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# Evolutionary graph theory revisited: when is an evolutionary process equivalent to the Moran process?

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Evolution in finite populations is often modelled using the classical Moran process. Over the last ten years this methodology has been extended to structured populations using evolutionary graph theory. An important question in any such population, is whether a rare mutant has a higher or lower chance of fixating (the fixation probability) than the Moran probability, i.e. that from the original Moran model, which represents an unstructured population. As evolutionary graph theory has developed, different ways of considering the interactions between individuals through a graph and an associated matrix of weights have been considered, as have a number of important dynamics. In this paper we revisit the original paper on evolutionary graph theory in light of these extensions to consider these developments in an integrated way. In particular we find general criteria for when an evolutionary graph with general weights satisfies the Moran probability for the set of six common evolutionary dynamics.

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# 1. Introduction

When modelling population evolution we are concerned with the spread of heritable characteristics in successive generations. The type of model that is used depends upon whether the population size is assumed to be finite or infinite. The majority of classical evolutionary models (see for example [1, 2]) use infinite populations, although finite population models are also well established, the most important models being those in [3, 4]. These models are stochastic, and are solved using classical Markov chain methodology [5, 6, 7]. See also [8, 9] for an extension 8 to evolutionary games in finite populations.

The populations in the models described above, however, were "well-mixed", i.e. every 10 individual was equally likely to encounter every other individual. Real populations of course 11 contain structural elements, such as geographical location or social relationship, which mean that 12 some pairs individuals are more likely to interact than others. In such circumstances we need to 13 14 be able to identify distinct individuals (or at least distinct classes of individuals), and considering finite populations is perhaps more natural than infinite ones (although finite structures each 15 containing an infinite number of individuals, so called "island models", were considered in [10]). 16 In [11] the modelling ideas of [3] were extended to consider such structured populations based 17 upon graphs, known as evolutionary graph theory. This has proved very successful, spawning a 18 large number of papers (for example [12, 13, 14, 15, 16, 17, 18, 19]). For informative reviews see 19 [20, 21]. 20

In an evolving population, we need to consider the mechanism of how the population changes, 21 22 called the dynamics. Informally, the dynamics specify the way in which heritable characteristics are passed on from one generation to the next. For infinite populations the classical replicator 23 equation [22] is often used (although there are a number of alternatives), and in the stochastic 24 model of [3] there is a natural replacement dynamics built in. For structured populations this issue 25 is actually considerably more complex, and the order of births and deaths, and where selection 26 acts, is of vital importance [23, 24]. We shall consider a set of dynamics that are commonly used 27 in evolutionary graph theory models. The relationship between dynamics and structure is of 28 key interest because the spread of heritable characteristics is directly dependent upon it. Whilst 29 having essentially no effect on populations with no structure, for constant fitness this relationship 30 potentially yields very different results on graphs. For non-constant fitness the results will vary 31 for different dynamics even in well-mixed populations [25]. 32

Under some circumstances it is, however, possible for the dynamics and structure to interact 33 in such a way that the spread of heritable characteristics behaves just as if the population was 34 homogeneous. This was a central theme of the classic paper [11], where two important results, 35 the circulation theorem and the isothermal theorem, were developed that addressed this question 36 (see also [26] for related work). In this paper we generalise the work of [11] to obtain a complete 37 classification of when the combination of a population structure and dynamics can be regarded 38 as equivalent to a homogeneous population in a precisely defined way, for the six most common 39 evolutionary dynamics and graphs with general weights. 40

#### 2. The Model 41

We shall first describe the population model of [11], which generalises the model of [3] by 42 incorporating a replacement structure. The notation used in this paper is summarised in Table 1. 43 The population has a constant size  $N \in \mathbb{Z}$ ,  $N \ge 2$ , consisting of individuals  $I_1, \ldots, I_N$ . Every individual 44 is either of type A or B. 45 This implies that there are  $2^N$  different states of the population given by the combination of 46 type A and B individuals. We represent each state by a set S such that  $n \in S$  if an individual  $I_n$ 47

is of type A. We can easily revert to using the number of type A individuals, |S|, if the population 48 is homogeneous. The states  $\emptyset$  and  $\mathcal{N} = \{1, 2, \dots, N\}$  have only type B and A individuals 49 respectively. 50

<sup>51</sup> Individuals have a constant fitness that may depend upon their type.

The fitness of individuals in state S is thus given by the vector  $\mathbf{F}(S) = (F_n(S))_{n=1,2,...,N}$ where

$$F_n(S) = \begin{cases} 1 & n \notin S, \\ r \in (0, \infty) & n \in S, \end{cases}$$

54 55

<sup>56</sup> is the fitness of  $I_n$ . Here the fitness r of a type A individual is given relative to the fitness of a type <sup>57</sup> B individual assumed to be 1.

<sup>58</sup> During a stochastic replacement event (that happens in an instant) an exact copy of an individual  $I_i$ <sup>59</sup> replaces an individual  $I_j$ .

The replacement events may be restricted in the sense that not all individuals can replace 60 one another. To enforce such restrictions, [11] imposed a replacement structure using a weighted 61 directed graph given by the tuple (D, w) where D = (V, E) is a directed graph, with sets V of 62 vertices and E of directed edges, and w is a map that assigns a weight to each edge such that 63  $w: V \times V \to [0,\infty): (i,j) \mapsto w_{ij}$ . Each vertex  $n \in V$  represents  $I_n$  therefore  $V = \{1, 2, \dots, N\}$  so 64 |V| = N. We assume that  $(i, j) \in E$  if and only if  $w_{ij} > 0$ , which indicates that  $I_i$  can replace  $I_j$ . 65 Note that we allow  $w_{ii} > 0$  and therefore  $I_i$  can replace itself. All the information contained within 66 the weighted digraph (D, w) is conveniently summarised by the  $N \times N$  weighted adjacency 67 matrix  $\mathbf{W} = (w_{ij})$  and therefore we will refer to (D, w) using  $\mathbf{W}$ , which we call the *replacement* 68 matrix. 69

The replacement events are stochastic which means that there is a probability  $\mathfrak{r}_{ij} = \mathfrak{r}_{ij}(\mathbf{F}(S), \mathbf{W})$  associated with (a copy of)  $I_i$  replacing  $I_j$ . There are several potential *evolutionary dynamics on graphs* that govern how the probability is determined. There three main types of dynamics that are summarised below, see also [21]. We use the convention that  $I_i$  is chosen for birth and  $I_j$  is chosen for death.

$b_i$ and then $(i, j) \in E_i$ is chosen with probability $d_{ij}$ , where $E_i$ are all edges starting in vertex <i>i</i> . $d_{ij}$ is used to signify that there is 'replacement by death'. Finally, $\mathfrak{r}_{ij} = b_i d_{ij}$ .
vertex <i>i</i> , $d_{ij}$ is used to signify that there is 'replacement by death'. Finally, $\mathbf{r}_{ij} = b_i d_{ij}$ .
$ = \sum_{i=1}^{n} \sum_{j=1}^{n} \sum_{i=1}^{n} \sum$
(ii) <i>Death-Birth</i> (DB): $I_j$ is chosen first then $I_i$ . We have that $j \in V$ is chosen with probability
$d_j$ and then $(i, j) \in E_j$ is chosen with probability $b_{ij}$ , where $E_j$ are all edges ending in
vertex <i>j</i> . $b_{ij}$ is used to signify that there is 'replacement by birth'. Finally, $r_{ij} = d_i b_{ij}$ .
(iii) Link (L): $I_i$ and $I_j$ are chosen simultaneously. In this case $(i, j) \in E$ is simply chosen with
probability $\mathfrak{r}_{ij}$ .

For each type of these dynamics, the natural selection can, through the fitness parameter, influence either the choice at birth (resulting in adding "B") or at death (adding "D"). It yields 6 kinds of evolutionary dynamics on graphs summarized in Table 2. These dynamics have been extensively studied, in particular, see [27] for a detailed comparison of them. Of these, the BDB and LB dynamics were used in [11].

## (a) The fixation probability

The fixation probability,  $\rho_S^A = \rho_S^A(*, \mathbf{W}, r)$ , is the probability that the population with initial state *S* is absorbed in  $\mathcal{N}$  where \* is the dynamics being used.

Given that the replacement events are random, the transitions between the states of the population are described by a stochastic process, which we denote  $\mathcal{E}$ . The properties of  $\mathcal{E}$ can be investigated once the state transition probabilities of moving from state *S* to *S'*,

 $P_{SS'} = P_{SS'}(*, \mathbf{W}, r)$ , are calculated using the replacement probabilities as follows: 94

$$P_{SS'} = \begin{cases} \sum_{\substack{i \notin S \\ i \notin S \\ i \in S \\ i \in S \\ i \in S \\ i, j \in S \\ \forall i, j \notin S \\ \forall i, j \notin S \\ \forall i, j \notin S \\ \end{cases} \text{ if } S' = S \cup \{j\} \text{ for some } j \notin S, \end{cases}$$

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The transition probabilities,  $P_{SS'}$ , satisfy the Markov property because they only depend upon 97 the state S, that is, the probability of transitioning from the present state to another state is 98 independent of any past and future state of the population. The stochastic process  $\mathcal{E}_{*,\mathbf{W},r}$  with 99 state transition matrix  $\mathbf{S} = \mathbf{S}(*, \mathbf{W}, r) = (P_{SS'})_{S,S' \subset \{1,2,\dots,N\}}$  is therefore a Markov chain. The 100 Markov chain  $\mathcal{E}_{*,\mathbf{W},r}$  is part of the class of evolutionary Markov chains described in [28]. 101

The absorbing states of  $\mathcal{E}_{*,\mathbf{W},r}$  are  $\emptyset, \mathcal{N}$ , which means that if the population is in either one of 102 these states then it remains there indefinitely. This property of  $\mathcal{E}_{*,\mathbf{W},r}$  can be used to measure the 103 success of a type A individual by calculating the probability that it fixates, that is, everyone in the 104 population is of type A. The fixation probability is then given by solving 105

106 
$$\rho_{S}^{A} = \sum_{S' \subset \{1, 2, \dots, N\}} P_{SS'} \rho_{S'}^{A}$$
(2.1)

