

$$S_r(m) = P(m, r) - P(r, r) = \sum_{k=1}^N \frac{\text{Pr}(k)B(km + (N - k)r)}{N} - C(m) - \left[\sum_{k=1}^N \frac{\text{Pr}(k)B(Nr)}{N} - C(r) \right]. \quad (3)$$

Differentiating with respect to the mutant trait value and evaluating at $m = r$ yields the selection gradient

$$S'_r(r) = \sum_{k=1}^N \frac{k \text{Pr}(k)B'(Nr)}{N} - C'(r) = \mu B'(Nr) - C'(r), \quad (4)$$

where μ is the average whole-group relatedness, defined as the expected value of fraction of the group that is i.b.d. to the focal individual (including the focal individual itself), i.e.

$$\mu = \sum_{k=1}^N \frac{k \text{Pr}(k)}{N}. \quad (5)$$

Given this selection gradient, we obtain the condition for convergence stability of an evolutionarily singular strategy

$$NB''(Nr^*)\mu - C''(r^*) < 0, \quad (6)$$

and the condition for evolutionary stability

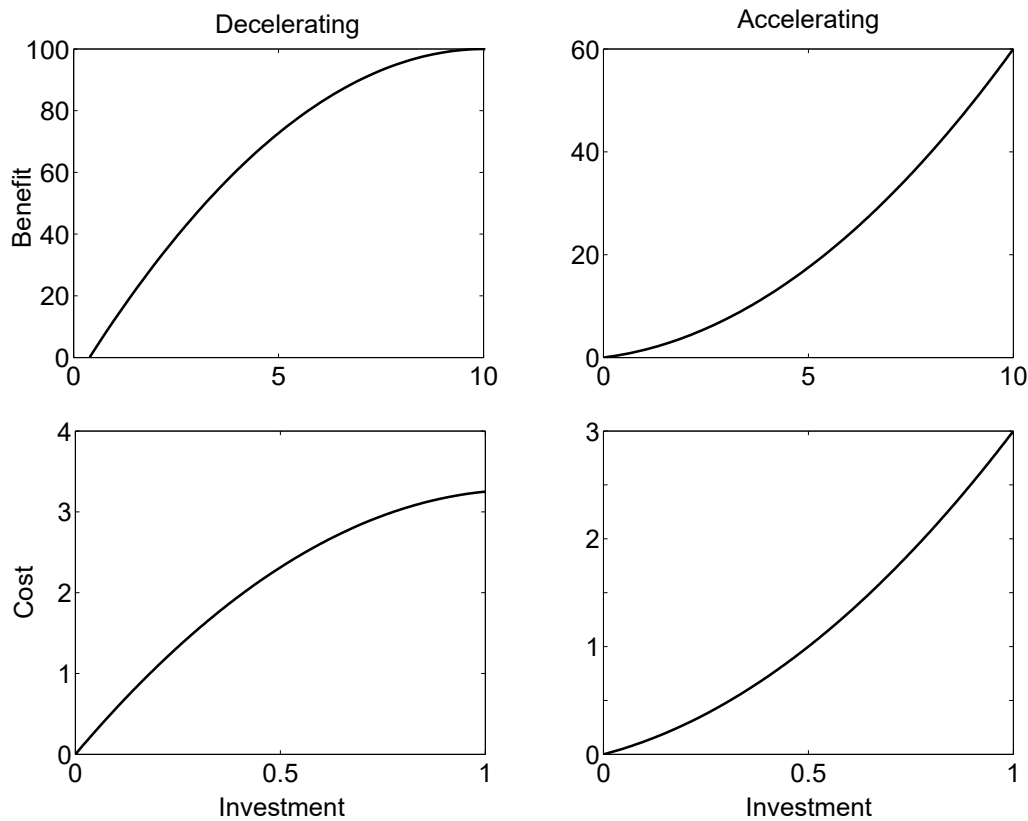


Figure 1. Examples of cost and benefit functions. Parameters used are for decelerating benefit $b_2 = -1$, $b_1 = 20$; for accelerating benefit $b_2 = 0.5$, $b_1 = 1$; for decelerating cost $c_2 = -2.75$, $c_1 = 6$; and for accelerating cost $c_2 = 2$, $c_1 = 1$. In choosing the horizontal range for the plots of the benefit functions, we have assumed $N = 10$.

$$NB''(Nr^*)[\mu^2 + \sigma^2] - C''(r^*) < 0, \quad (7)$$

where $\sigma^2 = \text{Var}[K/N]$, with K being the random variable with probability density function $\text{Pr}(k)$. For a derivation of these conditions, see the appendix of Cornforth et al. [4], as well as Appendix A.2.

