

Faster Monte-Carlo Algorithms for Fixation Probability of the Moran Process on Undirected Graphs

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Abstract

Evolutionary graph theory studies the evolutionary dynamics in a population structure given as a connected graph. Each node of the graph represents an individual of the population, and edges determine how offspring are placed. We consider the classical birth-death Moran process where there are two types of individuals, namely, the residents with fitness 1 and mutants with fitness r . The fitness indicates the reproductive strength. The evolutionary dynamics happens as follows: in the initial step, in a population of all resident individuals a mutant is introduced, and then at each step, an individual is chosen proportional to the fitness of its type to reproduce, and the offspring replaces a neighbor uniformly at random. The process stops when all individuals are either residents or mutants. The probability that all individuals in the end are mutants is called the fixation probability, which is a key factor in the rate of evolution. We consider the problem of approximating the fixation probability.

The class of algorithms that is extremely relevant for approximation of the fixation probabilities is the Monte-Carlo simulation of the process. Previous results present a polynomial-time Monte-Carlo algorithm for undirected graphs when r is given in unary. First, we present a simple modification: instead of simulating each step, we discard *ineffective* steps, where no node changes type (i.e., either residents replace residents, or mutants replace mutants). Using the above simple modification and our result that the number of effective steps is concentrated around the expected number of effective steps, we present faster polynomial-time Monte-Carlo algorithms for undirected graphs. Our algorithms are always at least a factor $O(n^2/\log n)$ faster as compared to the previous algorithms, where n is the number of nodes, and is polynomial even if r is given in binary. We also present lower bounds showing that the upper bound on the expected number of effective steps we present is asymptotically tight for undirected graphs.

1 Introduction

In this work we present faster Monte-Carlo algorithms for approximation of the fixation probability of the fundamental Moran process on population structures with symmetric interactions. We start with the description of the problem.

Evolutionary dynamics Evolutionary dynamics act on populations, where the composition of the population changes over time due to mutation and selection. Mutation generates new types and selection changes the relative abundance of different types. A fundamental concept in evolutionary dynamics is the fixation probability of a new mutant [5, 10, 12, 13]: Consider a population of n *resident* individuals, each with a fitness value 1. A single *mutant* with non-negative fitness value r is introduced in the population as the initialization step. Intuitively, the fitness represents the reproductive strength. In the classical Moran process the following *birth-death* stochastic steps are repeated: At each time step, one individual is chosen at random proportional to the fitness to reproduce and one other individual is chosen uniformly at random for death. The offspring of the reproduced individual replaces the dead individual. This stochastic process continues until either all individuals are mutants or all individuals are residents. The *fixation probability* is the probability that the mutants take over the population, which means all individuals are mutants. A standard calculation shows that the fixation probability is given by $(1 - (1/r)) / (1 - (1/r^n))$. The correlation between the relative fitness r of the mutant and the fixation probability is a measure of the effect of natural selection. The rate of evolution, which is the rate at which subsequent mutations accumulate in the population, is proportional to the fixation probability, the mutation rate, and the population size n . Hence fixation probability is a fundamental concept in evolution.

Evolutionary graph theory While the basic Moran process happens in well-mixed population (all individuals interact uniformly with all others), a fundamental extension is to study the process on population structures. Evolutionary graph theory studies this phenomenon. The individuals of the population occupy the nodes of a connected graph. The links (edges) determine who interacts with whom. Basically, in the birth-death step, for the death for replacement, a neighbor of the reproducing individual is chosen uniformly at random. Evolutionary graph theory describes evolutionary dynamics in spatially structured population where most interactions and competitions occur mainly among neighbors in physical space [11, 2, 6, 17]. Undirected graphs represent population structures where the interactions are symmetric, whereas directed graphs allow for asymmetric interactions. The fixation probability depends on the population structure [11, 1, 7, 3]. Thus, the fundamental computational problem in evolutionary graph theory is as follows: given a population structure (i.e., a graph), the relative fitness r , and $\epsilon > 0$, compute an ϵ -approximation of the fixation probability.

| | All steps | Effective steps |
|-----------------------|--|---|
| #steps in expectation | $O(n^2 \Delta^2)$ | $O(n\Delta)$ |
| Concentration bounds | $\Pr[\tau \geq \frac{n^2 \Delta^2 r x}{r-1}] \leq 1/x$ | $\Pr[\tau \geq \frac{6n\Delta x}{\min(r-1, 1)}] \leq 2^{-x}$ |
| Sampling a step | $O(1)$ | $O(\Delta)$ |
| Fixation algo | $O(n^6 \Delta^2 \epsilon^{-4})$ | $O(n^2 \Delta^2 \epsilon^{-2} (\log n + \log \epsilon^{-1}))$ |

Table 1: Comparison with previous work, for constant $r > 1$. We denote by n , Δ , τ , and ϵ , the number of nodes, the maximum degree, the random variable for the fixation time, and the approximation factor, respectively. The results in the column “All steps” is from [4], except that we present the dependency on Δ , which was considered as n in [4]. The results of the column “Effective steps” is the results of this paper

Monte-Carlo algorithms A particularly important class of algorithms for biologists is the Monte-Carlo algorithms, because it is simple and easy to interpret. The Monte-Carlo algorithm for the Moran process basically requires to simulate the process, and from the statistics obtain an approximation of the fixation probability. Hence, the basic question we address in this work is simple Monte-Carlo algorithms for approximating the fixation probability. It was shown in [16] that simple simulation can take exponential time on directed graphs and thus we focus on undirected graphs. The main previous algorithmic result in this area [4] presents a polynomial-time Monte-Carlo algorithm for undirected graphs when r is given in unary. The main result of [4] shows that for undirected graphs it suffices to run each simulation for polynomially many steps.

Our contributions In this work our main contributions are as follows:

1. *Faster algorithm for undirected graphs* First, we present a simple modification: instead of simulating each step, we discard *ineffective* steps, where no node changes type (i.e., either residents replace residents, or mutants replace mutants). We then show that the number of effective steps is concentrated around the expected number of effective steps. The sampling of each effective step is more complicated though than sampling of each step. We then present an efficient algorithm for sampling of the effective steps, which requires $O(m)$ preprocessing and then $O(\Delta)$ time for sampling, where m is the number of edges and Δ is the maximum degree. Combining all our results we obtain faster polynomial-time Monte-Carlo algorithms: Our algorithms are always at least a factor $n^2 / \log n$ times a constant (in most cases $n^3 / \log n$ times a constant) faster as compared to the previous algorithm, and is polynomial even if r is given in binary. We present a comparison in Table 1, for constant $r > 1$ (since the previous algorithm is not in polynomial time for r in binary). For a detailed comparison see Table 2 in the Appendix.

2. *Lower bounds* We also present lower bounds showing that the upper bound on the expected number of effective steps we present is asymptotically tight for undirected graphs.

Related complexity result While in this work we consider evolutionary graph theory, a related problem is evolutionary games on graphs (which studies the problem of frequency dependent selection). The approximation problem for evolutionary games on graphs is considerably harder (e.g., PSPACE-completeness results have been established) [9].

Technical contributions Note that for the problem we consider the goal is not to design complicated efficient algorithms, but simple algorithms that are efficient. By simple, we mean something that is related to the process itself, as biologists understand and interpret the Moran process well. Our main technical contribution is a simple idea to discard ineffective steps, which is intuitive, and we show that the simple modification leads to significantly faster algorithms. We show a gain of factor $O(n\Delta)$ due to the effective steps, then lose a factor of $O(\Delta)$ due to sampling, and our other improvements are due to better concentration results. We also present an interesting family of graphs for the lower bound examples. Technical proofs omitted due to lack of space are in the Appendix.

2 Moran process on graphs

Connected graph and type function We consider the population structure represented as a connected graph. There is a connected *graph* $G = (V, E)$, of n nodes and m edges, and two types $T = \{t_1, t_2\}$. The two types represent residents and mutants, and in the technical exposition we refer to them as t_1 and t_2 for elegant notation. We say that a node v is a *successor* of a node u if $(u, v) \in E$. The graph is *undirected* if for all $(u, v) \in E$ we also have $(v, u) \in E$, otherwise it is *directed*. There is a *type function* f mapping each node v to a type $t \in T$. Each type t is in turn associated with a positive integer $w(t)$, the type's fitness denoting the corresponding reproductive strength. Without loss of generality, we will assume that $r = w(t_1) \geq w(t_2) = 1$, for some number r (the process we consider does not change under scaling, and r denotes relative fitness). Let $W(f) = \sum_{u \in V} w(f(u))$ be the total fitness. For a node v let $\deg v$ be the degree of v in G . Also, let $\Delta = \max_{v \in V} \deg v$ be the maximum degree of a node. For a type t and type function f , let $V_{t,f}$ be the nodes mapped to t by f . Given a type t and a node v , let $f[v \rightarrow t]$ denote the following function: $f[v \rightarrow t](u) = t$ if $u = v$ and $f(u)$ otherwise.

Moran process on graphs We consider the following classical Moran birth-death process where a *dynamic evolution step* of the process changes a type function from f to f' as follows:

1. First a node v is picked at random with probability proportional to $w(f(v))$, i.e. each node v has probability of being picked equal to $\frac{w(f(v))}{W(f)}$.
2. Next, a successor u of v is picked uniformly at random.
3. The type of u is then changed to $f(v)$. In other words, $f' = f[u \rightarrow f(v)]$.