108

with boundary conditions  $\rho_{\emptyset}^A = 0$  and  $\rho_{\mathcal{N}}^A = 1$ . As demonstrated in [27], LB and LD dynamics may differ in time scale but they yield the 109 same fixation probabilities when fitness is constant (which is our case). Thus, for our purposes 110 the dynamics are the same and we will thus consider them together and denote them by L. 111

Fixation probability is not the only measure for evolutionary success and we can look at the 112 fixation time [29, 30] as well. 113

#### (b) The Moran Process

The Moran process [3], a stochastic birth-death process on finite fixed homogenous population, 115 can be reconstructed as  $\mathcal{E}_{BDB, \mathbf{W}_{\mathbf{H}}, r}$  for a constant replacement matrix 116

$$\mathbf{W}_{\mathbf{H}} = (1/N)_{i,j}.\tag{2.2}$$

For any  $r \in (0, \infty)$  and any  $S \subset \{1, \ldots, N\}$ , the fixation probability for this process, or *Moran* 119 probability, is given by 120

$$\rho_{S}^{A} = \begin{cases} \frac{1 - r^{-|S|}}{1 - r^{-N}} & \text{if } r \neq 1, \\ |S|/N & \text{if } r = 1. \end{cases}$$

We are interested in characterizing graphs (and evolutionary dynamics) that yield the same 123 fixation probabilities as the homogeneous matrix  $W_H$  given in (2.2). We note that for this matrix 124 all of the transition probabilities  $r_{ij}$  take the same value independent of i, j or the dynamics, and 125 consequently the fixation probability under each of the dynamics is the same. 126

### (c) Classes of Graphs/ Matrices

The set of all admissible replacement matrices is defined as follows 128

$$W = \{ \mathbf{W} : \text{for every } i, j, \text{ there is } n \text{ such that } (\mathbf{W}^n)_{i,j} > 0 \}.$$

This definition means that  $\mathbf{W}$  is strongly connected as for any pair of vertices *i* and *j*, there is 131

a path (of length n) going from i to j. Unless specified otherwise, we will consider admissible 132

replacement matrices only. 133

As in [11], for any **W** (admissible or not) we define the *in temperature* of  $I_n$ ,  $T_n^-$ , and the *out* temperature of  $I_n$ ,  $T_n^+$ , by

$$T_n^- = \sum_{j=1}^N w_{jn}$$
 and  $T_n^+ = \sum_{j=1}^N w_{nj}$ .

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<sup>138</sup> **W** is called a *circulation* if  $T_n^+ = T_n^-$ , for all  $n \in V$  and it is called *isothermal* if  $T_i^+ = T_j^-$ , for all <sup>139</sup>  $i, j \in V$ . **W** is called *right* stochastic if  $T_n^+ = 1$ , for all  $n \in V$  and it is called *left* stochastic if  $T_n^- = 1$ , <sup>140</sup> for all  $n \in V$ . The sets of all circulations, isothermal matrices, right stochastic matrices, and left <sup>141</sup> stochastic matrices, respectively are denoted by  $W_C, W_I, W_R$ , and  $W_L$  respectively.

The set  $C_N$  denotes the sets of matrices representing *cycles* of length N, more specifically, for  $(w_{ij}) \in C_N$  we have  $w_{ii} = 1/2$  for i = 1, 2, ..., N,  $w_{i_1i_2} = \cdots = w_{i_ni_{n+1}} = \cdots = w_{i_{N-1}i_N} = w_{i_Ni_1} = 1/2$  for some permutation  $i_1, i_2, ..., i_N$  of the sequence 1, 2, ..., N, and  $w_{ij} = 0$  otherwise. We also define the maps  $f_R : W \to W_R$ ,  $f_L : W \to W_L$ , and  $f' : W \to W$  respectively, by

$$f_{R}\left((w_{ij})\right) = \left(\frac{w_{ij}}{\sum_{n} w_{in}}\right), \quad f_{L}\left((w_{ij})\right) = \left(\frac{w_{ij}}{\sum_{n} w_{nj}}\right), \quad \text{and} \quad f'\left((w_{ij})\right) = \left(\frac{w_{ij}}{\sum_{n,k} w_{nk}}\right)$$

<sup>148</sup> Note that  $f_R$  preserves right stochastic matrices and  $f_L$  preserves left stochastic matrices. <sup>149</sup> Moreover,  $f_R(\mathbf{W}) = f_L(\mathbf{W})$  for all  $\mathbf{W} \in W_I$ . Also, since f' simply involves multiplying  $\mathbf{W}$  by <sup>150</sup> the constant  $1/\sum_{n,k} w_{nk}$ , it implies that  $\mathbf{W} \in W_C \Leftrightarrow f'(\mathbf{W}) \in W_C$ .

<sup>151</sup> When the dynamics \*, matrices  $\mathbf{W}_1$  and  $\mathbf{W}_2$ , and fitness r are given, we say that an <sup>152</sup> evolutionary Markov chain  $\mathcal{E}_{*,\mathbf{W}_1,r}$  is  $\rho$ -equivalent to  $\mathcal{E}_{*,\mathbf{W}_2,r}$  if for every  $S \subset \{1,\ldots,N\}$ , <sup>153</sup>  $\rho_S^A(*,\mathbf{W}_1,r) = \rho_S^A(*,\mathbf{W}_2,r)$ , in which case we write  $\mathbf{W}_1 \sim_{*,r} \mathbf{W}_2$ .

We are specifically interested in finding matrices equivalent to the Moran process. For a dynamics \*, we define

$$M_* = \{ \mathbf{W} : \mathbf{W} \sim_{*,r} \mathbf{W}_{\mathbf{H}} \text{ for all } r > 0 \}.$$

### 3. Results

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The map  $f_R$  preserves the equivalence classes of BDB and BDD dynamics,  $f_L$  preserves the equivalence classes of DBB and DBD dynamics and f' preserves the equivalence classes for link dynamics. Specifically, as one can see from the proofs in the Appendix, for any **W** and any r > 0

$$\begin{array}{c} {}_{\mathbf{162}} & {}_{\mathbf{W}} \sim_{\mathrm{BDB},r} f_R(\mathbf{W}), \qquad (3.1) \\ \\ {}_{\mathbf{163}} & {}_{\mathbf{W}} \sim_{\mathrm{BDD},r} f_R(\mathbf{W}), \\ \\ {}_{\mathbf{164}} & {}_{\mathbf{W}} \sim_{\mathrm{DBD},r} f_L(\mathbf{W}), \\ \\ \\ {}_{\mathbf{165}} & {}_{\mathbf{W}} \sim_{\mathrm{DBB},r} f_L(\mathbf{W}), \\ \\ \\ {}_{\mathbf{167}} & {}_{\mathbf{W}} \sim_{\mathrm{L},r} f'(\mathbf{W}). \end{array}$$

We thus obtain the following results, which completely specify the graphs which are equivalent
 to the homogeneous matrix W<sub>H</sub> for each of our evolutionary dynamics.

Proposition 1 (Link).  $M_L = W_C$ . More precisely, the following statements are equivalent:

(a) **W** is a circulation.

172 (b) For all r > 0,  $\mathbf{W} \sim_{L,r} \mathbf{W}_{\mathbf{H}}$ .

(c) There is r > 0 such that  $\mathbf{W} \sim_{L,r} \mathbf{W}_{\mathbf{H}}$ .

We note that  $W_C = f'^{-1}(W_C) = {\mathbf{W} : f'(\mathbf{W}) \in W_C}$  and thus, similarly to Proposition 2 below, Proposition 1 can be written as  $M_L = f'^{-1}(W_C)$ .

**Proposition 2** (BDB and DBD).  $M_{BDB} = f_R^{-1}(W_C)$  and  $M_{DBD} = f_L^{-1}(W_C)$ . More precisely, the following statements are equivalent:

- (a)  $f_R(\mathbf{W})$  is a circulation.
- (b) For all r > 0,  $\mathbf{W} \sim_{BDB, r} \mathbf{W}_{\mathbf{H}}$ .
- (c) There is r > 0 such that  $\mathbf{W} \sim_{BDB,r} \mathbf{W}_{\mathbf{H}}$

<sup>181</sup> The equivalent conditions for DBD are similar to the above for BDB but  $f_R$  is replaced by  $f_L$ .

Proposition 3 (BDD and DBB).  $M_{BDD} = f_R^{-1}(\{\mathbf{W}_{\mathbf{H}}\} \cup C_N)$  and  $M_{DBB} = f_L^{-1}(\{\mathbf{W}_{\mathbf{H}}\} \cup C_N)$ . More precisely, the following statements are equivalent:

- (a)  $f_R(\mathbf{W}) = \mathbf{W}_{\mathbf{H}} \text{ or } f_R(\mathbf{W}) \in C_N.$ (b) For all r > 0,  $\mathbf{W} \sim_{BDD,r} \mathbf{W}_{\mathbf{H}}.$
- <sup>186</sup> The equivalent conditions for DBB are similar to the above for BDD but  $f_R$  is replaced by  $f_L$ .

In particular, M<sub>BDD</sub> ⊂ M<sub>BDB</sub> and M<sub>DBB</sub> ⊂ M<sub>DBD</sub>. The sets M<sub>\*</sub> are illustrated in Table 2.
 Note that unlike in Propositions 1 and 2, Proposition 3 does not contain "any *r* implies all *r*".
 In fact, when *r* = 1, there is no selection and thus the dynamics BDB and BDD are the same (and also the dynamics DBB and DBD are the same). Consequently, by Proposition 2,

 $\mathbf{W} \sim_{\text{BDD},1} \mathbf{W}_{\mathbf{H}} \Leftrightarrow f_R(\mathbf{W}) \in W_C \Leftrightarrow \mathbf{W} \in M_{\text{BDB}},$  $\mathbf{W} \sim_{\text{DBB},1} \mathbf{W}_{\mathbf{H}} \Leftrightarrow f_L(\mathbf{W}) \in W_C \Leftrightarrow \mathbf{W} \in M_{\text{DBD}}.$ 

192 193

### (a) Our results in the context of known results

For the LB dynamics, Proposition 1 was stated and proved in [11] as the Circulation theorem. For the LD dynamics, Proposition 1 follows from the Circulation theorem and the result of [27] that the fixation probabilities for LB and LD are the same.