3. Results

We first carry out a comprehensive analysis of the effect of assortment on the evolution of cooperation with quadratic benefit and cost functions and then derive results for general cost and benefit functions. Our key findings are illustrated by Figure 2 and 3. These show the effects of increased average relatedness and increased variance in relatedness on the direction of selection and evolutionary stability of the cooperative investment. Note, in particular, that a singular strategy increases with increased relatedness if and only if it is convergence stable, and that increased relatedness as well as how increased uncertainty in relatedness can change the evolutionary stability of singular strategies. Finally we corroborate selected results using numerical simulations.

3.1. Quadratic benefit and cost functions

Recall that we assume benefit and cost functions, B and C , that are increasing, satisfy $B(0) = C(0) = 0$, and are given by quadratic polynomials, as shown in Figure 1. For these functions, we completely classify the evolutionary dynamics and its dependence on the mean and variance of relatedness. Our conclusions are corroborated by numerical investigations (Fig. 4).

With quadratic polynomials, the selection gradient, Eq. 2.2, is given by

$$S'_r(r) = \mu B'(Nr) - C'(r) = \mu(2Nb_2r + b_1) - 2c_2r - c_1. \quad (8)$$

Recalling from Sect. 2.1 that that we must have $b_1 \geq -2Nb_2$ for B to be increasing, we have $2Nb_2r + b_1 > 0$ and it follows as expected that increased relatedness always alters selection pressures in the direction of higher cooperative investments.

From the selection gradient, Eq. 8, we solve for the singular strategies, i.e., the values of r at which the selection gradient vanishes, and find that for each μ , there is at most one interior evolutionarily singular strategy in the allowed range of cooperative investments. It is given by

$$r^*(\mu) = \frac{c_1 - b_1\mu}{2(Nb_2\mu - c_2)}, \quad (9)$$

whenever this value lies between 0 and 1.

Momentarily leaving aside the constraints on μ and r^* imposed by their biological interpretation, we note that the graph of $r^*(\mu)$ crosses zero at $\mu_0 = c_1/b_1$, has a vertical asymptote at $\mu_V = c_2/(Nb_2)$ and a horizontal asymptote at $r_H = -b_1/(2Nb_2)$. The graph consists of two curves that are located above and below the horizontal asymptote, respectively. Recalling from Sect. 2.1 that we must have $b_1 \geq -2Nb_2$ for B to be increasing, we see that the horizontal asymptote is located outside of the range $0 \leq r < 1$. Specifically, for accelerating benefits we have $r_H > 1$ and for decelerating benefits we have $r_H < -1$. It follows that only one of the curves can be biologically relevant and this is the curve that crosses zero at $\mu_0 = c_1/b_1$.

The evolutionary dynamics now depends on whether the horizontal asymptote is located above or below the allowed trait range and on whether the graph of r^* crosses zero before or after the vertical asymptote, i.e. on whether $\mu_H > \mu_0$. As we have already noted, the answer to the former question depends on whether benefits are accelerating or decelerating. We have not found a similar straightforward interpretation of the latter condition, but we note that $\mu_0 > \mu_V$ whenever benefits and costs are not simultaneously accelerating or simultaneously decelerating. For each of the four possible combinations, we can characterize the evolutionary dynamics.

Theorem 1. *Let μ_0 , μ_V , and r_H be as defined above. Let $r^*(\mu)$ be the interior strategy given by Eq. 3.1 whenever $0 < r^*(\mu) < 1$. Otherwise, let $r^*(\mu)$ be the boundary strategy that results from directional selection. The following conclusions then hold.*

- Assume B is accelerating and that $\mu_0 > \mu_V$, as is the case if C is decelerating. For $\mu < \mu_V$, we have $r^*(\mu) = 0$ and selection is thus towards no cooperation. For $\mu_V < \mu < \mu_0$, $r^*(\mu)$ is decreasing and the evolutionary dynamics is bistable. For $\mu > \mu_0$ we have $r^*(\mu) = 1$ and selection is thus towards full cooperation.
- Assume B is accelerating and that $\mu_0 < \mu_V$. For $\mu < \mu_0$ we have $r^*(\mu) = 0$ and selection is thus towards no cooperation. For $\mu_0 < \mu < \mu_V$ we have $r^*(\mu)$ is increasing and convergence stable, i.e., an evolutionary attractor. For $\mu > \mu_V$ we have $r^*(\mu) = 1$ and selection is thus towards full cooperation.
- Assume B is decelerating and $\mu_0 > \mu_V$, as is the case if C is accelerating. For $\mu < \mu_0$ we have $r^*(\mu) = 0$ and selection is thus towards no cooperation. For $\mu > \mu_V$, $r^*(\mu)$ is increasing and convergence stable, i.e., an evolutionary attractor.
- Assume B is decelerating and $\mu_0 < \mu_V$. For $\mu_0 < \mu < \mu_V$, $r^*(\mu)$ is increasing and convergence stable, i.e., an evolutionary attractor. For $\mu > \mu_V$ we have $r^*(\mu) = 1$ and selection is thus towards full cooperation.