Fixation A type t *fixates* in a type function f if f maps all nodes to t . Given a type function f , repeated applications of the dynamic evolution step generate a sequence of type functions $f = f_1, f_2, \dots, f_\infty$. Note that if a type has fixated (for some type t) in f_i then it has also fixated in f_j for $i < j$. We say that a process has *fixation time* i if f_i has fixated but f_{i-1} has not. We say that an initial type function f has fixation probability p for a type t , if the probability that t eventually fixates (over the probability measure on sequences generated by repeated applications of the dynamic evolution step f)

Basic questions We consider the following basic questions:

1. *Fixation problem* Given a type t , what is the fixation probability of t averaged over the n initial type functions with a single node mapping to t ?
2. *Extinction problem* Given a type t , what is the fixation probability of t averaged over the n initial type functions with a single node *not* mapping to t ?
3. *Generalized fixation problem* Given a graph, a type t and an type function f what is the fixation probability of t in G , when the initial type function is f ?

Remark 1. Note that in the *neutral* case when $r = 1$, the fixation problem has answer $1/n$ and extinction problem has answer $1 - 1/n$. Hence, in the rest of the paper we will consider $r > 1$. Also, to keep the presentation focused, in the main article, we will consider fixation and extinction of type t_1 . In the Appendix we also present another algorithm for the extinction of t_2 .

Results We will focus on undirected graphs. For undirected graphs, we will give new FPRAS (fully polynomial, randomized approximation scheme) for the fixation and the extinction problem, and a polynomial-time algorithm for an additive approximation of the generalized fixation problem. There exists previous FPRAS for the fixation and extinction problems [4]. Our upper bounds are at least a factor of $O(\frac{n^2}{\log n})$ (most cases $O(\frac{n^3}{\log n})$) better and always in $\text{Poly}(n, 1/\epsilon)$, whereas the previous algorithms are not in polynomial time for r given in binary.

3 Discarding ineffective steps

We consider undirected graphs. Previous work by Diaz et al. [4] showed that the expected number of dynamic evolution steps till fixation is polynomial, and then used it to give a polynomial-time Monte-Carlo algorithm. Our goal is to improve the quite high polynomial-time complexity, while giving a Monte-Carlo algorithm. To achieve this we define the notion of effective steps.

Effective steps A dynamic evolution step, which changes the type function from f to f' , is *effective* if $f \neq f'$ (and *ineffective* otherwise). The idea is that steps in which no node changes type (because the two nodes selected in the dynamic evolution step already had the same type) can be discarded, without changing which type fixates/gets eliminated.

Two challenges The two challenges are as follows:

1. *Number of steps* The first challenge is to establish that the expected number of effective steps is asymptotically smaller than the expected number of all steps. We will establish a factor $O(n\Delta)$ improvement (recall Δ is the maximum degree).
2. *Sampling* Sampling an effective step is harder than sampling a normal step. Thus it is not clear that considering effective steps leads to a faster algorithm. We consider the problem of efficiently sampling an effective step in a later section, see Section 5. We show that sampling an effective step can be done in $O(\Delta)$ time (after $O(m)$ preprocessing).

Notation For a type function f , let $\Gamma_v(f)$ be the subset of successors of v , such that $u \in \Gamma_v(f)$ iff $f(v) \neq f(u)$. Also, let $W'(f) = \sum_u w(f(u)) \cdot \frac{|\Gamma_u(f)|}{\deg u}$.

Modified dynamic evolution step Formally, we consider the following *modified dynamic evolution step* (that changes the type function from f to f' and assumes that f does not map all nodes to the same type):

1. First a node v is picked at random with probability proportional to $p(v) = w(f(v)) \cdot \frac{|\Gamma_v(f)|}{\deg v}$ i.e. each node v has probability of being picked equal to $\frac{p(v)}{W'(f)}$.
2. Next, a successor u of v is picked uniformly at random among $\Gamma_v(f)$.
3. The type of u is then changed to $f(v)$, i.e., $f' = f[u \rightarrow f(v)]$.

In the following lemma we show that the modified dynamic evolution step corresponds to the dynamic evolution step except for discarding steps in which no change was made.

Lemma 1. Fix any type function f such that neither type has fixated. Let f_d (resp., f_m) be the next type function under dynamic evolution step (resp., modified dynamic evolution step). Then, $\Pr[f \neq f_d] > 0$ and for all type functions f' we have: $\Pr[f' = f_d \mid f \neq f_d] = \Pr[f' = f_m]$.

Potential function ψ Similar to [4] we consider the *potential function* $\psi = \sum_{v \in V_{t_1, f}} \frac{1}{\deg v}$ (recall that $V_{t_1, f}$ is the set of nodes of type t_1). We now lower bound the expected difference in potential per modified evolutionary step.

Lemma 2. Let f be a type function such that neither type has fixated. Apply a modified dynamic evolution step on f to obtain f' . Then,

$$\mathbb{E}[\psi(f') - \psi(f)] \geq \frac{r-1}{\Delta \cdot (r+1)} .$$

Proof. Observe that f differs from f' for exactly one node u . More precisely, let v be the node picked in line 1 of the modified dynamic evolution step and let u be the node picked in line 2. Then, $f' = f[u \rightarrow f(v)]$. The probability to select v is $\frac{p(v)}{W'(f)}$. The probability to then pick u is $\frac{1}{|\Gamma_v(f)|}$.

We have that

- If $f(u) = t_2$ (and thus, since it got picked $f(v) = t_1$), then $\psi(f') - \psi(f) = \frac{1}{\deg u}$.
- If $f(u) = t_1$ (and thus, since it got picked $f(v) = t_2$), then $\psi(f') - \psi(f) = -\frac{1}{\deg u}$.

Below we use the following notations:

$$E_{12} = \{(v, u) \in E \mid f(v) = t_1 \text{ and } f(u) = t_2\}; \quad E_{21} = \{(v, u) \in E \mid f(v) = t_2 \text{ and } f(u) = t_1\}.$$

Thus,

$$\begin{aligned} \mathbb{E}[\psi(f') - \psi(f)] &= \sum_{(v, u) \in E_{12}} \left(\frac{p(v)}{W'(f)} \cdot \frac{1}{|\Gamma_v(f)|} \cdot \frac{1}{\deg u} \right) - \sum_{(v, u) \in E_{21}} \left(\frac{p(v)}{W'(f)} \cdot \frac{1}{|\Gamma_v(f)|} \cdot \frac{1}{\deg u} \right) \\ &= \sum_{(v, u) \in E_{12}} \left(\frac{w(f(v))}{W'(f) \cdot (\deg u) \cdot (\deg v)} \right) - \sum_{(v, u) \in E_{21}} \left(\frac{w(f(v))}{W'(f) \cdot (\deg u) \cdot (\deg v)} \right) . \end{aligned}$$

Using that the graph is undirected we get,

$$\begin{aligned} \mathbb{E}[\psi(f) - \psi(f')] &= \sum_{(v, u) \in E_{12}} \left(\frac{w(f(v)) - w(f(u))}{W'(f) \cdot (\deg u) \cdot (\deg v)} \right) \\ &= \frac{1}{W'(f)} \sum_{(v, u) \in E_{12}} \left(\frac{r-1}{\min(\deg u, \deg v) \cdot \max(\deg u, \deg v)} \right) \\ &\geq \frac{r-1}{\Delta \cdot W'(f)} \sum_{(v, u) \in E_{12}} \frac{1}{\min(\deg u, \deg v)} = \frac{r-1}{\Delta \cdot W'(f)} \cdot S , \end{aligned}$$

where $S = \sum_{(v,u) \in E_{12}} \frac{1}{\min(\deg u, \deg v)}$. Note that in the second equality we use that for two numbers a, b , their product is equal to $\min(a, b) \cdot \max(a, b)$. By definition of $W'(f)$, we have

$$\begin{aligned} W'(f) &= \sum_u w(f(u)) \cdot \frac{|\Gamma_u(f)|}{\deg u} = \sum_u \sum_{v \in \Gamma_u(f)} \frac{w(f(u))}{\deg u} = \sum_{\substack{(v,u) \in E \\ f(u) \neq f(v)}} \frac{w(f(u))}{\deg u} \\ &= \sum_{(v,u) \in E_{12}} \left(\frac{w(f(u))}{\deg u} + \frac{w(f(v))}{\deg v} \right) \leq \sum_{(v,u) \in E_{12}} \frac{w(f(u)) + w(f(v))}{\min(\deg u, \deg v)} = (r+1) \cdot S . \end{aligned}$$

Thus, we see that $\mathbb{E}[\psi(f') - \psi(f)] \geq \frac{r-1}{\Delta \cdot (r+1)}$, as desired. This completes the proof. \square

Lemma 3. *Let $r = x\Delta$ for some number $x > 0$. Let f be a type function such that neither type has fixated. Apply a modified dynamic evolution step on f to obtain f' . The probability that $|V_{t_1, f'}| = |V_{t_1, f}| + 1$ is at least $\frac{x}{x+1}$ (otherwise, $|V_{t_1, f'}| = |V_{t_1, f}| - 1$).*