As shown in Appendix (a), BDB is the same as the LB dynamics for right stochastic matrices (in particular, for BDB dynamics, Proposition 2 can be seen as the Isothermal theorem from [11]). Proposition 2 thus follows from Proposition 1 thanks to (3.1). The natural symmetries between  $f_R$ and  $f_L$  and BDB and DBD dynamics allow us to extend the Isothermal theorem to DBD dynamics as well (see also [31]).

<sup>203</sup> Overall, Propositions 1 and 2 and the occurrence of  $W_C$  within them are consistent with the <sup>204</sup> claim made in [11] that the circulation criterion completely classifies all replacement matrices <sup>205</sup> where  $\mathcal{E}_{*,\mathbf{W},r}$  is  $\rho$ -equivalent to a Moran process.

Our most important new result is Proposition 3. It shows that the BDD and DBB dynamics require very strict conditions to yield the Moran process. Either the population structure is homogeneous, or it is a directed cycle. This latter structure is an interesting theoretical example, but is unlikely to apply to real populations, meaning that the homogeneous population is practically the only way to get the Moran process for a realistic population.

#### (b) The importance of self-loops in BDD and DBB dynamics

Proposition 3 by definition requires that  $w_{ii} > 0 \quad \forall i = 1, 2, ..., N$ . Without such self-loops,  $\mathcal{E}_{BDD,\mathbf{W},r}, \mathcal{E}_{DBB,\mathbf{W},r}$  cannot ever be  $\rho$ -equivalent to the Moran process. The ability of an individual to replace itself therefore plays an important role in the replacement structure of the population and cannot be discounted. For BD dynamics, when increasing the diagonal weights of **W**, the fixation probability decreases for BDB and increases for BDD. For DB dynamics, the increase in fixation probability DBB is greater than that for DBD. For LB dynamics, the fixation probability remains the same.

With BDD and DBB evolutionary dynamics on graphs one may encounter the following 219 problems if there are no self-loops. For DBB dynamics, a type A individual with almost infinite 220 fitness still has a fixation probability bounded away from 1 because even type A individuals 221 can be randomly picked for death and replaced by type B individuals [32, page 245]. With self-222 loops, however, a type A individual will almost always be replaced by itself (or another type A 223 individual) and therefore has a fixation probability approaching 1. Similarly, for BDD dynamics, 224 a type A individual with almost zero fitness does not have near probability 0 of fixating as type 225 226 A individuals can be randomly picked for birth and replace type B individuals [32, page 245]. With self-loops, such an individual will almost always pick itself (or another type A) to replace 227 and therefore its fixation probability is near 0. Thus the inclusion of self-loops removes some 228 problematic features of the BDD and DBB dynamics, and makes them more attractive dynamics 229 to use in models. 230

### 231 4. Discussion

In this paper we have considered an evolutionary graph theory model of a population involving 232 general weights and a variety of evolutionary dynamics based upon the work of [11], which was 233 a development of the classical population model of [3]. In such populations, the population size 234 is fixed at all times and at successive discrete time points one replacement event occurs. Like the 235 aforementioned papers we consider two types of individuals, where fitness depends upon type 236 but no other factors (i.e. there are no game-theoretic interactions). In particular the single most 237 important property of such a process is the fixation probability, the probability that a randomly 238 placed mutant individual of the second type will eventually completely replace the population of 239 the first type. 240

This fixation probability depends upon the fitnesses of the two types of individuals, but can also be heavily influenced by the population structure as given by the weights, and by the evolutionary dynamics used. These effects are commonly observed, although in some circumstances evolution proceeds as if as on a well-mixed population as from the original work of [3], dependent only upon the fitnesses of the two types, and some important results in this regard were already given in [11]. The aim of this paper was to provide a generalised set of conditions for when this would be the case.

By defining what is meant by fixation-equivalence to the Moran process, we provided a general result which, independent of the specific dynamics used, helps identify graphs that do not affect the fixation probability. With respect to each of the standard dynamics, we then classified sets of evolutionary graphs that have the same fixation probability as the Moran process (or well mixed population). These sets include graphs that are circulations and therefore generalises the work of [11].

An important new result shows that the set of weights for which we obtain fixation equivalence to the Moran process for the BDD and DBB dynamics is very restricted, and so that for most populations with any structure this equivalence will not hold for these dynamics. We note also that the inclusion of non-zero self weights  $w_{ii}$  eliminates some problematic features of these two dynamics (i.e. that individuals with 0 fitness could fixate or those with infinite fitness could be eliminated) and so improves the applicability of these dynamics.

Presenting evolutionary dynamics on graphs in the way that we have allows one to incorporate a variety of dynamics in their analysis, both of standard type and other definitions. This will improve our understanding of dynamics on graphs in general. We note that the list of dynamics in Table 2 is not exhaustive. For example, [33] used imitation dynamics, which is a class of DBB dynamics with an additional requirement  $w_{ii} > 0 \quad \forall i$ , and [34] consolidates the BDB and DBD dynamics such that one is chosen with a given probability.

In general the inclusion of non-zero self weights, in contrast to many earlier evolutionary graph theory works, allows for a greater flexibility of modelling. We note that this is consistent with the original work of [3], which allowed self-replacement as an integral part of the process. For well-mixed populations it does not matter much whether this possibility is included or not

- <sup>270</sup> (at least for sufficiently large populations with intermediate fitness values), and it is likely that
- 271 it has often been excluded for reasons of convenience because of this without the ramifications
- <sup>272</sup> being fully considered in many later works. It is thus important to consider whether to include
- <sup>273</sup> such self weights when modelling spatial structure using evolutionary graph theory.
- 274 Data accessibility. There is no supporting data for this article.
- <sup>275</sup> Conflict of interests. We have no competing interests.
- Author's contributions. KP developed the original concept in discussion with MB, and carried out
- $_{\it 277}$   $\,$  the majority of the analysis and writing. MB, JR and LJS have all been closely involved in refining the
- $_{278}$  paper in terms of both analysis and presentation. In particular JR did significant work on the proofs, MB
- <sup>279</sup> on the Introduction/Discussion and LJS on the scientific presentation. All authors gave final approval for <sup>280</sup> publication.
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# **Appendix**

364 A. Proofs

#### (a) BDB is the same as LB for right stochastic matrices

- For BDB dynamics we have  $r_{ij} = b_i d_{ij}$ . By definition  $\sum_{ij} b_i d_{ij} = 1$ , we can therefore write this as
- <sup>367</sup>  $\mathfrak{r}_{ij} = b_i d_{ij} / \sum_{n,k} b_n d_{n,k}$ . Substituting  $b_i = F_i / \sum_{m=1}^N F_m$  gives

$$\mathfrak{r}_{ij} = \frac{d_{ij}F_i \left/ \sum_{m=1}^N F_m}{\sum_{n,k} \left( d_{nk}F_n \left/ \sum_{m=1}^N F_m \right) \right)} = \frac{d_{ij}F_i}{\sum_{n,k} d_{nk}F_n}.$$

If W is right stochastic, i.e.  $\sum_{n=1}^{N} w_{in} = 1$  for all i = 1, 2, ..., N, for BDB dynamics we have 370 that  $d_{ij} = w_{ij} / \sum_{n=1}^{N} w_{in} = w_{ij}$  giving  $\mathfrak{r}_{ij} = w_{ij}F_i / \sum_{n,k} w_{nk}F_n$  which is the LB dynamics as 371 required. We also have that DBD is the same as LD for left stochastic matrices. The explanation 372 follows the same procedure as above. 373

#### (b) Lemma 1 (Forward Bias) 374

The key Lemma 1 stated below is used in the proofs of all propositions and it relies heavily on the 375 notion of *forward bias* of state S which is then given by the ratio of the probabilities of a forward 376 transition to a backward transition from S. A forward and backward transition from S occurs 377 when the number of type A individuals increase and decrease by one respectively, which happen 378 with probability 379

$$P_S^+ = \sum_{n \notin S} P_{S,S \cup \{n\}} \quad \text{and} \quad P_S^- = \sum_{n \in S} P_{S,S \setminus \{n\}}.$$

**Lemma 1** (Constant Forward Bias). Let  $\mathcal{E}$  be an evolutionary process on states  $S \subset \{1, 2, \dots, N\}$  with 382 transition probabilities  $P_{S,S'}$  that satisfy 383

•  $P_{S,S'} > 0$  only if S and S' differ in at most one element 384

• for every  $S \neq \emptyset$ ,  $\{1, \ldots, N\}$ , there are  $S^+$  and  $S^-$  such that  $|S^+| = |S| + 1$  and  $|S^-| = |S| - 1$ 385 and  $P_{S,S^+} > 0, P_{S,S^-} > 0.$ 

Then, the following are equivalent 387

a) There is a constant 
$$c > 0$$
 such that for all  $S \subset \{1, 2, ..., N\}$ 

$$\rho_{S}^{A} = \begin{cases} \frac{1 - c^{-|S|}}{1 - c^{-N}} & \text{if } c \neq 1, \\ |S|/N & \text{if } c = 1 \end{cases}$$

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*b)*  $\mathcal{E}$  has constant forward bias, that is, there is a constant d such that for all  $S \subset \{1, 2, \dots, N\}$ 

$$P_S^{+} / P_S^{-} = d$$

Moreover, if either (a) or (b) hold, then c = d. 394

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$$r \sum_{a \in S} \sum_{b \notin S} w_{ab} / \sum_{a \in S} \sum_{b \notin S} w_{ba} ,$$

which is what one gets when using Link dynamics, or BDB dynamics if  $\mathbf{W} \in W_R$ . Note that in 398 Lemma 1 the forward bias is defined independent of the dynamics and therefore applies to all 399 dynamics that satisfy the assumptions. 400

Note that a similar result is given in [11, 20] where the forward bias is explicitly defined as