Proof. The conclusions follow from the geometric observations that precede the theorem and the fact, also discussed above, that the selection gradient is increasing function of μ . \square

We next investigate the effects of relatedness on evolutionary stability. The condition for evolutionary stability, Eq. 7, is

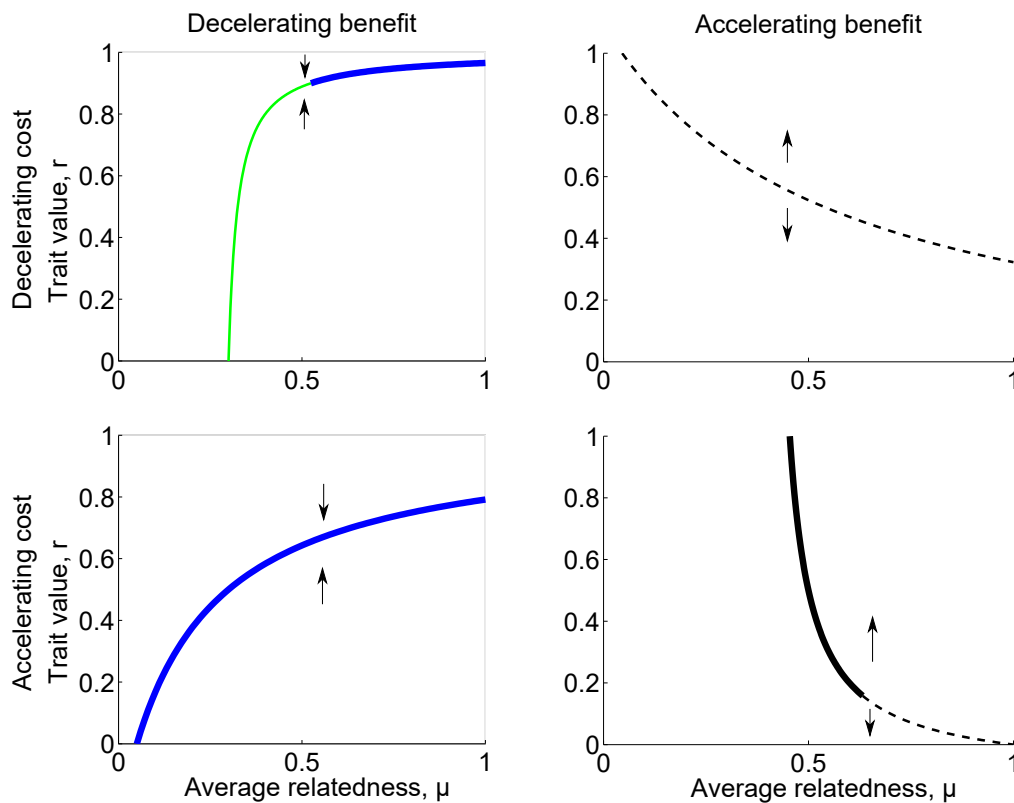


Figure 2. Bifurcation diagrams showing how the evolutionary dynamics depend on average relatedness for all four possible combinations of the accelerating and decelerating benefit and cost functions in Fig. 1. Green thin line represents convergence stability without evolutionary stability, blue thick line represents convergence stability with evolutionary stability, black dotted line represents neither convergence stability nor evolutionary stability, and black thick line represents evolutionary stability without convergence stability. We see that the four cases are qualitatively different. We also see that when B and C are both decelerating, a certain range of relatedness leads to evolutionary branching. The parameters used were in all cases $N = 10$, and b_2, b_1, c_2, c_1 as in Figure 1.

$$182 \quad Nb_2(\mu^2 + \sigma^2) - c_2 < 0. \quad (10)$$

183 We thus see that both increased average relatedness (μ) and increased variance in relatedness (σ^2)
 184 can affect the evolutionary stability of a singular strategy. We consider four cases depending on the
 185 signs of b_2 and c_2 . When benefits are accelerating ($b_2 > 0$) and costs decelerating ($c_2 < 0$), the singular
 186 strategy is never evolutionarily stable. As Theorem 1 shows, any interior singular strategy will lack
 187 convergence stability and there will thus not be any evolutionary branching points. Diversification
 188 may still be possible, but would have to be established through a large change in strategy. When
 189 benefits are decelerating ($b_2 < 0$) and costs accelerating ($c_2 > 0$), any interior singular strategy is
 190 always evolutionarily stable. If benefits and costs are both accelerating ($b_2 > 0, c_2 > 0$), either is
 191 possible and any change with increased relatedness μ or variance σ^2 is a loss of evolutionary stability.
 192 Finally, if benefits and costs are both decelerating ($b_2 < 0, c_2 < 0$), either is possible and any change
 193 with increased relatedness μ or variance σ^2 is a gain of evolutionary stability.