Proof. Consider any type function f . Let m' be the number of edges (u, v) , such that $f(u) \neq f(v)$. We will argue that the total weight of nodes of type t_1 , denoted W_1 , is at least xm' and that the total weight of nodes of type t_2 , denoted W_2 , is at most m' . We see that as follows:

$$W_1 = \sum_{v \in V_{t_1, f}} w(f(v)) \cdot \frac{|\Gamma_v(f)|}{\deg v} \geq \frac{x\Delta}{\Delta} \sum_{v \in V_{t_1, f}} |\Gamma_v(f)| = xm' ,$$

using that $\deg v \leq \Delta$ and $w(f(v)) = r = x\Delta$ in the inequality. Also,

$$W_2 = \sum_{v \in V_{t_2, f}} w(f(v)) \cdot \frac{|\Gamma_v(f)|}{\deg v} \leq \sum_{v \in V_{t_2, f}} |\Gamma_v(f)| = m' ,$$

using that $\deg v \geq 1$ and $w(f(v)) = 1$ in the inequality. We see that we thus have a probability of at least $\frac{x}{x+1}$ to pick a node of type t_1 . Because we are using effective steps, picking a member of type t will increment the number of that type (and decrement the number of the other type). \square

Lemma 4. *Consider an upper bound ℓ , for each starting type function, on the expected number of (effective) steps to fixation. Then for any starting type function the probability that fixation requires more than $2 \cdot \ell \cdot x$ (effective) steps is at most 2^{-x} .*

Proof. By Markov's inequality after $2 \cdot \ell$ (effective) steps the Moran process fixates with probability at least $\frac{1}{2}$, irrespective of the initial type function. We now split the steps into blocks of length $2 \cdot \ell$. In every block, by the preceding argument, there is a probability of at least $\frac{1}{2}$ to fixate in some step of that block, given that the process has not fixated before that block. Thus, for any integer $x \geq 1$, the probability to not fixate before the end of block x , which happens at step $2 \cdot \ell \cdot x$ is at most 2^{-x} . \square

We now present the main theorem of this section, which we obtain using the above lemmas, and techniques from [4].

Theorem 5. *Let t_1 and t_2 be the two types, such that $r = w(t_1) > w(t_2) = 1$. Let Δ be the maximum degree. Let k be the number of nodes of type t_2 in the initial type function. The following assertions hold:*

- Bounds dependent on r
 1. Expected steps *The process requires at most $3k\Delta / \min(r - 1, 1)$ effective steps in expectation, before fixation is reached.*
 2. Probability *For any integer $x \geq 1$, after $6xn\Delta / \min(r - 1, 1)$ effective steps, the probability that the process has not fixated is at most 2^{-x} , irrespective of the initial type function.*
- Bounds independent on r
 1. Expected steps *The process requires at most $2nk\Delta^2$ effective steps in expectation, before fixation is reached.*
 2. Probability *For any integer $x \geq 1$, after $4xn^2\Delta^2$ effective steps, the probability that the process has not fixated is at most 2^{-x} , irrespective of the initial type function.*
- Bounds for $r \geq 2\Delta$
 1. Expected steps *The process requires at most $3k$ effective steps in expectation, before fixation is reached.*
 2. Probability *For any integer $x \geq 1$, after $6xn$ effective steps, the probability that the process has not fixated is at most 2^{-x} , irrespective of the initial type function.*

4 Lower bound for undirected graphs

In this section, we will argue that our bound on the expected number of effective steps is essentially tight, for fixed r .

We construct our lower bound graph $G^{\Delta, n}$, for given Δ, n (sufficiently large), but fixed $r > 1$, as follows. We will argue that fixation of $G^{\Delta, n}$ takes $\Omega(k\Delta)$ effective steps, if there are initially exactly k members of type t_2 . For simplicity, we consider $\Delta > 2$ and $n > 4\Delta$ (it is easy to see using similar techniques that for lines, where $\Delta = 2$, the expected fixation time is $\Omega(k)$ - basically because t_1 is going to fixate with pr. $\approx 1 - 1/r$, using a proof like Lemma 6, and converting the k nodes of type t_2 takes at least k efficient steps). There are two parts to the graph: A line of $\approx n/2$ nodes and a stars-on-a-cycle graph of $\approx n/2$. There is 1 edge from the one of the stars in the stars-on-a-cycle graph to the line. More formally, the graph is as follows: Let $x := \lfloor n/(2\Delta - 2) \rfloor$. There are nodes $V_C = \{c_1, \dots, c_x\}$, such that c_i is connected to c_{i-1} and c_{i+1} for $1 < i < x$. Also, c_1 is connected to c_x . The nodes V_C are the centers of the

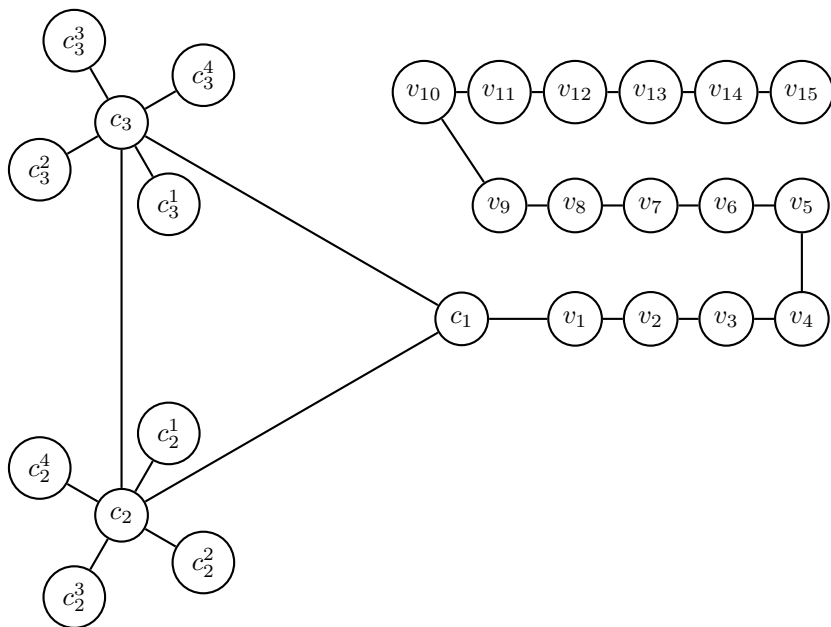


Figure 1: Example of a member of the family that attains the lower bound for undirected graphs. (Specifically, it is $G^{6,31}$)

stars in the stars-on-a-cycle graph. For each i , such that $2 \leq i \leq x$, the node c_i is connected to a set of leaves $V_C^i = \{c_i^1, \dots, c_i^{\Delta-2}\}$. The set $V_C \cup \bigcup_{i=2}^x V_C^i$ forms the stars-on-a-cycle graph. Note that c_1 is only connected to c_2 and c_x in the stars-on-a-cycle graph. We have that the stars-on-a-cycle graph consists of $s = (x-1) \cdot (\Delta-1) + 1 \approx n/2$ nodes. There are also nodes $V_L = \{\ell_1, \dots, \ell_{n-s}\}$, such that node ℓ_i is connected to ℓ_{i-1} and ℓ_{i+1} for $1 < i < n/2$. The nodes V_L forms the line and consists of $n-s \geq n/2$ nodes. The node c_1 is connected to ℓ_1 . There is an illustration of $G^{6,31}$ in Figure 1.

We first argue that if at least one of $V_L' = \{\ell_{\lceil n/4 \rceil}, \dots, \ell_{n-s}\}$ is initially of type t_1 , then with pr. lower bounded by a number depending only on r , type t_1 fixates (note that $|V_L'| \geq n/4$ and thus, even if there is only a single node of type t_1 initially placed uniformly at random, it is in V_L' with pr. $\geq 1/4$).

Lemma 6. *With pr. above $\frac{1-1/r}{2}$ if at least one of V_L' is initially of type t_1 , then t_1 fixates.*

The proof is based on applying the gambler's ruin twice. Once to find out that the pr. that V_L eventually becomes all t_1 is above $\frac{1-1/r}{2}$ (it is nearly $1-1/r$ in fact) and once to find out that if V_L is at some point all t_1 , then the pr. that t_2 fixates is exponentially small with base r and exponent $n-s$. See the appendix for the proof.

Whenever a node of V_C^i , for some i , changes type, we say that a *leaf-step*

occurred. We will next consider the pr. that an effective step is a leaf-step.

Lemma 7. *The pr. that an effective step is a leaf-step is at most $\frac{r}{\Delta}$.*

The proof is quite direct and considers that the probability that a leaf gets selected for reproduction over a center node in the stars-on-a-cycle graph. See the appendix for the proof.

We are now ready for the theorem.