401 *Proof.* "(a) 
$$\Rightarrow$$
 (b)": Take any  $S \subset \{1, 2, \dots, N\}$ . It is known that

$$\rho_{S}^{A} = \sum_{S'} P_{S,S'} \rho_{S'}^{A} = P_{S,S} \rho_{S}^{A} + \sum_{n \notin S} \left( P_{S,S \cup \{n\}} \rho_{S \cup \{n\}}^{A} \right) + \sum_{n \in S} \left( P_{S,S \setminus \{n\}} \rho_{S \setminus \{n\}}^{A} \right)$$

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and using  $P_{S,S} = 1 - P_S^+ - P_S^-$  gives 404

$$0 = \sum_{n \notin S} \left( P_{S, S \cup \{n\}} \left( \rho_{S \cup \{n\}}^{A} - \rho_{S}^{A} \right) \right) + \sum_{n \in S} \left( P_{S, S \setminus \{n\}} \left( \rho_{S \setminus \{n\}}^{A} - \rho_{S}^{A} \right) \right).$$
(A 1)

For  $c \neq 1$ , equation (A 1) simplifies to 407

$$0 = \frac{1 - c^{-|S| - 1} - 1 + c^{-|S|}}{1 - c^{-N}} P_S^+ + \frac{1 - c^{-|S| + 1} - 1 + c^{-|S|}}{1 - c^{-N}} P_S^- =$$

$$P_{S}^{+} / P_{S}^{-} = \frac{c^{-|S|} - c^{-|S|+1}}{c^{-|S|-1} - c^{-|S|}} = \frac{1-c}{c^{-1}-1} = c$$

For c = 1, equation (A 1) simplifies to 411

$$\begin{array}{c} _{^{412}} \\ _{^{413}} \end{array} \qquad 0 = (|S|+1-|S|)P_S^+ + (|S|-1-|S|)P_S^- \Rightarrow P_S^+ \left/ P_S^- = 1. \right.$$

"(b)  $\leftarrow$  (a)": The state transition matrix  $\mathbf{S} = (P_{S,S'})$  can be scaled to give  $\mathbf{S}' = (P'_{S,S'})$  such that 414  $P'_{S,S} = 0$  and  $P'_{S,S'} = P_{S,S'}/(1 - P_{S,S}) = P_{S,S'}/(P_S^+ + P_S^-)$  where S is a non-absorbing state. 415 The fixation probability  $\rho_S^A$  will be the same whether S' or S is used. This is because equation 416 (2.1) can be rearranged as follows 417

418 
$$\rho_S^A = \sum_{S'} P_{SS'} \rho_{S'}^A \Rightarrow \quad \rho_S^A = P_{SS} \rho_S^A + \sum_{S': S' \neq S} P_{SS'} \rho_{S'}^A \Rightarrow$$

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 $\rho_{S}^{A}(1-P_{SS}) = \sum_{S' \cdot S' \neq S} P_{SS'} \rho_{S'}^{A} \Rightarrow \quad \rho_{S}^{A} = \sum_{S' \cdot S' \neq S} \frac{P_{SS'}}{P_{S}^{+} + P_{S}^{-}} \rho_{S'}^{A}.$ 420

Let  $\{S_0, S_1, \ldots, S_N\}$  be a partition of the states S such that  $S \in S_i$  if |S| = i. The probability 421  $P_{i,j}(S)$  of transitioning from state  $S \in S_i$  to *lumped state*  $S_j$  with respect to **S'** is 422

$$P_{i,j}(S) = \begin{cases} 0 & j \neq i \pm 1, \\ 1/(d+1) & j = i-1, \\ d/(d+1) & j = i+1 \end{cases}$$
(A 2)

This can be easily verified, for example, take j = i - 1 then 425

$$P_{i,i-1}(S) = \sum_{S' \in \mathcal{S}_{i-1}} P'_{S,S'} = \sum_{S' \in \mathcal{S}_{i-1}} \frac{P_{S,S'}}{P_S^+ + P_S^-} = \frac{P_S^-}{P_S^+ + P_S^-} = \frac{1}{1+d}$$

since the forward bias is equal to d. Equation (A 2) satisfies the necessary and sufficient condition 428 for the Markov chain with state transition matrix  $\mathbf{S}'$  to be lumpable with respect to the partition 429  $\{S_0, S_1, \dots, S_N\}$  (Theorem 6.3.2 page 124, [35]). Let  $\hat{\mathbf{S}} = (P_{i,j})$  be the state transition matrix for 430 this lumped Markov chain then the probability  $P_{i,j}$  of transitioning from lumped states  $S_i$  to  $S_j$ 431 is given by 432

$$P_{i,j} = P_{i,j}(S)$$

The state transition matrix  $\hat{\mathbf{S}}$  describes a random walk with absorbing barriers and therefore the 435 probability  $\rho_i^A$  of type A individuals fixating when the population starts in lumped state  $S_i$  is 436 calculated using the methods in [5] to give 437

$$\rho_i^A = 1 + \sum_{j=1}^{i-1} \prod_{k=1}^j \frac{P_{k,k-1}}{P_{k,k+1}} / 1 + \sum_{j=1}^{N-1} \prod_{k=1}^j \frac{P_{k,k-1}}{P_{k,k+1}} .$$

In this case, 440

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$$\rho_i^A = \begin{cases} \frac{1-d^{-i}}{1-d^{-N}} & d \neq 1, \\ i/N & d = 1 \end{cases}$$

since  $P_{k,k-1}/P_{k,k+1} = 1/r$  for k = 1, 2, ..., N - 1. By definition,  $\rho_S^A = \rho_i^A$  where i = |S| as 443 required. 444

#### (c) Proposition 1 (Link) 445

The following statements are equivalent: 446

- (a) **W** is a circulation. 447
- (b) For all r > 0,  $\mathbf{W} \sim_{\mathrm{L},r} \mathbf{W}_{\mathbf{H}}$ . 448
- (c) There is r > 0 such that  $\mathbf{W} \sim_{\mathbf{L},r} \mathbf{W}_{\mathbf{H}}$ . 449
- (d) For all r > 0 and for all  $S \subset \{1, 2, ..., N\}$ , the forward bias of  $\mathcal{E}_{L, \mathbf{W}, r}$  is r, i.e. 450
- $P_S^+ \left/ P_S^- = r. \right.$ 451 452
- (e) There is r > 0 such that for all  $a \in \{1, 2, ..., N\}$ , the forward bias of the one element set 453  $S = \{a\}$  is r, i.e. 454

$$\frac{\displaystyle\sum_{b\neq a} P_{\{a\},\{a,b\}}}{\displaystyle P_{a,\emptyset}} = r.$$

*Proof.* For LB dynamics the forward bias is given by 457

$$\frac{P_S^+}{P_S^-} = \frac{\sum\limits_{a \in S} \sum\limits_{b \notin S} \frac{w_{ab}F_a}{\sum\limits_{n,k} w_{nk}F_n}}{\sum\limits_{a \in S} \sum\limits_{b \notin S} \frac{w_{ba}F_b}{\sum\limits_{n,k} w_{nk}F_n}} = \frac{r \sum\limits_{a \in S} \sum\limits_{b \notin S} w_{ab}}{\sum\limits_{a \in S} \sum\limits_{b \notin S} w_{ba}}$$

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For LD dynamics the forward bias is given by 460

$$\frac{P_S^+}{P_S^-} = \frac{\sum_{a \in S} \sum_{b \notin S} \frac{w_{ab}/F_b}{\sum w_{nk}/F_k}}{\sum_{a \in S} \sum_{b \notin S} \frac{w_{ba}/F_a}{\sum \sum_{n,k} w_{nk}/F_k}} = \frac{r \sum_{a \in S} \sum_{b \notin S} w_{ab}}{\sum_{a \in S} \sum_{b \notin S} w_{ba}}.$$

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"(a)  $\Rightarrow$  (d)": W is a circulation i.e.  $T_n^+ = T_n^-$  for all  $n \in \{1, \dots, N\}$  and thus 463

$$\sum_{a \in S} \sum_{b \notin S} w_{ab} = \sum_{a \in S} \left( \sum_{n} w_{an} - \sum_{k \in S} w_{ak} \right) = \sum_{a \in S} \left( T_a^+ - \sum_{k \in S} w_{ak} \right) \Rightarrow$$

$$\sum_{a \in S} \sum_{b \notin S} w_{ab} = \sum_{a \in S} \left( T_a^- - \sum_{k \in S} w_{ka} \right) = \sum_{a \in S} \left( \sum_n w_{na} - \sum_{k \in S} w_{ka} \right) \Rightarrow$$

$$\sum_{a \in S} \sum_{b \notin S} w_{ab} = \sum_{a \in S} \sum_{b \notin S} w_{ba}.$$

Note that  $\sum_{a \in S} \sum_{b \notin S} w_{ab} \neq 0$  because W is admissible and represents a strongly connected 468 graph. Thus, the forward bias for both LB and LD is equal to *r*. 469

"(d) $\Rightarrow$ (e)" is trivial as (d) is much stronger than (e). 470

"(e) $\Rightarrow$ (a)" Let *a* and *r* is fixed. By above calculations of the forward bias, we have 471

$$\sum_{472} \sum_{b \notin S = \{a\}} w_{ab} = \sum_{b \notin S = \{a\}} w_{ba} \Rightarrow -w_{aa} + \sum_{i=1}^{N} w_{ai} = -w_{aa} + \sum_{i=1}^{N} w_{ia} \Rightarrow \sum_{i=1}^{N} w_{ai} = \sum_{i=1}^{N} w_{ia}$$

therefore **W** is a circulation. 474

"(d) $\Rightarrow$ (b)" follows from Lemma 1. 475

"(b) $\Rightarrow$ (c)" is trivial. 476

"(c) $\Rightarrow$ (e)" follows from Lemma 1. 477

### (d) Proposition 2 (BDB and DBD)

- <sup>479</sup> More precisely, the following statements are equivalent:
- (a)  $f_R(\mathbf{W})$  is a circulation.
- (b) For all r > 0,  $\mathbf{W} \sim_{\text{BDB},r} \mathbf{W}_{\mathbf{H}}$ .
- (c) There is r > 0 such that  $\mathbf{W} \sim_{\text{BDB},r} \mathbf{W}_{\mathbf{H}}$
- (d) For all r > 0 and for all  $S \subset \{1, 2, ..., N\}$ , the forward bias of  $\mathcal{E}_{\text{BDB}, \mathbf{W}, r}$  is r, i.e.