194 Figure 2 shows typical outcomes for four combinations of accelerating and decelerating costs and
 195 benefits. The outcomes are typical and consistent with our conclusions, though it should be noted that
 196 Theorem 1 allows for alternative outcomes of directional selection when benefits and costs are both
 197 accelerating or both decelerating.

198 Finally, we investigate the potential for evolutionary branching when benefits and costs are both
 199 accelerating or both decelerating. Solving Eq. 6 for μ , the boundary between convergence stability and
 200 not convergence stability becomes

$$201 \quad \mu_c = \frac{c_2}{Nb_2}, \quad (11)$$

202 assuming $b_2 \neq 0$. If B is decelerating, then $\mu > \mu_c$ implies convergence stability, and if B is accelerating,
 203 then $\mu < \mu_c$ implies convergence stability.

204 Solving Eq. 10 for σ^2 , we get the boundary between evolutionary stability and instability

$$205 \quad \sigma_e^2 = \frac{c_2 - Nb_2\mu^2}{Nb_2}, \quad (12)$$

206 once again assuming $b_2 \neq 0$. If B is decelerating, then $\sigma^2 > \sigma_e^2$ implies evolutionary stability, and if B
 207 is accelerating, then $\sigma^2 < \sigma_e^2$ implies evolutionary stability. Instead solving Eq. 10 for μ we see that the
 208 singular strategy gains evolutionary stability at

$$209 \quad \mu_e = \sqrt{\frac{c_2 - Nb_2\sigma_e^2}{Nb_2}}, \quad (13)$$

210 If B is decelerating, then $\mu > \mu_e$ implies evolutionary stability, and if B is accelerating, then $\mu < \mu_e$
 211 implies evolutionary stability. For μ_e to equal μ_c implies $\sigma_e^2 = \mu_c(1 - \mu_c)$, that is, with maximum
 212 variance (see A.3), convergence stability and evolutionary stability switch at the same time. Assuming
 213 on the other hand that $\mu_e < \mu_c$, this would imply $\sigma_e^2 > \mu_e(1 - \mu_e)$ which is not possible, see Appendix
 214 A.3. In other words, evolutionary stability will switch as soon as or after convergence stability switches
 215 when increasing the average relatedness, depending on whether the variance is at its maximum or not.

216 There is a region in the $b_2 - c_2$ -plane where μ_e is undefined. This is when $b_2 < 0$, $c_2 > Nb_2\sigma^2$; or
 217 $b_2 > 0$, $c_2 < Nb_2\sigma^2$. When the first one of these is the case, the condition for evolutionary stability will
 218 always hold, and hence the strategy will always be evolutionarily stable. Conversely, when the second
 219 is the case, the strategy will never be evolutionarily stable.

220 Using (12), we can plot σ^2 against μ to find out the effects of increased variance in relatedness
 221 on the evolutionary stability, as in Figure 3. We see that increased variance has a noticeable effect:
 222 The higher the variance, the sooner the switch to evolutionary stability. We also see that increased
 223 relatedness can make the evolutionary dynamics bistable.

224 We have carried out numerical simulations to corroborate our findings results, using the method
 225 described in Appendix B. Our analytical investigation predicts evolutionary branching at $\mu = 0.4$ and
 226 evolutionary stability at $\mu = 0.8$ for B and C decelerating, and bistability when $\mu = 0.5$ for B and C
 227 accelerating. This is consistent with the results from the simulations, see Figure 4.

228 3.2. General cost and benefit functions

229 Having analyzed the case of quadratic cost and benefit functions, we now turn to the case of
 230 general strictly increasing non-linear functions. As our first result, we note that the position of an
 231 interior singular strategy, if it exists, depends only on the average relatedness and not on any higher
 232 moment of the probability distribution. To see this, we recall that an interior singular strategy r^* is by
 233 definition a point at which the selection gradient vanishes,

$$234 \quad \mu B'(Nr^*) - C'(r^*) = 0.$$

235 To see how increased relatedness affects an interior singular strategy, we differentiate implicitly with
 236 respect to μ which gives,

$$237 \quad r^{*'}(\mu) = \frac{B'(Nr^*)}{C''(r^*) - \mu NB''(Nr^*)}. \quad (14)$$