Theorem 8. *Let $r > 1$ be some fixed constant. Consider $\Delta > 2$ (the maximum degree of the graph), $n > 4\Delta$ (sufficiently big), and some k such that $0 < k < n$. Then, if there are initially k members of type t_2 placed uniformly at random, the expected fixation time of $G^{\Delta, n}$ is above $\frac{k\Delta(1-1/r)}{32r}$ effective steps.*

Proof. Even if $k = n - 1$, we have that with pr. at least $\frac{1}{4}$, the lone node of type t_1 is initially in V'_L . If so, by Lemma 6, type t_1 is going to fixate with pr. at least $\frac{1-1/r}{2}$. Note that even for $\Delta = 3$, at least $\frac{n}{4}$ nodes of the graphs are in $V' := \bigcup_{i=2}^x V_C^i$ (i.e. the leaves of the stars-on-a-cycle graph). In expectation $\frac{k}{4}$ nodes of V' are thus initially of type t_2 . For fixation for t_1 to occur, we must thus make that many leaf-steps. Any effective step is a leaf-step with pr. at most $\frac{r}{\Delta}$ by Lemma 7. Hence, with pr. $\frac{1}{4} \cdot \frac{1-1/r}{2}$ ($\frac{1}{4}$ is the probability that at least one node of type t_1 is in V'_L and $\frac{1-1/r}{2}$ is a lower bound on the fixation probability if a node of V'_L is of type t_1) we must make $\frac{k\Delta}{4r}$ effective steps before fixation in expectation, implying that the expected fixation time is at least $\frac{k\Delta(1-1/r)}{32r}$ effective steps. \square

5 Sampling an effective step

In this section, we consider the problem of sampling an effective step. It is quite straightforward to do so in $O(m)$ time. We will present a data-structure that after $O(m)$ preprocessing can sample and update the distribution in $O(\Delta)$ time. For this result we assume that a uniformly random number can be selected between 0 and x for any number $x \leq n \cdot w(t)$ in constant time, a model that was also implicitly assumed in previous works [4]¹.

Remark 2. If we consider a weaker model, that requires constant time for each random bit, then we need $O(\log n)$ random bits in expectation and additional $O(\Delta)$ amortized time, using a similar data-structure (i.e., a total of $O(\Delta + \log n)$ amortized time in expectation). The argument for the weaker model is presented in the Appendix. In this more restrictive model [4] would use $O(\log n)$ time per step for sampling.

¹The construction of [4] was to store a list for t_1 and a list for t_2 and then first decide if a t_1 or t_2 node would be selected in this step (based on r and the number of nodes of the different types) and then pick a random such node. This works when all nodes of a type has the same weight but does not generalize to the case when each node can have a distinct weight based on the nodes successors like here

Sketch of data-structure We first sketch a list data-structure that supports (1) inserting elements; (2) removing elements; and (3) finding a random element; such that each operation takes (amortized or expected) $O(1)$ time. The idea based on dynamic arrays is as follows:

1. **Insertion** Inserting elements takes $O(1)$ amortized time in a dynamic array, using the standard construction.
2. **Deletion** Deleting elements is handled by changing the corresponding element to a null-value and then rebuilding the array, without the null-values, if more than half the elements have been deleted since the last rebuild. Again, this takes $O(1)$ amortized time.
3. **Find random element** Repeatedly pick a uniformly random entry. If it is not null, then output it. Since the array is at least half full, this takes in expectation at most 2 attempts and thus expected $O(1)$ time.

At all times we keep a doubly linked list of empty slots, to find a slot for insertion in $O(1)$ time.

Data-structure The idea is then as follows. We have 2Δ such list data-structures, one for each pair of type and degree. We also have a weight associated to each list, which is the sum of the weight of all nodes in the list, according to the modified dynamic evolution step. When the current type function is f , we represent each node v as follows: The corresponding list data-structure contains $|\Gamma_v(f)|$ copies of v (and v keeps track of the locations in a doubly linked list). Each node v also keeps track of $\Gamma_v(f)$, using another list data-structure. It is easy to construct the initial data-structure in $O(m)$ time (note: $\sum_v |\Gamma_v(f)| \leq 2m$).

Updating the data-structure We can then update the data-structure when the current type function f changes to $f[u \rightarrow t]$ (all updates have that form for some t and u), by removing u from the list data-structure $(f(u), \deg u)$ containing it and adding it to $(t, \deg u)$. Note that if we removed x' copies of u from $(f(u), \deg u)$ we add $\deg u - x'$ to $(t, \deg u)$. Also, we update each neighbor v of u (by deleting or adding a copy to $(f(v), \deg v)$, depending on whether $f(v) = t$). We also keep the weight corresponding to each list updated and $\Gamma_v(f)$ for all nodes v . This takes at most 4Δ data-structure insertions or deletions, and thus $O(\Delta)$ amortized time in total.

Sampling an effective step Let f be the current type function. First, pick a random list L among the 2Δ lists, proportional to their weight. Then pick a random node v from L . Then pick a node at random in $\Gamma_v(f)$. This takes $O(\Delta)$ time in expectation.

Remark 3. Observe that picking a random list among the 2Δ lists, proportional to their weight takes $O(\Delta)$ time to do naively: E.g. consider some ordering of

the lists and let w_i be the total weight of list i (we keep this updated so it can be found in constant time). Pick a random number x between 1 and the total weight of all the lists (assumed to be doable in constant time). Iterate over the lists in order and when looking at list i , check if $x < \sum_{j=1}^i w_j$. If so, pick list i , otherwise continue to list $i + 1$. By making a binary, balanced tree over the lists (similar to what is used for the more restrictive model, see the Appendix), the time can be brought down to $O(\log \Delta)$ for this step - however the naive approach suffices for our application, because updates requires $O(\Delta)$ time.

This leads to the following theorem.

Theorem 9. *An effective step can be sampled in (amortized and expected) $O(\Delta)$ time after $O(m)$ preprocessing, if a uniformly random integer between 0 and x , for any $0 < x \leq n \cdot w(t)$, can be found in constant time.*

6 Algorithms for approximating fixation probability

We present the algorithms for solving the fixation, extinction, and generalized fixation problems.

The Meta-simulation algorithm Similar to [4], the algorithms are instantiating the following meta-simulation algorithm, that takes a distribution over initial type functions \mathcal{D} , type t and natural numbers u and z as input:

Function MetaSimulation(t, z, u, \mathcal{D})

```

Let  $y \leftarrow 0$ ;
for ( $i \in \{1, \dots, z\}$ ) do
    Initialize a new simulation  $I$  with initial type function  $f$  picked
    according to  $\mathcal{D}$ ;
    Let  $j \leftarrow 0$ ;
    while ( $I$  has not fixated) do
        if ( $j \geq u$ ) then
            return Simulation took too long;
        Set  $j \leftarrow j + 1$ ;
        Simulate an effective step in  $I$ ;
    if ( $t$  fixated in  $I$ ) then
        Set  $y \leftarrow y + 1$ ;
return  $y/z$ ;

```

Basic principle of simulation Note that the meta-simulation algorithm uses $O(uz\Delta)$ time (by Theorem 9). In essence, the algorithm runs z simulations

of the process and terminates with “Simulation took too long” iff some simulation took over u steps. Hence, whenever the algorithm returns a number it is the mean of z binary random variables, each equal to 1 with probability $\Pr[\mathcal{F}_t \mid \mathcal{E}_u]$, where \mathcal{F}_t is the event that t fixates and \mathcal{E}_u is the event that fixation happens before u effective steps, when the initial type function is picked according to \mathcal{D} (we note that the conditional part was overlooked in [4], moreover, instead of steps we consider only effective steps). By ensuring that u is high enough and that the approximation is tight enough (basically, that z is high enough), we can use $\Pr[\mathcal{F}_t \mid \mathcal{E}_u]$ as an approximation of $\Pr[\mathcal{F}_t]$, as shown in the following lemma.

Lemma 10. *Let $0 < \epsilon < 1$ be given. Let \mathcal{X}, \mathcal{E} be a pair of events and x a number, such that $\Pr[\mathcal{E}] \geq 1 - \frac{\epsilon \cdot \Pr[\mathcal{X}]}{4}$ and that $x \in [(1 - \epsilon/2) \Pr[\mathcal{X} \mid \mathcal{E}], (1 + \epsilon/2) \Pr[\mathcal{X} \mid \mathcal{E}]]$. Then*

$$x \in [(1 - \epsilon) \cdot \Pr[\mathcal{X}], (1 + \epsilon) \cdot \Pr[\mathcal{X}]] .$$

The value of u : $u_{z,r}$ Consider some fixed value of z . The value of u is basically just picked so high that $\Pr[\mathcal{E}_u] \geq 1 - \frac{\epsilon \cdot \Pr[\mathcal{F}_t]}{4}$ (so that we can apply Lemma 10) and such that after taking union bound over the z trials, we have less than some constant probability of stopping. The right value of u is thus sensitive to r , but in all cases at most $O(n^2 \Delta^2 \max(\log z, \log \epsilon^{-1}))$, because of Theorem 5. More precisely, we let

$$u_{z,r} = \begin{cases} 30n \cdot \max(\log z, \log \epsilon^{-1}) & \text{if } r \geq 2\Delta \\ \frac{30n\Delta}{\min(r-1, 1)} \cdot \max(\log z, \log \epsilon^{-1}) & \text{if } 1 + \frac{1}{n \cdot \Delta} \leq r < 2\Delta \\ 20n^2 \Delta^2 \cdot \max(\log z, \log \epsilon^{-1}) & \text{if } r < 1 + \frac{1}{n \cdot \Delta} . \end{cases}$$

Algorithm Algo1 We consider the fixation problem for t_1 . Algorithm Algo1 is as follows:

1. Let \mathcal{D} be the uniform distribution over the n type functions where exactly one node is t_1 .
2. Return $\text{MetaSimulation}(t_1, z, u_{z,r}, \mathcal{D})$, for $z = 48 \cdot \frac{n}{\epsilon^2}$.