$$P_S^+ \left/ P_S^- = r \right.$$

(e) There is r > 0 such that for all  $a \in \{1, 2, ..., N\}$ , the forward bias of  $\mathcal{E}_{\text{BDB}, \mathbf{W}, r}$  of the one element set  $S = \{a\}$  is r, i.e.

$$\frac{\displaystyle\sum_{b\neq a} P_{\{a\},\{a,b\}}}{P_{a,\emptyset}} = r$$

<sup>490</sup> *Proof.* Let  $\mathbf{U} = (u_{ij}) = f_R(\mathbf{W}) = (w_{ij}/\sum_n w_{in})$  then for BDB dynamics the forward bias of <sup>491</sup>  $\mathcal{E}_{BDB,\mathbf{W},r}$  is given by

$$\frac{P_S^+}{P_S^-} = \frac{\sum\limits_{a \in S} \sum\limits_{b \notin S} \frac{F_a}{\sum F_n} \frac{w_{ab}}{\sum w_{an}}}{\sum\limits_{a \in S} \sum\limits_{b \notin S} \frac{F_b}{\sum F_n} \frac{w_{ba}}{\sum w_{ba}}} = \frac{r \sum\limits_{a \in S} \sum\limits_{b \notin S} u_{ab}}{\sum\limits_{b \notin S} \sum\limits_{a \in S} u_{ba}}$$

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and therefore the forward bias of  $\mathcal{E}_{\text{BDB},\mathbf{W},r}$  is the same as forward bias of  $\mathcal{E}_{\text{BDB},\mathbf{U},r}$ .

Similarly, with almost identical working as above, when  $\mathbf{V} = f_L(\mathbf{W})$ , the forward bias of  $\mathcal{E}_{\text{DBD},\mathbf{W},r}$  is the same as forward bias of  $\mathcal{E}_{\text{DBD},\mathbf{V},r}$  and is given by

$$-\frac{P_{S}^{+}}{P_{S}^{-}} = \frac{\sum_{a \in S} \sum_{b \notin S} \frac{1/F_{b}}{\sum 1/F_{n}} \frac{w_{ab}}{\sum w_{nb}}}{\sum_{a \in S} \sum_{b \notin S} \frac{1/F_{a}}{\sum n} \frac{w_{ba}}{\sum n}} = \frac{\sum_{a \in S} \sum_{b \notin S} v_{ab}}{\frac{1}{r} \sum_{a \in S} \sum_{b \notin S} v_{ba}}.$$

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and the proof of the Proposition for DBD closely follows the one for BDB given below with U and  $f_R$  appropriately replaced by V and  $f_L$ .

"(a)  $\Rightarrow$  (d)": If  $\mathbf{U} = f_R(\mathbf{W}) \in W_C$ , i.e. if  $\mathbf{U}$  is doubly stochastic, then the forward bias (for  $S \neq \emptyset, \mathcal{N}$ ) is equal to

$$\frac{P_{S}^{+}}{P_{S}^{-}} = \frac{r \sum_{a \in S} \left(\sum_{n} (u_{an}) - \sum_{k \in S} (u_{ak})\right)}{\sum_{a \in S} \left(\sum_{n} (u_{na}) - \sum_{k \in S} (u_{ka})\right)} = \frac{r \left(|S| - \sum_{a \in S} \sum_{k \in S} u_{ak}\right)}{|S| - \sum_{a \in S} \sum_{k \in S} u_{ka}} = r$$

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 $(d) \Rightarrow (e)$  is trivial as (d) is stronger than (e).

<sup>507</sup> "(e) $\Rightarrow$ (a)" Let *a* and *r* is fixed. By above calculations of the forward bias, we have

$$\sum_{a \in S} \sum_{b \notin S} u_{ab} = \sum_{a \in S} \sum_{b \notin S} u_{ba}$$

Consider the states  $S = \{a\}$  in which there is only one individual of type A then 510

$$\sum_{a} \sum_{b \notin S} u_{ab} = \sum_{b \notin S} u_{ba} \Rightarrow -u_{aa} + \sum_{i=1}^{N} u_{ai} = -u_{aa} + \sum_{i=1}^{N} u_{ia} \Rightarrow 1 = \sum_{i=1}^{N} u_{ia}$$

is true for all a = 1, 2, ..., N and therefore **U** is doubly stochastic and thus  $f_R(\mathbf{W})$  is a circulation. 513

- "(d) $\Rightarrow$ (b)" follows from Lemma 1. 514
- "(b) $\Rightarrow$ (c)" is trivial. 515
- "(c) $\Rightarrow$ (e)" follows from Lemma 1. 516

#### (e) Proposition 3 (BDD and DBB) 517

The following statements are equivalent: 518

(a) 
$$f_R(\mathbf{W}) = \mathbf{W}_{\mathbf{H}}$$
 or  $f_R(\mathbf{W}) \in C_N$ .  
(b) For all  $r > 0$ ,  $\mathbf{W} \sim_{\text{BDD},r} \mathbf{W}_{\mathbf{H}}$ .

*Proof.* The replacement probabilities  $r_{ij}(\mathbf{F}(S), \mathbf{W})$  for BDD dynamics can be rewritten as 521  $\mathfrak{r}_{ij}(\mathbf{F}(S), \mathbf{U})$  where  $\mathbf{U} = (u_{ij}) = f_R(\mathbf{W}) = (w_{ij} / \sum_n w_{in})$  by multiplying the numerator and 522 denominator with  $\sum_n w_{in}$  as follows 523

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$$\mathfrak{r}_{ij}(\mathbf{F}(S), \mathbf{W}) = \frac{1}{N} \frac{w_{ij}/F_j(S)}{\sum_n w_{in}/F_n(S)} = \frac{1}{N} \frac{w_{ij}/\left(F_j(S)\sum_n w_{in}\right)}{\sum_n w_{in}/\left(F_n(S)\sum_n w_{in}\right)} \Rightarrow$$

$$\sum_{525}_{526} \frac{u_{ij}/F_j(S)}{\sum_n u_{in}/F_n(S)} = \mathfrak{r}_{ij}(\mathbf{F}(S), \mathbf{U})$$

and therefore we have that  $\mathbf{W} \sim_{\text{BDD},r} \mathbf{U}$ , for all r > 0. The forward bias using  $\mathbf{U}$  for state S is 527 given by 528

$$\frac{P_{S}^{+}}{P_{S}^{-}} = \frac{\sum_{a \in S} \sum_{b \notin S} \frac{1}{N} \frac{u_{ab}/F_{b}}{\sum_{n} u_{an}/F_{n}}}{\sum_{a \in S} \sum_{b \notin S} \frac{1}{N} \frac{u_{ba}/F_{a}}{\sum_{n} u_{bn}/F_{n}}} = \frac{\sum_{a \in S} \sum_{b \notin S} \frac{u_{ab}}{\sum_{n} u_{an}/F_{n}}}{\frac{1}{r} \sum_{a \in S} \sum_{b \notin S} \frac{u_{ba}}{\sum_{n} u_{bn}/F_{n}}}.$$
 (A 3)

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Similarly, let  $\mathbf{V} = (v_{ij}) = f_L(\mathbf{W}) = (w_{ij} / \sum_n w_{nj})$ . Then for DBB dynamics we have 531

$$b_{ij} = \frac{w_{ij}F_i}{\sum_n w_{nj}F_n} = \frac{w_{ij}F_i / \sum_n w_{nj}}{\sum_n w_{nj}F_n / \sum_n w_{nj}} = \frac{v_{ij}F_i}{\sum_n v_{nj}F_n}$$

and therefore the forward bias when using  ${\bf V}$  is given by 534

$$\frac{P_S^+}{P_S^-} = \frac{\sum\limits_{a \in S} \sum\limits_{b \notin S} \frac{1}{N} \frac{v_{ab} F_a}{\sum\limits_n v_{nb} F_n}}{\sum\limits_{a \in S} \sum\limits_{b \notin S} \frac{1}{N} \frac{v_{ba} F_b}{\sum\limits_n v_{na} F_n}} = \frac{r \sum\limits_{a \in S} \sum\limits_{b \notin S} \frac{v_{ab}}{\sum\limits_n v_{nb} F_n}}{\sum\limits_{a \in S} \sum\limits_{b \notin S} \frac{1}{\sum\limits_n v_{na} F_n}}$$

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The proof of the Proposition for DBB closely follows the one for BDD given below with U and  $f_R$ 537 appropriately replaced by  $\mathbf{V}$  and  $f_L$ . 538

#### (i) If $\mathbf{U} \in C_N$ , then $\mathbf{U} \sim_{\mathsf{BDD},r} \mathbf{W}_{\mathbf{H}}$ 539

If  $\mathbf{U} \in C_N$  then there are only two nonzero elements in each row. In particular, in row *i* of **U** we 540

have that  $u_{ii}, u_{ik_i} = 1/2$  for some  $k_i \neq i$ . In the numerator of equation (A 3) for  $a \in S$ ,  $b \notin S$  and 541

 $k_a \neq a$  we have that for all *S* 542

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$$\frac{u_{ab}}{\sum_{n} u_{an}/F_n(S)} = \frac{u_{ab}}{u_{aa}/F_a(S) + u_{ak_a}/F_{k_a}(S)} = \begin{cases} 0 & \text{if } b \neq k_a, \\ \frac{1/2}{1/2r + 1/2} & \text{if } b = k_a. \end{cases}$$

Similarly, in the denominator of equation (A 3) for  $a \in S$ ,  $b \notin S$  and  $k_b \neq b$  we have that for all S 545

$$\frac{u_{ba}}{\sum_{n} u_{bn}/F_n(S)} = \frac{u_{ba}}{u_{bb}/F_b(S) + u_{bk_b}/F_{k_b}(S)} = \begin{cases} 0 & \text{if } a \neq k_b, \\ \frac{1/2}{1/2 + 1/2r} & \text{if } a = k_b. \end{cases}$$