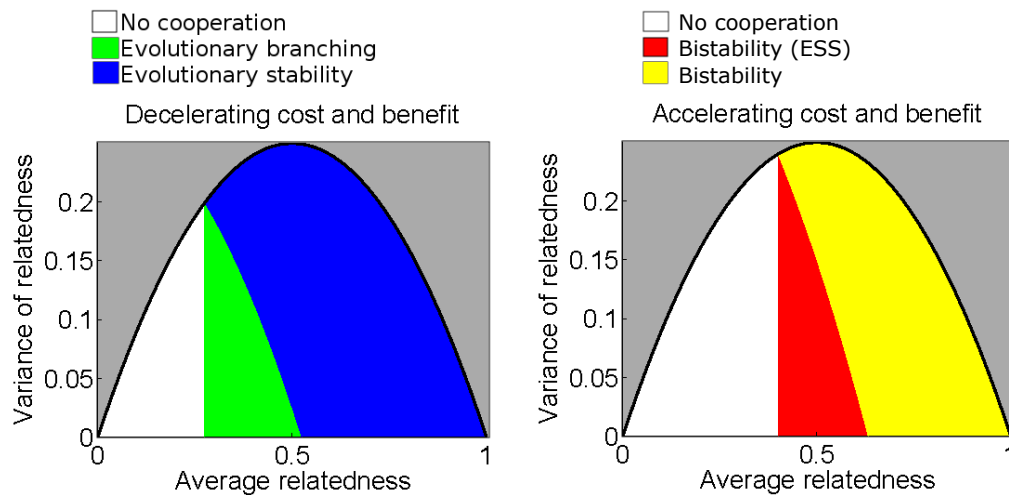


Figure 3. Plots of variance in relatedness against average relatedness. The grey region represents combinations of mean and variance that are not logically possible (see Appendix A.3). We see that for decreasing cost and benefit, the region with evolutionary branching shrinks with increased variance in relatedness. We also see that for increasing cost and benefit, the region with neither convergence stability nor evolutionary stability shrinks with increased variance in relatedness. In the latter case, with increased average relatedness or variance in relatedness, the singular strategy moves from evolutionary stability without convergence stability to neither evolutionary stability nor convergence stability. The parameters used are in both cases $N = 10$, and b_2, b_1, c_2, c_1 as in Figure 1

238 From Eq. 6 we see that the denominator is positive if and only if the interior singular strategy is
 239 convergence stable. Since the benefit function is strictly increasing, we conclude that increased
 240 relatedness has opposite effect on interior singular strategies depending on whether they are
 241 evolutionary attractors or repellers. As illustrated in Fig. 2, evolutionary attractors increase with
 242 relatedness while evolutionary repellers decrease with relatedness.

243 Next, we restrict attention to benefit and cost functions that are either accelerating or decelerating
 244 for the full range of strategies considered, giving a total of four combinations to be explored. For two
 245 cases, we give a full classification of the evolutionary dynamics, allowing for the possibility of large
 246 mutational steps.

247 **Theorem 2.** Assume that benefits are decelerating and costs are accelerating, more specifically that $B''(x) < 0$
 248 and $C''(x) > 0$ for all $x \in [0, 1]$. Then there is exactly one continuously stable strategy r^* . Furthermore, we
 249 have that $S_r(m) < 0$ if $m < r \leq r^*$ or $m > r \geq r^*$. The continuously stable strategy is always evolutionarily
 250 stable and increases with the degree of relatedness.

251 In short, the theorem asserts that the evolutionary dynamics will lead to a monomorphic
 252 population that, once established, cannot be invaded by any other strategy.

253 **Proof.** We have already shown in the analysis of quadratic benefit and cost functions that all three
 254 outcomes can in fact occur, hence we only need to assert that no additional outcomes are possible. To
 255 first see that there can never be more than one interior singular strategy, we show that the selection
 256 gradient is a strictly declining function of the resident strategy. Differentiating the selection gradient
 257 with respect to the resident trait value gives,

$$258 \quad \frac{d}{dr} S'_r(r) = \mu N B''(Nr) - C''(r) < 0 \text{ for all } r \in [0, 1].$$

259 Thus, we see that there can be maximally one interior singular strategy which can be either a boundary
 260 strategy, $r^* = 0$ (no investment), $r^* = 1$ (maximal investment) or an interior strategy, $0 < r^* < 1$
 261 (intermediate investment). As the selection gradient is a declining function of the resident strategy,
 262 we also conclude that directional selection will small evolutionary steps will eventually lead to the
 263 vicinity of this strategy, i.e., the singular strategy is convergence stable. To see that large mutational
 264 steps will also leads towards the strategy and that, once established, it cannot be invaded by any other
 265 strategy, we write the invasion fitness as

$$266 \quad S_r(m) = \int_r^m S'_r(x) dx = \int_r^m \left(S'_r(r) + \int_r^x S''_r(y) dy \right) dx = (r - m)S'_r(r) + \int_r^m \int_r^x S''_r(y) dy dx. \quad (15)$$

267 It is easily seen that $S''_r(m) < 0$ and hence the double integral in the second term is always negative
 268 since x is intermediate between r and m . It follows that $S_r(m) < 0$ whenever $(r - m)S'_r(r) < 0$, which
 269 is precisely when $m < r \leq r^*$ or $m > r \geq r^*$. Thus, the strategy is continuously stable as asserted. \square

270 The other case which can similarly be classified is that of accelerating costs and decelerating
 271 benefits. In this case, coexistence of cooperators and defectors is possible whenever the evolutionary
 272 dynamics is bistable.