Algorithm Algo2 We consider the extinction problem for t_1 . Algorithm Algo2 is as follows:

1. Let \mathcal{D} be the uniform distribution over the n type functions where exactly one node is t_2 .
2. Return $\text{MetaSimulation}(t_1, z, u_{z,r}, \mathcal{D})$, for $z = 24/\epsilon^2$.

Algorithm Algo3 We consider the problem of (additively) approximating the fixation probability given some type function f and type t . Algorithm Algo3 is as follows:

1. Let \mathcal{D} be the distribution that assigns 1 to f .
2. Return $\text{MetaSimulation}(t, z, u_{z,r}, \mathcal{D})$, for $z = 6/\epsilon^2$.

Theorem 11. *Let G be a connected undirected graph of n nodes with the highest degree Δ , divided into two types of nodes t_1, t_2 , such that $r = w(t_1) > w(t_2) = 1$. Given $\frac{1}{2} > \epsilon > 0$, let $\alpha = n^2 \cdot \Delta \cdot \epsilon^{-2} \cdot \max(\log n, \log \epsilon^{-1})$ and $\beta = n \cdot \Delta \cdot \epsilon^{-2} \cdot \log \epsilon^{-1}$. Consider the running times:*

$$T(x) = \begin{cases} O(x) & \text{if } r \geq 2\Delta \\ O\left(\frac{x \cdot \Delta}{\min(r-1, 1)}\right) & \text{if } 1 + \frac{1}{n \cdot \Delta} \leq r < 2\Delta \\ O(n \cdot \Delta^2 \cdot x) & \text{if } 1 < r < 1 + \frac{1}{n \cdot \Delta} \end{cases} .$$

- **Fixation (resp. Extinction) problem for t_1** *Algorithm Algo1 (resp. Algo2) is an FPRAS algorithm, with running time $T(\alpha)$ (resp. $T(\beta)$), that with probability at least $\frac{3}{4}$ outputs a number in $[(1 - \epsilon) \cdot \rho, (1 + \epsilon) \cdot \rho]$, where ρ is the solution of the fixation (resp. extinction) problem for t_1 .*
- **Generalized fixation problem** *Given an initial type function f and a type t , there is an (additive approximation) algorithm, Algo3, with running time $T(\beta)$, that with probability at least $\frac{3}{4}$ outputs a number in $[\rho - \epsilon, \rho + \epsilon]$, where ρ is the solution of the generalized fixation problem given f and t .*

Remark 4. There exists no known FPRAS for the generalized fixation problem and since the fixation probability might be exponentially small such an algorithm might not exist. (It is exponentially small for fixation of t_2 , even in the Moran process (that is, when the graph is complete) when there initially is 1 node of type t_2)

Alternative algorithm for extinction for t_2 We also present an alternative algorithm for extinction for t_2 when r is big. This is completely different from the techniques of [4]. The alternative algorithm is based on the following result where we show for big r that $1/r$ is a good approximation of the extinction probability for t_2 , and thus the algorithm is polynomial even for big r in binary.

Theorem 12. *Consider an undirected graph G and consider the extinction problem for t_2 on G . If $r \geq \max(\Delta^2, n)/\epsilon$, then $\frac{1}{r} \in [(1 - \epsilon) \cdot \rho, (1 + \epsilon) \cdot \rho]$, where ρ is the solution of the extinction problem for t_2 .*

Proof sketch We present a proof sketch, and details are in the Appendix. We have two cases:

- By [4, Lemma 4], we have $\rho \geq \frac{1}{n+r}$. Thus, $(1 + \epsilon) \cdot \rho \geq \frac{1}{r}$, as desired, since $n/\epsilon \leq r$.
- On the other hand, the probability of fixation for t_2 in the first effective step is at most $\frac{1}{r+1} < \frac{1}{r}$ (we show this in Lemma 20 in the Appendix).

The probability that fixation happens for t_2 after the first effective step is at most ϵ/r because of the following reason: By Lemma 3, the probability of increasing the number of members of t_2 is at most $p := \frac{1}{r/\Delta+1}$ and otherwise it decrements. We then model the problem as a Markov chain M with state space corresponding to the number of members of t_1 , using p as the probability to decrease the current state. In M the starting state is state 2 (after the first effective step, if fixation did not happen, then the number of members of t_1 is 2). Using that $\Delta^2/\epsilon \leq r$, we see that the probability of absorption in state 0 of M from state 2 is less than ϵ/r . Hence, ρ is at most $(1 + \epsilon)/r$ and $(1 - \epsilon)\rho$ is thus less than $1/r$.

□

Remark 5. While Theorem 12 is for undirected graphs, a variant (with larger r and which requires the computation of the pr. that t_1 goes extinct in the first step) can be established even for directed graphs, see the Appendix.

Concluding remarks In this work we present faster Monte-Carlo algorithms for approximating fixation probability for undirected graphs (see Remark 6 in the Appendix for detailed comparison). An interesting open question is whether the fixation probability can be approximated in polynomial time for directed graphs.

References

- [1] B. Adlam, K. Chatterjee, and M. Nowak. Amplifiers of selection. In *Proc. R. Soc. A*, volume 471, page 20150114. The Royal Society, 2015.
- [2] F. Débarre, C. Hauert, and M. Doebeli. Social evolution in structured populations. *Nature Communications*, 2014.
- [3] J. Díaz, L. A. Goldberg, G. B. Mertzios, D. Richerby, M. Serna, and P. G. Spirakis. On the fixation probability of superstars. *Proceedings of the Royal Society A: Mathematical, Physical and Engineering Science*, 469(2156), 2013.
- [4] J. Díaz, L. A. Goldberg, G. B. Mertzios, D. Richerby, M. Serna, and P. G. Spirakis. Approximating Fixation Probabilities in the Generalized Moran Process. *Algorithmica*, 69(1):78–91, 2014 (Conference version SODA 2012).
- [5] W. Ewens. *Mathematical Population Genetics 1: I. Theoretical Introduction*. Interdisciplinary Applied Mathematics. Springer, 2004.
- [6] M. Frean, P. B. Rainey, and A. Traulsen. The effect of population structure on the rate of evolution. *Proceedings of the Royal Society B: Biological Sciences*, 280(1762), 2013.
- [7] A. Galanis, A. Göbel, L. A. Goldberg, J. Lapinskas, and D. Richerby. Amplifiers for the Moran Process. In *43rd International Colloquium on Automata, Languages, and Programming (ICALP 2016)*, volume 55, pages 62:1–62:13, 2016.
- [8] A. L. Hill, D. G. Rand, M. A. Nowak, and N. A. Christakis. Emotions as infectious diseases in a large social network: the SISa model. *Proceedings of the Royal Society B: Biological Sciences*, 277:3827–3835, 2010.
- [9] R. Ibsen-Jensen, K. Chatterjee, and M. A. Nowak. Computational complexity of ecological and evolutionary spatial dynamics. *Proceedings of the National Academy of Sciences*, 112(51):15636–15641, 2015.
- [10] S. Karlin and H. M. Taylor. *A First Course in Stochastic Processes, Second Edition*. Academic Press, 2 edition, Apr. 1975.
- [11] E. Lieberman, C. Hauert, and M. A. Nowak. Evolutionary dynamics on graphs. *Nature*, 433(7023):312–316, Jan. 2005.
- [12] P. A. P. Moran. *The Statistical Processes of Evolutionary Theory*. Oxford University Press, Oxford, 1962.
- [13] M. A. Nowak. *Evolutionary Dynamics: Exploring the Equations of Life*. Harvard University Press, 2006.
- [14] M. A. Nowak and R. M. May. Evolutionary games and spatial chaos. *Nature*, 359:826, 1992.

- [15] H. Ohtsuki, C. Hauert, E. Lieberman, and M. A. Nowak. A simple rule for the evolution of cooperation on graphs and social networks. *Nature*, 441:502–505, 2006.
- [16] M. Serna, D. Richerby, L. A. Goldberg, and J. Díaz. Absorption time of the moran process. *Random Structures & Algorithms*, 48(1):137–159, 2016.
- [17] P. Shakarian, P. Roos, and A. Johnson. A review of evolutionary graph theory with applications to game theory. *Biosystems*, 107(2):66 – 80, 2012.

Appendix

7 Details of Section 3

In this section, we prove Lemma 1 and Theorem 5.

Lemma 1. Fix any type function f such that neither type has fixated. Let f_d (resp., f_m) be the next type function under dynamic evolution step (resp., modified dynamic evolution step). Then, $\Pr[f \neq f_d] > 0$ and for all type functions f' we have: $\Pr[f' = f_d \mid f \neq f_d] = \Pr[f' = f_m]$.

Proof. Let f, f_d, f_m be as in the lemma statement. First note that $\Pr[f_d \neq f] > 0$ since G is connected and no type has fixated in f . Therefore, there must be some edge $(u, v) \in E$, such that $f(u) \neq f(v)$. There is a positive probability that u and v is selected by the unmodified evolution step, in which case $f_d \neq f$. Thus $\Pr[f_d \neq f] > 0$.