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This means that equation (A 3) for all S can be written as 548

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$$\frac{x/2}{1/2r+1/2} \left/ \frac{1}{r} \frac{y/2}{1/2+1/2r} = rx/y \right.$$

where x(y) is the number of nonzero  $u_{ab}(u_{ba})$  terms in the numerator (denominator). If we 551 partition the vertices of the digraph of U into any two sets  $V_1, V_2$  then the number of edges e(i, j)552 and e(j, i) for  $i \in V_1$  and  $j \in V_2$  are by definition the same because it is a cycle. This means that 553 for  $a \in S$  and  $b \notin S$  the number of nonzero  $u_{ab}, u_{ba}$  terms in the numerator and denominator 554 respectively are the same hence x = y and rx/y = r as required. As per Lemma 1,  $\mathcal{E}_{BDD,U,r}$  is 555  $\rho$ -equivalent to the Moran process. 556

#### (ii) If $\mathbf{U} \sim_{\mathsf{BDD},r} \mathbf{W}_{\mathbf{H}}$ for all r > 0, then $\mathbf{U} = \mathbf{W}_{\mathbf{H}}$ or $\mathbf{U} \in C_N$ 557

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By Lemma 1, the forward bias (A 3) is equal to *r* for all  $S \subset \{1, ..., N\}$  giving 558

$$\sum_{a \in S} \sum_{b \notin S} \frac{u_{ab}}{\sum_{n} u_{an}/F_n} = \sum_{a \in S} \sum_{b \notin S} \frac{u_{ba}}{\sum_{n} u_{bn}/F_n} \Rightarrow$$

$$\sum_{a \in S} \frac{\sum_{b \notin S} u_{ab}}{\sum_{j \notin S} u_{aj} + \frac{1}{r} \sum_{i \in S} u_{ai}} = \sum_{b \notin S} \frac{\sum_{a \in S} u_{ba}}{\sum_{j \notin S} u_{bj} + \frac{1}{r} \sum_{i \in S} u_{bi}}.$$
(A 4)

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Note that if r = 1, (A 4) holds for all  $U \in W_C$ . From now, we will consider  $r \neq 1$  only. For clarity, 562

the remainder of this section of the proof is broken down into the following six steps. 563

#### Step 1: Derivation of general state dependent row-sum equation 564

Let 
$$U(a, S) = \sum_{i \in S} u_{ai}$$
, i.e.  $1 - U(a, S) = \sum_{j \notin S} u_{aj}$ . Equation (A 4) thus becomes

$$\sum_{a \in S} \frac{1 - U(a, S)}{1 - U(a, S) + U(a, S)/r} = \sum_{b \notin S} \frac{U(b, S)}{1 - U(b, S) + U(b, S)/r} \Rightarrow$$

$$\sum_{a \in S} \frac{1}{1 + U(a, S)(1/r - 1)} = \sum_{n=1}^{N} \frac{U(n, S)}{1 + U(n, S)(1/r - 1)}.$$
 (A 5)

Equation (A 5) can be written as a Taylor series as follows 569

$$\sum_{a \in S} \sum_{k=0}^{\infty} (-1)^{k} (1/r - 1)^{k} [U(a, S)]^{k} = \sum_{n=1}^{N} U(n, S) \sum_{k=0}^{\infty} (-1)^{k} (1/r - 1)^{k} [U(n, S)]^{k} \Rightarrow$$

$$\sum_{a \in S} \sum_{k=0}^{\infty} (1 - 1/r)^{k} [U(a, S)]^{k} = \sum_{n=1}^{N} \sum_{k=0}^{\infty} (1 - 1/r)^{k} [U(n, S)]^{k+1}$$
(A)

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(A 6)

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For equation (A 6) to hold for all r the coefficients of  $(1 - 1/r)^k$  should be same, that is, for all k

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$$\sum_{a \in S} [U(a,S)]^k = \sum_{n=1}^N [U(n,S)]^{k+1}.$$
 (A 7)

#### $_{\rm 576}$ Step 2: The diagonal of U consists of non-zero elements

<sup>577</sup> Consider the state  $S = \{a\}$  then equation (A 7) gives

$$u_{aa}^{k} = \sum_{n=1}^{N} u_{na}^{k+1}.$$
 (A 8)

<sup>580</sup> If  $u_{aa} = 0$  or 1 then (A 8) implies that all off-diagonal terms in column n are zero which is a <sup>581</sup> contradiction with **W** (and thus also  $\mathbf{U} = f_R(\mathbf{W})$ ) being strongly connected, which means that <sup>582</sup>  $0 < u_{aa} < 1$ .

### Step 3: The $n^{ ext{th}}$ column of ${f U}$ contains $m_n$ nonzero elements, all equal to $1/m_n$

Since  $0 < u_{aa} < 1$ , we can divide equation (A 8) by  $u_{aa}^k$  giving

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$$1 = \sum_{n=1}^{N} u_{na} \left(\frac{u_{na}}{u_{aa}}\right)^{k}.$$
 (A 9)

587 We have that

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$$\lim_{k \to \infty} \left( \frac{u_{na}}{u_{aa}} \right)^k = \begin{cases} \infty & u_{na} > u_{aa}, \\ 1 & u_{na} = u_{aa}, \\ 0 & u_{na} < u_{aa}, \end{cases}$$
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and therefore (A 9) implies that  $0 \le u_{na} \le u_{aa}$ . There must be  $n \ne a$  such that  $u_{na} = u_{aa}$  as otherwise, by (A 9), we would have  $u_{aa} = 1$ . Let  $C_a = \{i : u_{ia} = u_{aa}\}$ . (A 9) becomes

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593 
$$1 = \left(\sum_{i \in C_a} u_{aa}\right) + \left(\sum_{j \notin C_a} \frac{u_{ja}^{k+1}}{u_{aa}^k}\right) = |C_a|u_{aa} + \left(\sum_{j \notin C_a} \frac{u_{ja}^{k+1}}{u_{aa}^k}\right).$$
(A 10)

As  $k \to \infty$ , (A 10) implies that  $u_{aa} = 1/|C_a|$ . Thus, again by (A 10),  $u_{ja} = 0$  for all  $j \notin C_a$ . This means that in column n of **U** there should be  $m_n = |C_n|$  with  $2 \le m_n \le N$  nonzero elements, including  $u_{nn}$ , that are all equal to  $1/m_n$ .

#### 597 Step 4: $m_n$ is the same for all n

<sup>598</sup> Considering state  $S = \{i, j\}$  and using  $u_{aa} = 1/m_{a}$ , (A 7) can be written as follows

<sup>599</sup> 
$$(u_{ii} + u_{ij})^k + (u_{ji} + u_{jj})^k = \alpha \frac{1}{m_i^{k+1}} + \beta \frac{1}{m_j^{k+1}} + \gamma \left(\frac{1}{m_i} + \frac{1}{m_j}\right)^{k+1}$$
 (A 11)

where  $\alpha, \beta, \gamma$  are the number of rows where  $1/m_i$  is adjacent to 0, 0 is adjacent to  $1/m_j$ , and  $1/m_i$ is adjacent to  $1/m_j$  in columns *i* and *j* respectively. More precisely,  $\alpha$  is the cardinality of the set  $K_{ij}^i = \{n : u_{ni} = 1/m_i, u_{nj} = 0\}, \beta$  is the cardinality of the set  $K_{ij}^j = \{n : u_{ni} = 0, u_{nj} = 1/m_j\}$ and  $\gamma$  is the cardinality of the set  $K_{ij}^{ij} = \{n : u_{ni} = 1/m_i, u_{nj} = 1/m_j\}$ .

Since  $C_i = K_{ij}^i \cup K_{ij}^{ij}$  and  $C_j = K_{ij}^j \cup K_{ij}^{ij}$ , we have that  $m_i = \alpha + \gamma$  and  $m_j = \beta + \gamma$ . Since  $K_{ij}^i, K_{ij}^j, K_{ij}^{ij}$  are disjoint, we have  $\alpha + \beta + \gamma \leq N$ . Now, consider the different possibilities we con have on the left-hand side of equation (A 11).

#### Case 1: 608

 $u_{ii} = 1/m_i, \ u_{ij} = 0$  in row i and  $u_{ji} = 1/m_i, \ u_{jj} = 1/m_j$  in row j. Thus  $\alpha, \gamma \ge 1$  and therefore 609 equation (A 11) gives 610

$$_{611} \quad \frac{1}{m_i^k} + \left(\frac{m_i + m_j}{m_i m_j}\right)^k = \frac{\alpha}{m_i^{k+1}} + \frac{\beta}{m_j^{k+1}} + \gamma \left(\frac{m_i + m_j}{m_i m_j}\right)^{k+1} \Rightarrow$$

$$\frac{1}{(\alpha+\gamma)^{k}} + \left(\frac{\alpha+\beta+2\gamma}{(\alpha+\gamma)(\beta+\gamma)}\right)^{k} = \frac{\alpha}{(\alpha+\gamma)^{k+1}} + \frac{\beta}{(\beta+\gamma)^{k+1}} + \gamma \left(\frac{\alpha+\beta+2\gamma}{(\alpha+\gamma)(\beta+\gamma)}\right)^{k+1} \Rightarrow$$

$$^{613} \quad \frac{(\beta+\gamma)^{\kappa}+(\alpha+\beta+2\gamma)^{\kappa}}{[(\alpha+\gamma)(\beta+\gamma)]^{k}} = \frac{\alpha(\beta+\gamma)^{\kappa+1}+\beta(\alpha+\gamma)^{\kappa+1}+\gamma(\alpha+\beta+2\gamma)}{[(\alpha+\gamma)(\beta+\gamma)]^{k+1}}$$

$$_{^{614}} \quad (\beta+\gamma)^k + (\alpha+\beta+2\gamma)^k = \frac{\alpha(\beta+\gamma)^{k+1} + \beta(\alpha+\gamma)^{k+1} + \gamma(\alpha+\beta+2\gamma)^{k+1}}{(\alpha+\gamma)(\beta+\gamma)} \Rightarrow$$

$$_{615} \quad (\beta+\gamma)^k + (\alpha+\beta+2\gamma)^k = \frac{\alpha(\beta+\gamma)^k}{\alpha+\gamma} + \frac{\beta(\alpha+\gamma)^k}{\beta+\gamma} + \frac{(\alpha\gamma+\beta\gamma+2\gamma^2)(\alpha+\beta+2\gamma)^k}{\alpha\beta+\alpha\gamma+\beta\gamma+\gamma^2} \Rightarrow$$

$$\underset{_{617}}{_{616}} \quad \frac{\gamma(\beta+\gamma)^k}{\alpha+\gamma} = \frac{\beta(\alpha+\gamma)^k}{\beta+\gamma} + \frac{(\gamma^2-\alpha\beta)(\alpha+\beta+2\gamma)^k}{\alpha\beta+\alpha\gamma+\beta\gamma+\gamma^2}.$$