273 **Theorem 3.** *Assume that benefits are accelerating and costs are decelerating, more specifically that $B''(x) < 0$
 274 and $C''(x) > 0$ for all $x \in [0, 1]$. Then selection is either towards full cooperation, towards no cooperation,
 275 or there is exactly one interior singular stable strategy r^* in the sense that $S_r(m) > 0$ if $m < r \leq r^*$ or
 276 $m > r \geq r^*$. In the case of an interior singular strategy, we have that $S_0(1) > 0$ and $S_1(0) > 0$, enabling a
 277 protected dimorphism or cooperators and defectors. The interior singular strategy is never evolutionarily stable
 278 and decreases with the degree of relatedness.*

279 **Proof.** The first part of the proof is nearly identical to the previous case, but with opposite signs.
 280 Specifically, we show by differentiation that the selection gradient is increasing which implies that
 281 there can be at most one interior singular strategy, r^* . Noting that $S''_r(m) > 0$, we conclude from Eq. 15
 282 with $r = r^*$ that $S_r(m) > 0$ if $m < r \leq r^*$ or $m > r \geq r^*$.

283 It remains to show that a protected dimorphism of cooperators and defectors is possible whenever
 284 we have an interior singular strategy, $0 < r^* < 1$. Note that in this case, $S'_1(1) > 0$ and $S'_0(0) < 0$ due
 285 to the bistable evolutionary dynamics. Furthermore noting that the double integral in Eq. 15 is positive
 286 independent of whether $m > r$ or $r < m$, we have that

$$287 \quad S_1(0) = (1 - 0)S'_1(1) + \int_0^1 \int_x^1 S''_r(y) dy dx > 0, \quad (16)$$

288 and

$$289 \quad S_0(1) = (0 - 1)S'_0(0) + \int_0^1 \int_0^x S''_r(y) dy dx > 0, \quad (17)$$

290 showing that a protected dimorphism of cooperators and defectors is possible. \square

291 4. Discussion

292 In this paper, we have carried out a systematic investigation of the effects of relatedness on the
 293 evolution of cooperation in non-linear public goods game with continuous investments. As expected,
 294 we found that relatedness is beneficial for cooperation. When the evolutionary outcome is a single
 295 intermediate level of cooperation, higher relatedness increases that level. If, on the other hand, the
 296 evolutionary outcome is bistable such that an initially uncooperative population evolves to lower
 297 levels of cooperation and an initially cooperative society evolves to higher levels of cooperation, higher
 298 relatedness decreases the threshold level of cooperation that separates the two outcomes. The degree
 299 of relatedness also affects the potential for evolutionary branching, i.e., the emergence of two or more

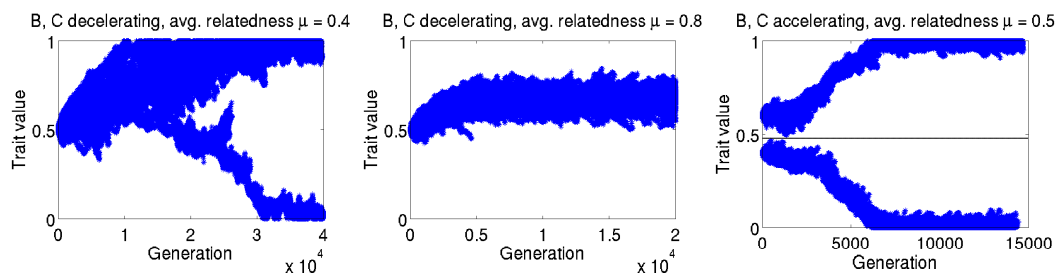


Figure 4. Simulations using group size $N = 10$, and B and C both decelerating or both accelerating. As predicted by the analytical investigation, when B and C are decelerating, the population undergoes evolutionary branching when $\mu = 0.4$, and it reaches evolutionary stability when $\mu = 0.8$. When B and C are accelerating, bistability occurs when $\mu = 0.5$. In all three cases, $\sigma^2 = 0$.

300 coexisting strategies. We find that relatedness reduces the scope for evolutionary branching, making it
 301 more likely that all individual evolve the same strategy.

302 Our framework allow us to also investigate the consequences of uncertainty in the degree to which
 303 an individual is related with his or her interaction partners, and we find that such uncertainty also
 304 reduces the scope for evolutionary branching. This can be compared with an earlier result showing that
 305 uncertainty in the size of interaction groups reduces the scope for evolutionary branching when the
 306 payoff structure has an additive form but not when it has a multiplicative form [20]. In this study, we
 307 have considered only payoff functions with an additive form and by analogy it appears plausible that
 308 uncertainty in relatedness might increase the scope for evolutionary branching under multiplicative
 309 payoff structures.