We consider the node selection in the two cases:

1. Let v be the node picked in the first part of the unmodified dynamic evolution step and u be the node picked in the second part.
2. Similarly, let v' be the node picked in the first part of the modified dynamic evolution step and u' be the node picked in the second part.

Observe that $(u, v), (u', v') \in E$. We will argue that for all $(u'', v'') \in E$, we have that

$$\Pr[u = u'' \cup v = v'' \mid f(u) \neq f(v)] = \Pr[u' = u'' \cup v' = v''].$$

This clearly implies the lemma statement, since $f_d = f[u \rightarrow f(v)]$ and $f_m = f[u' \rightarrow f(v')]$, by the last part of the unmodified and modified dynamic evolution step. If $f(u'') = f(v'')$, then $\Pr[u = u'' \cup v = v'' \mid f(u) \neq f(v)] = 0$, because $f(u'') = f(u) \neq f(v) = f(v'')$ which contradicts that $f(u'') = f(v'')$. Also, if $f(u'') = f(v'')$, then $\Pr[u' = u'' \cup v' = v''] = 0$, because $f(u'') = f(u') \neq f(v') = f(v'')$ (note $f(u') \neq f(v')$ because u' was picked from $\Gamma_{v'}$), again contradicting that $f(u'') = f(v'')$.

We therefore only need to consider that $(u'', v'') \in E$ and $f(u'') \neq f(v'')$. The probability to pick v and then pick u in an unmodified dynamic evolution step is $\frac{w(f(v))}{W(f)} \cdot \frac{1}{\deg v}$, for any $(u, v) \in E$ and especially the ones for which $f(u) \neq f(v)$. Hence, $\Pr[u = u'' \cup v = v''] = \frac{f(v'')}{W(f)} \cdot \frac{1}{\deg v''}$. Thus, also, $\Pr[(u = u'' \cup v = v'') \cap (f(u) \neq f(v))] = \frac{w(f(v''))}{W(f)} \cdot \frac{1}{\deg v''}$. We also have that

$$\Pr[f(u) \neq f(v)] = \sum_{\substack{(u,v) \in E \\ f(u) \neq f(v)}} \frac{w(f(u))}{W(f)} \cdot \frac{1}{\deg u} = \sum_{v \in V} \frac{w(f(v))}{W(f)} \cdot \frac{1}{\deg v} \cdot |\Gamma_v(f)| = \frac{W'(f)}{W(f)},$$

where the second equality comes from that $\Gamma_v(f)$ is the set of nodes u such that $(u, v) \in E$ and $f(u) \neq f(v)$. Thus,

$$\Pr[u = u'' \cup v = v'' \mid f(u) \neq f(v)] = \frac{\Pr[(u = u'' \cup v = v'') \cap (f(u) \neq f(v))]}{\Pr[f(u) \neq f(v)]} = \frac{w(f(v''))}{W'(f)} \cdot \frac{1}{\deg v''}$$

The probability to pick v' and then pick u' in a modified dynamic evolution step, for some $(u', v') \in E$ is

$$\frac{w(f(v')) \cdot \frac{|\Gamma_{v'}(f)|}{\deg v'}}{W'(f)} \cdot \frac{1}{|\Gamma_{v'}(f)|} = \frac{w(f(v'))}{W'(f)} \cdot \frac{1}{\deg v'}$$

Hence,

$$\Pr[u' = u'' \cup v' = v''] = \frac{w(f(v''))}{W'(f)} \cdot \frac{1}{\deg v''} = \Pr[u = u'' \cup v = v'' \mid f(u) \neq f(v)]$$

This completes the proof of the lemma. \square

Next, the proof of Theorem 5.

Theorem 5. Let t_1 and t_2 be the two types, such that $r = w(t_1) > w(t_2) = 1$. Let Δ be the maximum degree. Let k be the number of nodes of type t_2 in the initial type function. The following assertions hold:

- *Bounds dependent on r*
 1. *Expected steps* The process requires at most $3k\Delta / \min(r-1, 1)$ effective steps in expectation, before fixation is reached.
 2. *Probability* For any integer $x \geq 1$, after $6xn\Delta / \min(r-1, 1)$ effective steps, the probability that the process has not fixated is at most 2^{-x} , irrespective of the initial type function.
- *Bounds independent on r*
 1. *Expected steps* The process requires at most $2nk\Delta^2$ effective steps in expectation, before fixation is reached.
 2. *Probability* For any integer $x \geq 1$, after $4xn^2\Delta^2$ effective steps, the probability that the process has not fixated is at most 2^{-x} , irrespective of the initial type function.
- *Bounds for $r \geq 2\Delta$*
 1. *Expected steps* The process requires at most $3k$ effective steps in expectation, before fixation is reached.
 2. *Probability* For any integer $x \geq 1$, after $6xn$ effective steps, the probability that the process has not fixated is at most 2^{-x} , irrespective of the initial type function.

Proof. Observe that if $k = 0$, then fixation has been reached in 0 (effective) steps. Thus assume $k \geq 1$. We first argue about the first item of each case (i.e., about expected steps) and then about the second item (i.e., probability) of each case.

Expected steps of first item In every step, for $r > 1$, the potential increases by $-k_2 = \frac{r-1}{\Delta \cdot (r+1)}$, unless fixation has been achieved, in expectation by Lemma 2. Let $n' = \sum_{v \in V} \frac{1}{\deg v}$ be the maximum potential. Let f be the initial type function. Observe that the potential initially is $\psi(f) = \sum_{v \in V_{t_1, f}} \frac{1}{\deg v} = \sum_{v \in V} \frac{1}{\deg v} - \sum_{v \in V_{t_2, f}} \frac{1}{\deg v} \geq n' - k$. Hence, the potential just needs to increase by at most k . Similar to [4], we consider a modified process where if t_2 fixates, then the next type function f' has a uniformly random node mapped to t_1 (note that this only increases the time steps till t_1 fixates). We see that in every step, unless fixation happens for t_1 , the potential increases by $-k_2$ (in case fixation for t_2 has happened in the previous step, the potential increases by at least $\frac{1}{\Delta} > -k_2$, since $\frac{r-1}{r+1} < 1$). Applying [4, Theorem 6], with $k_1 = k$ and $-k_2$, we have that the number of effective steps in expectation, when starting with the type function f , is at most $\frac{k\Delta(r+1)}{(r-1)}$, for the modified process. Thus the upper bound also follows for the original process, which fixates faster. Note that for $r \geq 2$ we have that $\frac{r+1}{r-1} \leq 3$ and that for $1 < r < 2$ we have that $\frac{r+1}{r-1} < \frac{3}{r-1}$. Thus, $\frac{r+1}{r-1} \leq \frac{3}{\min(r-1, 1)}$. This establishes the expected number of effective steps for the first item.

Expected steps of second item The fact that $2nk\Delta^2$ effective steps is sufficient in expectation for r close to 1 can be seen as follows. Let f^* denote the type function that assigns all nodes to t_1 . Let f be the initial type function, and we denote by \bar{f} its complement, i.e., \bar{f} maps every node to the opposite type as compared to f . Similar to the proof² of [4, Theorem 11] we get that $\Delta^2((\psi(f^*))^2 - (\psi(f))^2)$ is sufficient in expectation, where f is the initial type function. The change as compared to the proof of [4, Theorem 11] consists of using that $\mathbb{E}[(\psi(f') - \psi(f''))^2] \geq \Delta^{-2}$ instead of [4, Equation (3)], where f' is a fixed type function and f'' the following type function according to a modified dynamic evolution step. Let a be $\psi(f)$ and b be $\psi(f^*) - \psi(f) = \psi(\bar{f})$. Hence, $\Delta^2((\psi(f^*))^2 - (\psi(f))^2) = \Delta^2((a+b)^2 - a^2) = \Delta^2(2ab + b^2)$. This is monotone increasing in a and b (since, a, b, Δ are positive). For all type functions \hat{f} , we have that $\psi(\hat{f})$ is at most the number of nodes assigned to t_1 by \hat{f} . Hence, we have that $a = \psi(f) \leq n - k$ and $b = \psi(\bar{f}) \leq k$. Thus

$$\Delta^2((\psi(f^*))^2 - (\psi(f))^2) = \Delta^2(2ab + b^2) \leq \Delta^2(2(n-k)k + k^2) = \Delta^2(2nk - k^2) \leq 2nk\Delta^2,$$

as desired.

Expected steps of third item We can consider the potential function ψ' , which is $\psi'(f') = |V_{t_1, f'}|$. We see that in every effective step the potential increases by at least $\frac{1}{3}$ in expectation, by Lemma 3. Similar to the first item, we can then define a modified process and apply [4, Theorem 6], and see that $3k$ effective steps suffices in expectation.

²Note that the proof of [4, Theorem 11] does not directly use that $r = 1$, even though the statement they end up with does

Probability bounds of the three cases Follows from the above three items and Lemma 4. \square

8 Details of Section 4

In this section we prove Lemma 6 and Lemma 7.

Lemma 6. With pr. above $\approx \frac{1-1/r}{2}$ if at least one of V'_L is initially of type t_1 , then t_1 fixates.