As  $k \to \infty$ , we get  $(\beta + \gamma)^k \neq (\alpha + \gamma)^k \pm (\alpha + \beta + 2\gamma)^k$  since  $\alpha + \beta + 2\gamma > \beta + \gamma$ ,  $\alpha + \gamma$  hence we want  $\gamma^2 = \alpha\beta$  to get rid off  $(\alpha + \beta + 2\gamma)^k$ . This implies that  $\beta + \gamma = \alpha + \gamma \Rightarrow \alpha = \beta \Rightarrow \alpha = \beta$ 618 619  $\beta = \gamma$  giving  $m_i = m_j$ . 620

#### Case 2: 621

 $u_{ii} = 1/m_i$ ,  $u_{ij} = 1/m_j$  in row *i* and  $u_{ji} = 0$ ,  $u_{jj} = 1/m_j$  in row *j*. This case is symmetrical to 622 Case 1 and therefore we get that  $\alpha = \beta = \gamma$  giving  $m_i = m_j$ . 623

 $u_{ii} = 1/m_i$ ,  $u_{ij} = 1/m_j$  in row *i* and  $u_{ji} = 1/m_i$ ,  $u_{jj} = 1/m_j$  in row *j*. Thus  $\gamma \ge 2$  and therefore 625 equation (A 11) gives 626

$$^{627} \qquad 2\left(\frac{m_i+m_j}{m_im_j}\right)^k = \frac{\alpha}{m_i^{k+1}} + \frac{\beta}{m_j^{k+1}} + \gamma\left(\frac{m_i+m_j}{m_im_j}\right)^{k+1} \Rightarrow$$

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$$2\left(\frac{\alpha+\beta+2\gamma}{(\alpha+\gamma)(\beta+\gamma)}\right)^{k} = \frac{\alpha(\beta+\gamma)^{k+1} + \beta(\alpha+\gamma)^{k+1} + \gamma(\alpha+\beta+2\gamma)^{k+1}}{[(\alpha+\gamma)(\beta+\gamma)]^{k+1}}$$

$$^{629} \qquad 2\left(\alpha+\beta+2\gamma\right)^{k} = \frac{\alpha(\beta+\gamma)^{k+1}+\beta(\alpha+\gamma)^{k+1}+\gamma(\alpha+\beta+2\gamma)^{k+1}}{(\alpha+\gamma)(\beta+\gamma)} \Rightarrow$$

$$2\left(\alpha+\beta+2\gamma\right)^{k} = \frac{\alpha(\beta+\gamma)^{k}}{\alpha+\gamma} + \frac{\beta(\alpha+\gamma)^{k}}{\beta+\gamma} + \frac{(\alpha\gamma+\beta\gamma+2\gamma^{2})(\alpha+\beta+2\gamma)^{k}}{\alpha\beta+\alpha\gamma+\beta\gamma+\gamma^{2}} \Rightarrow$$

$$\underset{\text{\tiny 631}}{\text{\tiny 631}} \qquad \qquad \frac{(2\alpha\beta + \alpha\gamma + \beta\gamma)(\alpha + \beta + 2\gamma)^k}{\alpha\beta + \alpha\gamma + \beta\gamma + \gamma^2} = \frac{\alpha(\beta + \gamma)^k}{\alpha + \gamma} + \frac{\beta(\alpha + \gamma)^k}{\beta + \gamma}.$$

As  $k \to \infty$ , we get  $(\alpha + \beta + 2\gamma)^k \neq (\beta + \gamma)^k + (\alpha + \gamma)^k$  since  $\alpha + \beta + 2\gamma > \beta + \gamma$ ,  $\alpha + \gamma$  hence 633 we want  $2\alpha\beta + \alpha\gamma + \beta\gamma = 0 \Rightarrow \alpha, \beta = 0$  giving  $m_i = m_i$ . 634

 $u_{ii} = 1/m_i, \ u_{ij} = 0$  in row i and  $u_{ji} = 0, \ u_{jj} = 1/m_j$  in row j. Thus  $\alpha, \beta \ge 1$  and therefore 636 equation (A 11) gives 637

 $\frac{1}{(\alpha+\gamma)^k} + \frac{1}{(\beta+\gamma)^k} = \frac{\alpha}{(\alpha+\gamma)^{k+1}} + \frac{\beta}{(\beta+\gamma)^{k+1}} + \gamma \left(\frac{\gamma+\beta+2\gamma}{(\alpha+\gamma)(\beta+\gamma)}\right)^{k+1} \Rightarrow$ 

$$_{^{638}} \qquad \qquad 1/m_i^k + 1/m_j^k = \frac{\alpha}{m_i^{k+1}} + \frac{\beta}{m_j^{k+1}} + \gamma \left(\frac{m_i + m_j}{m_i m_j}\right)^{k+1} \Rightarrow$$

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$$\frac{(\beta+\gamma)^k+(\alpha+\gamma)^k}{[(\alpha+\gamma)(\beta+\gamma)]^k}=\frac{\alpha(\beta+\gamma)^{k+1}+\beta(\alpha+\gamma)^{k+1}+\gamma(\alpha+\beta+2\gamma)^{k+1}}{[(\alpha+\gamma)(\beta+\gamma)]^{k+1}}\Rightarrow$$

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$$(\beta+\gamma)^k + (\alpha+\gamma)^k = \frac{\alpha(\beta+\gamma)^{k+1} + \beta(\alpha+\gamma)^{k+1} + \gamma(\alpha+\beta+2\gamma)^{k+1}}{(\alpha+\gamma)(\beta+\gamma)} \Rightarrow$$

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$$(\beta+\gamma)^k + (\alpha+\gamma)^k = \frac{\alpha(\beta+\gamma)^k}{\alpha+\gamma} + \frac{\beta(\alpha+\gamma)^k}{\beta+\gamma} + \frac{\gamma(\alpha+\beta+2\gamma)^{k+1}}{\alpha\beta+\alpha\gamma+\beta\gamma+\gamma^2}.$$

As  $k \to \infty$ , we get  $0 \neq (\alpha + \beta + 2\gamma)^k$  since  $\alpha, \beta \ge 1$  hence we require that  $\gamma = 0$  to get an equality.

#### 645 Conclusion from all the cases above

We see that  $m_i \neq m_j$  is potentially possible only in Case 4. However, **U** is strongly connected. If one connects *i* and *j* by a path  $i = i_0, i_1, i_2, \dots i_n = j$ , then one has  $m_{i_k} = m_{i_{k+1}}$  as  $i_k$  and  $i_{k+1}$ must fall into Case 1, Case 2 or Case 3 above. Thus  $m_i = m_j$ . This implies that every column of **U** has  $2 \le m \le N$  nonzero elements, including  $u_{nn}$ , that are all equal to 1/m. This is also true for every row of **U** because it is right stochastic by definition.

### Step 5: There exists state S such that $\mathcal{C}_a = \mathcal{C}_{a'}$ for all $a, a' \in S$

We can define the state  $\mathcal{R}_x = \{n : u_{xn} = u_{xx}\}$  then, by definition,  $x \in \mathcal{R}_x$  and  $|\mathcal{R}_x| = m$  since there are *m* nonzero elements in row *x* of **U**. Consider the state  $S = \mathcal{R}_x \setminus \{y\}$  for  $y \in \mathcal{R}_x \setminus \{x\}$ . For this *S* (as well as any other state), we have that

$$\begin{array}{c} \text{if } n \in S \text{ then } 1/m \\ \text{if } n \notin S \text{ then } 0 \end{array} \right\} \leq U(n,S) \leq \frac{\min(m,|S|)}{m}.$$

<sup>657</sup> We can therefore write equation (A 7) in the form

$$\sum_{i=1}^{\min(m,|S|)} \lambda_S(i) \left(\frac{i}{m}\right)^k = \sum_{i=0}^{\min(m,|S|)} \lambda'_S(i) \left(\frac{i}{m}\right)^{k+1}$$
(A 12)

where  $\lambda_S(i)$  is the number of U(n, S) terms equal to i/m for  $n \in S$  and  $\lambda'_S(i)$  is the number of U(n, S) terms equal to i/m for  $n \in \mathcal{N}$ , which means that  $\lambda'_S(i) \ge \lambda_S(i)$  for  $i \neq 0$ . The ratio of the left-hand side and right-hand side of equation (A 12) should always be equal to one. Therefore, as  $k \to \infty$ , we require that

$$\lambda_S(i_{\max}) = \lambda_S'(i_{\max}) \frac{i_{\max}}{m}$$

where  $i_{\max}$  is the largest i such that  $\lambda_S(i) > 0$ .

We have that  $i_{\max} = m - 1$  in equation (A 12) because |S| = m - 1 so U(x, S) = (m - 1)/m. This means that for state S, as  $k \to \infty$ , we require that

$$\lambda_{S}(m-1) = \lambda'_{S}(m-1) \frac{m-1}{m}$$

<sup>671</sup> Since  $\lambda_S(m-1)$  is an integer,  $\lambda'_S(m-1)$  has to be a multiple of m and the only possible value <sup>672</sup> that satisfies this criteria is  $\lambda'_S(m-1) = m$  hence  $\lambda_S(m-1) = m - 1$ .