310 In a related study, Molina and Earn [23] rigorously analyzed a similar public goods game with
 311 non-linear benefits and linear costs. While they did not consider relatedness or assortment per se,
 312 one of their results, Theorem 4.4 in which they give conditions for the existence of a continuously
 313 stable strategy, allow for finite proportions of mutants and can be recast as a result on assortment.
 314 Our results differ by asserting global convergence stability and global evolutionary stability as well
 315 as in allowing for non-linear cost functions. The latter is important, as one cannot reduce to the case
 316 of linear costs by measuring investments in units of fitness costs, i.e., by redefining trait values as
 317 $\tilde{x} := C(x)$, with C being the cost function. Although this change of unit would make the cost function
 318 linear, the benefit function B would no longer depend on the sum of individual investments. Hence,
 319 the results by Molina and Earn [23] do not extend to public goods games with non-linear costs.

320 There are several directions in which the work presented here could be extended and we
 321 particularly wish to highlight two. First, our assumptions that the public goods are formed by
 322 adding individual contributions and that the payoff can be expressed as a difference of benefit and
 323 cost function are rather limiting and are unlikely to adequately describe all relevant settings involving
 324 public goods. Recently, Ito et al. [21] introduced and considered a large class of possible interaction
 325 structures in cooperative games. We think it would be interesting to extend our work to this larger
 326 framework and, in particular, systematically explore the effects of relatedness under the different
 327 options for aggregating rewards. Second, we base our results on a simplified representation of
 328 assortment. While simple representations have advantages, we think it would be good to complement
 329 our approach with more mechanistically-grounded studies of how assortment arises and influences
 330 the evolution of cooperation.

331 Our conclusion that increased assortment facilitates the evolution of cooperation is expected,
 332 in-line with empirical studies (e.g., [28]), and only a few exceptions to this rule has been reported in
 333 the literature (see for example [29] which shows how increased competition may prevent selection
 334 for altruistic genes). The value of our study thus do not lie as much in its overall conclusion as in
 335 revealing in detail how increased assortment helps to promote cooperation, for example by showing
 336 how boundaries in bistable evolutionary regimes shift and revealing that the scope for evolutionary

337 diversification is reduced by both increased assortment and increased uncertainty in assortment. As
 338 such, our findings should prove valuable both to theoreticians and empiricists striving to interpret
 339 results on the evolution of cooperation from laboratory experiments and field observations.

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 343 name. Å.B. conceived and designed the study; K.C.G. performed the numerical investigations. Å.B. and K.C.G.
 344 jointly analysed the case of quadratic cost and benefit functions. Å.B. proved the results on general benefit and
 345 cost functions; Å.B. and K.C.G. jointly discussed the results and wrote the paper.

346 Appendix A. Analytical investigations

347 Appendix A.1. Derivation of selection gradient

348 We start with

$$349 \quad S_r(m) = \lim_{x_m \rightarrow 0^+} \left[\frac{dx_m/dt}{x_m} \right], \quad (\text{A1})$$

350 and from the definitions of dx_m/dt , P_m , and \bar{P} , we get

$$\begin{aligned} 351 \quad S_r(m) &= \lim_{x_m \rightarrow 0^+} \left[\frac{dx_m/dt}{x_m} \right] \\ 352 &= \lim_{x_m \rightarrow 0^+} \left[\frac{x_m \cdot (P_m - \bar{P})}{x_m} \right] \\ 353 &= \lim_{x_m \rightarrow 0^+} [P_m - \bar{P}] \\ 354 &= \lim_{x_m \rightarrow 0^+, x_r \rightarrow 1^-} [P_m - (x_r \cdot P_r + x_m \cdot P_m)] \\ 355 &= \lim_{x_m \rightarrow 0^+, x_r \rightarrow 1^-} [x_r \cdot P(m, r) + x_m \cdot P(m, m) - \\ 356 &\quad (x_r \cdot (x_r \cdot P(r, r) + x_m \cdot P(r, m)) + x_m \cdot (x_r \cdot P(m, r) + x_m \cdot P(m, m)))] \\ 357 &= 1 \cdot P(m, r) + 0 \cdot P(m, m) - (1 \cdot (1 \cdot P(r, r) + 0 \cdot P(r, m)) + 0 \cdot (1 \cdot P(m, r) + 0 \cdot P(m, m))) \\ 358 &= P(m, r) - P(r, r). \end{aligned}$$