Proof. In this proof we consider that there is initially 1 member of t_1 . As shown by [16], the fixation probability is increasing in the number of members.

If the graph just consisted of the nodes in V_L (and v_1 was only connected to v_2), then if the initial state had a single member of type t_1 , the pr. that t_1 fixated would be $\frac{1-1/r}{1-r^{x-n}} \approx 1-1/r$ (using that it is in essence the gambler's ruin problem when the probability for winning each round for t_2 is $\frac{1}{r+1}$, but t_2 starts with $n-x-1$ pennies compared to 1 for t_1). We see that in the original graph, the only difference is that v_1 is connected to the stars-on-a-cycle graph. Thus, we see that as long as v_1 is always a member of t_2 , there are no differences. We can use that as follows: If in the initial type function the lone member of t_1 is in V'_L , then with pr. above $1/2$, the node v_{n-x} becomes of type t_1 before v_1 , because at all times (as long as neither v_1 or v_{n-x} has ever been of type t_1), the nodes of type t_1 forms an interval $\{i, \dots, j\}$ and the pr. that in the next iteration v_{i-1} becomes of type t_1 is equal to the pr. that in the next iteration v_{j+1} becomes of type t_1 . Hence, we get that with pr. above $\frac{1-1/r}{2(1-r^{x-n})} \approx \frac{1-1/r}{2}$ at some point the members of t_1 is exactly V_L .

But if t_2 fixates after all members of V_L has become of type t_1 , V_L must thus eventually go from being all t_1 to being all t_2 . We will now argue that it is exponentially unlikely to happen. For V_L to change to being all t_2 , we must have that in some step, v_1 has become a member of type t_2 . We can now consider a stronger version of t_2 (that thus have more pr. to fixate), where v_1 can only become of type t_1 if it was reproduced to from v_2 . We will say that t_2 starts an attempt whenever v_1 becomes of type t_2 . We say that t_2 wins the attempt if V_L eventually becomes only of type t_2 and we say that t_2 loses the attempt if V_L becomes of type t_1 . Note that t_2 needs to win an attempt to fixate. Again, using gambler's ruin, we see that t_2 wins any attempt with pr. $1 - \frac{1-r^{x-n-1}}{1-r^{x-n}} = 1 - \frac{r^{n-x}-r^{-1}}{r^{n-x}-1} \approx \frac{1}{r^{n-x}-1}$. But the process fixates, as we showed earlier, in $3k\Delta/\min(r-1, 1)$ effective steps, which is then an upper bound on the number of attempts. But then the pr. that t_2 ever wins an attempt is at most

$$3k\Delta/\min(r-1, 1) \cdot \left(1 - \frac{r^{n-x} - r^{-1}}{r^{n-x} - 1}\right) \approx \frac{3k\Delta}{r^{n-x}},$$

which is exponentially small for fixed r and we thus see that the pr. that t_1 eventually has all of V_L (if one of $\{\ell_{\lceil n/4 \rceil}, \dots, \ell_{n-x}\}$ is initially of type t_1) is approximately the pr. that t_1 fixates. \square

Recall that whenever a node of V_C^i , for some i , changes type, we say that a leaf-step occurred.

Lemma 7. The pr. that an effective step is a leaf-step is at most $\frac{r}{\Delta}$.

Proof. Consider some initial type function f . We will consider the conditional probability of a leaf-step conditioned on a node (but not c_1) in the stars-on-a-cycle graph is selected for reproduction (clearly, if no such node is selected for reproduction, the pr. of a leaf-step is 0). For simplicity, we will furthermore condition on the node that gets selected for reproduction is in $\{c_i\} \cup V_C^i$ for some i (this is without loss of generality because all nodes in the stars-on-a-cycle graph is in one of those sets). Let $t := f(c_i)$ and let t' be the other type. We will consider that $x > 0$ nodes of V_C^i is of type t' (if no nodes of V_C^i is of type t' then the pr. of a leaf-step, conditioned on a node of $\{c_i\} \cup V_C^i$ being selected for reproduction is 0). For a leaf step to occur, we need to do as follows: we must select c_i for reproduction and we must select one of the nodes of V_C^i for death. Let $x' \in \{x, x+1, x+2\}$ (the +2 is because of the other center nodes) be the number of neighbors of c_i which are of type t' . The fitness of c_i is then $w(t) \cdot \frac{x'}{\Delta}$. The fitness of each node of type t' in V_C^i is $w(t')$. If c_i is picked, the pr. of a leaf-step is $\frac{x'}{x}$ (because if a center neighbor is selected, no leaf-step occurred). The pr. of a leaf step is then

$$\frac{w(t) \cdot \frac{x'}{\Delta} \cdot \frac{x}{x'}}{xw(t') + w(t)} = \frac{w(t)}{\Delta(w(t') + w(t)/x)} \leq \frac{w(t)}{\Delta w(t')}$$

Since in the worst case $w(t) = r$ and $w(t') = 1$, we get our result. □

9 Details of Section 5

In this section we consider the weaker model, where getting even a single random bit costs constant time.

We can use the same data-structure as in Section 5, since the data-structure is deterministic. Thus, we just need to argue how to sample an effective step given the data-structure. (i.e. like in the paragraph **Sampling an effective step** of Section 5).

There are thus 3 things we need to consider, namely (1) picking random list L among the 2Δ lists, proportional to their weight; (2) pick a random node v from L and (3) pick a node at random in $\Gamma_v(f)$.

Picking a random list among the 2Δ lists, proportional to their weight takes $O(\Delta)$ time and uses $O(\log \Delta)$ random bits, e.g. as follows: We can build a binary, balanced tree where each leaf corresponds to a list. We can then annotate each node of the tree with the sum of the weights of all lists below it. This can be done in $O(\Delta)$ time. Afterwards, we can then find a random list proportional to its weight using $O(\log \Delta)$ bits in expectation as follows: Start at the root of the tree.

1. When looking at an internal node v do as follows: Let y be the number annotated on v and let z be the number annotated on the left child. Construct a random number x bit for bit, starting with the most significant, between 0 and $w = 2^{\lceil \log y \rceil}$. Let x_i be the random variable denoting x after i bits have been added. Let $x_{i,1}$ be $x_i + w - 2^i - 1$, i.e. the greatest number x can become if the remaining bits are all 1. If, after having added the i 'th bit, $x_{i,1} \leq z$, continue to the left child. If, after having added the i 'th bit, $x_i > z$ and $x_{i,1} \leq y$ continue to the right child. If, after having added the i 'th bit, $x_i > y$, start over at node v . Otherwise, add bit $i + 1$ to x .

When adding a bit to x in step i , at least one of $x_{i,1} \leq z$ or $x_i > z$ becomes satisfied with pr. $1/2$. Similar, at least one of $x_{i,1} \leq y$ or $x_i > y$ becomes satisfied with pr. $1/2$. Thus, in expectation, we need four steps before we continue to either the left, the right or back to v . Since $2^{\lceil \log y \rceil} < 2y$, we have that we start over at v with pr. at most $1/2$. Thus, in expectation, after less than 8 steps, we go to either the left or the right child. Hence, after $O(\log \Delta)$ steps we reach a list.

2. When looking at a list, output it.

We can find a random element in the atleast half-full list L by repeatedly finding a random number x between 0 and $2^{\lceil \log s \rceil}$, where $s \leq n$ is the size of L . If x is below s and is in L , output it, otherwise, find the next random number.

Note that $2^{\lceil \log s \rceil} < 2s$. Thus, $x \leq s$ with pr. $1/2$. If so, we have another $1/2$ that entry x will be filled. Hence, we need to pick 4 numbers in expectation. Note that this thus takes $O(\log n)$ time/random bits.

We can do similar to find an element in $\Gamma_v(f)$ in $O(\log(\Delta))$ time/random bits.

This leads to the following theorem.

Theorem 13. *If no more than one random bit can be accessed in constant time, then an effective step can be sampled in (amortized and expected) $O(\Delta + \log n)$ time, using $O(\log n)$ expected random bits, after deterministic $O(m)$ preprocessing.*

10 Details of Section 6

In this section, we prove Lemma 10 and give the correctness and time-bound arguments for our algorithms, which implies Theorem 11.

Lemma 10. Let $0 < \epsilon < 1$ be given. Let \mathcal{X}, \mathcal{E} be a pair of events and x a number, such that $\Pr[\mathcal{E}] \geq 1 - \frac{\epsilon \cdot \Pr[\mathcal{X}]}{4}$ and that $x \in [(1 - \epsilon/2) \Pr[\mathcal{X} \mid \mathcal{E}], (1 + \epsilon/2) \Pr[\mathcal{X} \mid \mathcal{E}]]$. Then

$$x \in [(1 - \epsilon) \cdot \Pr[\mathcal{X}], (1 + \epsilon) \cdot \Pr[\mathcal{X}]] .$$

Proof. Fix $0 < \epsilon < 1$. Let \mathcal{X}, \mathcal{E} be a pair of events, such that $\Pr[\mathcal{E}] \geq 1 - \frac{\epsilon \cdot \Pr[\mathcal{X}]}{4}$ and that $x \in [(1 - \epsilon/2) \Pr[\mathcal{X} \mid \mathcal{E}], (1 + \epsilon/2) \Pr[\mathcal{X} \mid \mathcal{E}]]$.