Since 
$$\lambda'_S(m-1) = m$$
 there exist  $m$  rows  $j_1, j_2, \dots, j_m$  such that  $U(j_n, S) = (m-1)/m$ , that is,  
 $u_{j_n a} = 1/m \ \forall a \in S$ . This means that  $C_a = \{j_1, j_2, \dots, j_m\} \ \forall a \in S$  hence  $C_a = C_{a'}$  for all  $a, a' \in S$ .

### <sup>675</sup> Step 6: m = 2 or m = N

By contradiction, assume that 2 < m < N. We can consider another state  $S' = \mathcal{R}_x \setminus \{z\}$  such that  $z \in \mathcal{R}_x \setminus \{x, y\}$ . We then have that  $i_{\max} = m - 1$  in equation (A 12) because |S'| = m - 1so U(x, S') = (m - 1)/m. As before, this means that  $C_a = C_{a'}$  for all  $a, a' \in S'$ . Since  $x \in S, S'$ and  $\mathcal{R}_x = S \cup S'$  we have that  $C_a = C_{a'}$  for all  $a, a' \in \mathcal{R}_x$ . For 2 < m < N this implies that vertices  $i \in \mathcal{R}_x$  are disconnected from  $j \in \mathcal{N} \setminus \mathcal{R}_x$  and we therefore have disconnected graphs, a contradiction.

	Su	mmary of Notation
Symbol	Definition	Description
N	$\in \mathbb{Z}^+ \setminus \{0,1\}$	Population size.
A, B		The two types of individuals in population.
$I_n$		Individual <i>n</i> .
S	$= \{n : I_n \text{ of type } A\}$	State of the population.
$\mathcal{N}$	$=\{1, 2, \dots, N\}$	State in which all $I_n$ of type A.
r	$\in (0,\infty)$	Fitness of a type <i>A</i> individual.
	$\in \{1, r\}$	Fitness of $I_n$ in state S.
D	=(V,E)	Replacement digraph with vertices $V$ where $ V  = N$ and directed edges $E$ .
$w_{ij}$	$\in [0,\infty)$	Edge weight such that $w_{ij} > 0$ if and only if $(i, j) \in E$ .
$\mathbf{W}^{ij}$	$=(w_{ij})$	<i>Replacement matrix:</i> $N \times N$ weighted adjacency matrix of tuple $(D, w)$ .
$T_n^+$	$=\sum_{j=1}^{N} w_{nj}$	<i>Out temperature:</i> Sum of all outgoing edge weights of vertex $n \in V$ .
$T_n^-$	$=\sum_{i=1}^{N} w_{in}$	<i>In temperature</i> : Sum of all incoming edge weights of vertex $n \in V$ .
$b_i$	$\in [0,1]$	Probability $I_i$ chosen for birth.
	$\in [0,1]$	Probability a copy of $I_i$ replaces $I_j$ given $I_i$ was chosen for birth, i.e. replacement by death.
$d_i$	$\in [0,1]$	Probability $I_j$ chosen for death.
$b_{ij}$	$\in [0, 1]$	Probability a copy of $I_i$ replaces $I_j$ given $I_j$ is chosen for death, i.e. replacement by birth.
<b>r</b>	$\in [0,1]$	Probability a copy of $I_i$ replaces $I_j$ .
	$\in [0, 1]$ $\in [0, 1]$	State transition probability.
S S	$= (P_{SS'})$	State transition matrix.
$\mathcal{E}_{*,\mathbf{W},r}$	- (1 55')	Stochastic process with state transition matrix $S$ such that * dynamics are used on graph $W$ and type $A$ individuals have fitness $r$ .
$ ho_S^A$	$\in [0,1]$	Fixation probability of type <i>A</i> individual given initial state <i>S</i> .
W		Set of all strongly connected replacement matrices.
	$\{\mathbf{W}: T_n^+ = T_n^- \forall n\}$	Replacement matrices that are circulations.
	$\{\mathbf{W}: T_n^+ = T_j^- \ \forall i, j\}$	Replacement matrices that are isothermal.
	$\{\mathbf{W}: T_i^+ = I_j^+ \forall i, j\}$ $\{\mathbf{W}: T_n^+ = 1 \;\forall n\}$	Right stochastic replacement matrices.
$W_R W_L$	$\{\mathbf{W}: T_n = 1 \forall n\} \\ \{\mathbf{W}: T_n^- = 1 \forall n\}$	Left stochastic replacement matrices.
$C_N$	$\{\mathbf{v}\mathbf{v}:I_n=1\ \forall h\}$	Replacement matrices whose digraphs are cycles of length N.
$f_R$	$(w_{ij}) \mapsto (w_{ij} / \sum_n w_{in})$	Map from $W$ to $W_R$ .
$f_L$	$(w_{ij}) \mapsto (w_{ij} / \sum_n w_{nj})$	Map from W to $W_L$ .
f'	$(w_{ij}) \mapsto (w_{ij} / \sum_{n,k} w_{nk})$	Map from $W$ to $W_L$ .
$M_*$	$(-ij) \sim (\omega ij) \bigtriangleup n, k \ \omega nk)$	Replacement matrices for which $\mathcal{E}_*$ is $\rho$ -equivalent to a
111*		Moran process when $*$ dynamics are used.

Summary of Notation

Table 1: Notation used in this paper.

Process	$\mathbb{P}(I_i \text{ replaces } I_j)$	Order chosen	$\mathbb{P}(\text{Chosen first})$	$\mathbb{P}(\text{Chosen second})$	Definition of <i>M</i> *	Illustration of $M_*$
BDB [11, 13, 14, 16, 19, 33, 36]	$\mathfrak{r}_{ij}=b_id_{ij}$	$I_i$ then $I_j$	$b_i = \frac{F_i(S)}{\sum_n F_n(S)}$	$d_{ij} = \frac{w_{ij}}{\sum_{n} w_{in}}$	$M_{\text{BDB}} = \{ \mathbf{W} : f_R(\mathbf{W}) \in W_C \}$ = $f_R^{-1}(W_C)$ $M_{\text{BDD}} = \{ \mathbf{W} : f_R(\mathbf{W}) \in \{ \mathbf{W}_{\mathbf{H}} \} \cup C_N \}$ = $f_R^{-1}(\{ \mathbf{W}_{\mathbf{H}} \} \cup C_N)$	
BDD [27]	$\mathfrak{r}_{ij}=b_id_{ij}$	$I_i$ then $I_j$	$b_i = \frac{1}{N}$	$d_{ij} = \frac{w_{ij}/F_j(S)}{\sum_n w_{in}/F_n(S)}$	$M_{\text{BDD}} = \{ \mathbf{W} : f_R(\mathbf{W}) \in \{ \mathbf{W}_{\mathbf{H}} \} \cup C_N \}$ $= f_R^{-1}(\{ \mathbf{W}_{\mathbf{H}} \} \cup C_N)$	
DBD [12, 13, 36, 37]	$\mathfrak{r}_{ij}=d_ib_{ij}$				$M_{\text{DBD}} = \{ \mathbf{W} : f_L(\mathbf{W}) \in W_C \}$ $= f_L^{-1}(W_C)$	
DBB [16, 33, 38, 39, 40]	$\mathfrak{r}_{ij} = d_i b_{ij}$				$M_{\text{DBB}} = \{ \mathbf{W} : f_L(\mathbf{W}) \in \{ \mathbf{W}_{\mathbf{H}} \} \cup C_N \}$ $= f_L^{-1}(\{ \mathbf{W}_{\mathbf{H}} \} \cup C_N)$	
LB [11, 13, 36]	$\mathfrak{r}_{ij} = \frac{w_{ij}F_i(S)}{\sum_{r,h} w_{nk}F_n(S)}$	Simultaneous	N/A	n N/A	$M_{\text{LB}} = \{ \mathbf{W} : f'(\mathbf{W}) \in W_C \}$ $= f'^{-1}(W_C) = W_C$	
LD [23]	$\mathfrak{r}_{ij} = \frac{w_{ij}/F_j(S)}{\sum_{n,k} w_{nk}/F_k(S)}$	Simultaneous	N/A	N/A	$M_{\text{LD}} = \{ \mathbf{W} : f'(\mathbf{W}) \in W_C \}$ $= f'^{-1}(W_C) = W_C$	

Key for Illustration of  $M_*$ :

W		$W_1 = W_I \cap f_R^{-1}(\{\mathbf{W}_{\mathbf{H}}\} \cup C_N)$
	$W_8$	$= W_I \cap f_{L_*}^{-1}(\{\mathbf{W}_{\mathbf{H}}\} \cup C_N)$
		$W_2 = W_I \setminus f_R^{-1}(\{\mathbf{W}_{\mathbf{H}}\} \cup C_N)$
	$ W_6 W_1 W_4 $	$= W_I \setminus f_L^{-1}(\{\mathbf{W}_{\mathbf{H}}\} \cup C_N)$
		$W_3 = W_C \setminus W_I$
	$ W_7 W_2 W_5 $	$W_4 = \left(f_R^{-1}(W_C) \setminus W_C\right) \cap f_R^{-1}(\{\mathbf{W}_{\mathbf{H}}\} \cup C_N)$
	117	$W_5 = \left(f_R^{-1}(W_C) \setminus W_C\right) \setminus f_R^{-1}(\{\mathbf{W}_{\mathbf{H}}\} \cup C_N)$
	$W_3$	$W_6 = \left(f_L^{-1}(W_C) \setminus W_C\right) \cap f_L^{-1}(\{\mathbf{W}_{\mathbf{H}}\} \cup C_N)$
		$W_7 = \left( f_L^{-1}(W_C) \setminus W_C \right) \setminus f_L^{-1}(\{\mathbf{W}_{\mathbf{H}}\} \cup C_N)$
		$W_8 = W \setminus \bigcup_{i=1}^7 W_i$

The key on the left gives the definition of partitions  $W_1, W_2, \ldots, W_8$  of W. The partitions  $W_i$  that make up  $M_*$  are highlighted for each of the dynamics in the last column. The partition of W where  $\mathcal{E}$  is  $\rho$ -equivalent to a Moran process regardless of the standard dynamics being used is given by  $M_L \cap M_{BDB} \cap M_{BDD} \cap M_{DBD} \cap M_{DBB} \equiv M_{BDD} \cap M_{DBB} \equiv W_1$ .