359 Appendix A.2. Condition for evolutionary stability

360 Starting from the condition in [4], we want to find a condition in terms of μ and σ^2 . We use K
 361 as the random variable having probability density function $\text{Pr}(k)$, and the notation $\langle K \rangle$ for expected
 362 value. Following the appendix from [4], we get to the condition

$$363 \quad \frac{B''(Nr^*)}{N} (\langle K \rangle^2 + \text{Var}[K]) - C''(r^*) < 0. \quad (\text{A2})$$

364 Changing variables from K to $\rho = K/N$, we get

$$365 \quad \frac{B''(Nr^*)}{N} (\langle N\rho \rangle^2 + \text{Var}[N\rho]) - C''(r^*) < 0. \quad (\text{A3})$$

366 Because of the rules of expected value and variance, this is the same as

$$367 \quad \frac{B''(Nr^*)}{N} (N^2 \langle \rho \rangle^2 + N^2 \text{Var}[\rho]) - C''(r^*) < 0, \quad (\text{A4})$$

368 which we simplify to

$$369 \quad NB''(Nr^*) (\langle \rho \rangle^2 + \text{Var}[\rho]) - C''(r^*) < 0, \quad (\text{A5})$$

370 and the desired form is achieved when noticing that $\langle \rho \rangle = \mu$ and $\text{Var}(\rho) = \sigma^2$.

371 Appendix A.3. Impossible region in the $\mu - \sigma^2$ -plane

372 What we want to find out is the upper bound on σ^2 in terms of μ . Firstly, since μ and σ^2 are
 373 defined through the fraction k/N where $N \geq k$ and $k \geq 1$, we note that $\mu, \sigma^2 \in [0, 1]$. Next, given
 374 $\mu \in [0, 1]$, by the definition of variance, the maximum variance will be achieved when all the values are
 375 at the endpoints. This is because as long as any value is in the interior of $[0, 1]$, the sum of the squares
 376 of the distances from μ , and hence the variance, will be lower. In other words, the maximum variance
 377 is achieved when we are dealing with a Bernoulli distribution, in which case $\sigma^2 = \mu(1 - \mu)$. So, given
 378 $\mu \in [0, 1]$, we must have $\sigma^2 \in [0, \mu(1 - \mu)]$.

379 Appendix A.4. Implicit differentiation

380 We want to find out the effects of relatedness on the location of the singular strategy in the
 381 general case, but having no explicit formula for r^* in terms of μ , we need to differentiate implicitly.
 382 We view r^* as a function of μ and we want to find the derivative $r^{*'}(\mu)$. To do this, we use the
 383 relationship $\mu B'(Nr^*) - C'(r^*) = 0$, and differentiate both sides implicitly with respect to μ : $B'(Nr^*) +$
 384 $\mu B''(Nr^*)Nr^{*'}(\mu) - C''(r^*)r^{*'}(\mu) = 0$, from which we get

$$385 \quad r^{*'}(\mu) = \frac{B'(Nr^*)}{C''(r^*) - \mu NB''(Nr^*)}. \quad (\text{A6})$$

386 Appendix B. Individual-based simulations

387 We follow the method from [18], with a few alterations. A population size N_{pop} and a virtual
 388 group size N is fixed, and each player has the two attributes generation and trait value. Next the
 389 following procedure is repeated until the generation of the offspring is higher than a chosen bound: A
 390 random focal individual with trait value x is chosen to be replaced by an offspring. The x individual
 391 gets to interact with another random individual with trait value u . The payoff to the focal individual,
 392 $P_x = P(x, u)$, is then computed as

$$393 \quad P(x, u) = \sum_{k=1}^N \frac{\text{Pr}(k)}{N} B(kx + (N - k)u) - C(x), \quad (\text{A7})$$

394 where the probabilities $\text{Pr}(k)$ are chosen such that $\sum_{k=1}^N \text{Pr}(k) = 1$, and such that $\sum_{k=1}^N k\text{Pr}(k)/N$ gets
 395 the desired value (e.g. 0.4 or 0.8). A third random individual with trait value y is chosen for an
 396 interaction with a fourth random individual with trait value v . The payoff to this other individual, P_y ,
 397 is computed in a similar way as P_x . P_x is then compared to P_y to see which individual is the parent to
 398 the offspring replacing the x individual. If $P_y < P_x$ then the x individual is the parent. Otherwise the y
 399 individual is the parent with a probability $w = (P_y - P_x)/\alpha$, where

$$400 \quad \alpha = \max_{0 \leq x, u, y, v \leq 1} |P(x, u) - P(y, v)| \quad (\text{A8})$$

401 to ensure $w \in [0, 1]$. The offspring inherits the parent's trait value at a high probability, otherwise
 402 the offspring's trait value is taken from a normal distribution centered on the parent's trait value. If
 403 the trait value gets lower than 0 then it's set to 0, and if it gets higher than 1 then it's set to 1. The
 404 generation and trait value of each individual is saved to a text file in order to keep the time complexity
 405 at a minimum.

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