We have that

$$\begin{aligned}
\Pr[\mathcal{X}] &= \Pr[\mathcal{E}] \cdot \Pr[\mathcal{X} \mid \mathcal{E}] + (1 - \Pr[\mathcal{E}]) \cdot \Pr[\mathcal{X} \mid \neg\mathcal{E}] \\
&\geq \left(1 - \frac{\epsilon \cdot \Pr[\mathcal{X}]}{4}\right) \cdot \Pr[\mathcal{X} \mid \mathcal{E}] \\
&\geq \Pr[\mathcal{X} \mid \mathcal{E}] - \frac{\epsilon \cdot \Pr[\mathcal{X}]}{4} .
\end{aligned}$$

Also,

$$\begin{aligned}
\Pr[\mathcal{X}] &= \Pr[\mathcal{E}] \cdot \Pr[\mathcal{X} \mid \mathcal{E}] + (1 - \Pr[\mathcal{E}]) \cdot \Pr[\mathcal{X} \mid \neg\mathcal{E}] \\
&\leq \Pr[\mathcal{X} \mid \mathcal{E}] + \frac{\epsilon \cdot \Pr[\mathcal{X}]}{4} \cdot \Pr[\mathcal{X} \mid \neg\mathcal{E}] \\
&\leq \Pr[\mathcal{X} \mid \mathcal{E}] + \frac{\epsilon \cdot \Pr[\mathcal{X}]}{4} .
\end{aligned}$$

We now see that

$$\begin{aligned}
x &\in [(1 - \epsilon/2) \Pr[\mathcal{X} \mid \mathcal{E}], (1 + \epsilon/2) \Pr[\mathcal{X} \mid \mathcal{E}]] \\
\Rightarrow x &\in [(1 - \epsilon/2)(1 - \epsilon/4) \Pr[\mathcal{X}], (1 + \epsilon/2)(1 + \epsilon/4) \Pr[\mathcal{X}]]
\end{aligned}$$

Note that

$$(1 - \epsilon/2) \cdot (1 - \epsilon/4) = 1 + \epsilon^2/8 - 3\epsilon/4 > 1 - \epsilon$$

and

$$(1 + \epsilon/2) \cdot (1 + \epsilon/4) = 1 + \epsilon^2/8 + 3\epsilon/4 < 1 + 7\epsilon/8 < 1 + \epsilon ,$$

using that $0 < \epsilon^2 < \epsilon < 1$. Hence,

$$x \in [(1 - \epsilon) \Pr[\mathcal{X}], (1 + \epsilon) \Pr[\mathcal{X}]] ,$$

as desired. \square

We will use the following lemmas from previous works.

Lemma 14 ([5]). *The solution of the fixation problem for t_1 is at least $\frac{1}{n}$ and the solution of the extinction problem for t_1 is at least $1 - \frac{1}{n}$.*

Lemma 15 ([4]). *The solution of the extinction problem is greater than $\frac{1}{n+r}$ for t_2 .*

We next show the following lemma, which will be useful in the correctness proofs of our algorithms.

Lemma 16. *Consider running $\text{MetaSimulation}(t, z, u, \mathcal{D})$ on an instance where $w(t_1) = r$, for some distribution \mathcal{D} , type t and numbers z , $0 < \epsilon < \frac{1}{2}$, when $u = u_{z,r}$. Then, $1 - \Pr[\mathcal{E}_u] \leq \min(z^{-5}, \epsilon^5)$ and the probability that MetaSimulation does not output a number is smaller than $\min(z^{-4}, \epsilon^4)$. The run time T is*

$$T = \begin{cases} O(n\Delta z \cdot \max(\log z, \log \epsilon^{-1})) & \text{if } r \geq 2\Delta \\ O(n\Delta^2 \frac{z}{\min(r-1, 1)} \cdot \max(\log z, \log \epsilon^{-1})) & \text{if } 1 + \frac{1}{n\Delta} \leq r < 2\Delta \\ O(n^2 \Delta^3 z \cdot \max(\log z, \log \epsilon^{-1})) & \text{if } r < 1 + \frac{1}{n\Delta} \end{cases}$$

Proof. Let $t, z, \mathcal{D}, r, \epsilon$ be as in the lemma statement. Also, let $u = u_{z,r}$. The run time of $\text{MetaSimulation}(t, z, u, \mathcal{D})$ is $O(uz\Delta)$ (by Theorem 9), which leads to the desired run time when inserting $u_{z,r}$. By choice of u , we have that

$$1 - \Pr[\mathcal{E}_u] \leq 2^{-5 \max(\log z, \log \epsilon^{-1})} = \min(z^{-5}, \epsilon^5) ,$$

using Theorem 5. Thus, any single simulation takes more than u time with probability less than $\min(z^{-5}, \epsilon^5)$. Union bounding the z simulations, we see that the probability that any one takes more than u time is at most $\min(z^{-4}, \epsilon^{-4})$. Note that only in case one simulation takes more than u time does the algorithm output something which is not a number. \square

The remainder of this section is about showing the correctness and running times of the various algorithms.

Lemma 17. *Algorithm Algo1 is correct and runs in time T_1 , for*

$$T_1 = \begin{cases} O(n^2 \cdot \Delta \cdot \epsilon^{-2} \cdot \max(\log n, \log \epsilon^{-1})) & \text{if } r \geq 2\Delta \\ O\left(\frac{n^2 \cdot \Delta^2}{\epsilon^2 \cdot \min(r-1, 1)} \cdot \max(\log n, \log \epsilon^{-1})\right) & \text{if } 1 + \frac{1}{n \cdot \Delta} \leq r < 2\Delta \\ O(n^3 \cdot \Delta^3 \cdot \epsilon^{-2} \cdot \max(\log n, \log \epsilon^{-1})) & \text{if } 1 < r < 1 + \frac{1}{n \cdot \Delta} \end{cases}$$

Proof. Recall if $r = 1$, then the fixation probability of t_1 is $\frac{1}{n}$. We consider the other cases. Otherwise, the output of **Algo1** is $\text{MetaSimulation}(t_1, z, u_{z,r}, \mathcal{D})$, for $z = 48 \cdot \frac{n}{\epsilon^2}$. The run time then follows from Lemma 16.

We will utilize Lemma 10 to show the correctness of **Algo1**. We thus need to ensure that $\Pr[\mathcal{E}_u] \geq 1 - \frac{\epsilon \cdot \Pr[\mathcal{F}_{t_1}]}{4}$ and that with probability $\frac{3}{4}$ our algorithm outputs x , for

$$x \in [(1 - \epsilon/2) \Pr[\mathcal{F}_{t_1} \mid \mathcal{E}_u], (1 + \epsilon/2) \Pr[\mathcal{F}_{t_1} \mid \mathcal{E}_u]] .$$

According to Lemma 16, we have that

$$\Pr[\mathcal{E}_u] \geq 1 - \min(z^{-5}, \epsilon^5) = 1 - z^{-5} \geq 1 - \frac{\epsilon}{4n} \geq 1 - \frac{\epsilon \cdot \Pr[\mathcal{F}_{t_1}]}{4} ,$$

using that $\Pr[\mathcal{F}_{t_1}]$ is at least $\frac{1}{n}$ according to Lemma 14 as desired.

Error sources We see that there are two cases where **Algo1** does not output a number in $[(1 - \epsilon) \cdot \Pr[\mathcal{F}_{t_1}], (1 + \epsilon) \cdot \Pr[\mathcal{F}_{t_1}]]$:

1. The algorithm does not output a number. This happens with probability at most

$$\min(z^{-4}, \epsilon^4) = \left(48 \cdot \frac{n}{\epsilon^2}\right)^{-4} < \left(48 \cdot \frac{2}{(1/2)^2}\right)^{-4} = 384^{-4} ,$$

according to Lemma 16.

2. The algorithm outputs a number outside $[(1 - \epsilon) \cdot \Pr[\mathcal{F}_{t_1}], (1 + \epsilon) \cdot \Pr[\mathcal{F}_{t_1}]]$.

Simulations are far from expectation Let the output of the algorithm, conditioned on it being a number, be the random variable \mathcal{B} . Observe that if

$$\mathcal{B} \in [(1 - \epsilon/2) \cdot \Pr[\mathcal{F}_{t_1} \mid \mathcal{E}_u], (1 + \epsilon/2) \cdot \Pr[\mathcal{F}_{t_1} \mid \mathcal{E}_u]] ,$$

then, according to Lemma 10, we have that $\mathcal{B} \in [(1 - \epsilon) \cdot \Pr[\mathcal{F}_{t_1}], (1 + \epsilon) \cdot \Pr[\mathcal{F}_{t_1}]]$, since we already argued that $\Pr[\mathcal{E}_u] \geq 1 - \frac{\epsilon \cdot \Pr[\mathcal{F}_{t_1}]}{4}$. We are thus interested in the probability

$$\Pr[|\mathcal{B} - \Pr[\mathcal{F}_{t_1} \mid \mathcal{E}_u]| \geq \epsilon/2 \mid \mathcal{E}_u] .$$

Applying Lemma 10, with $\mathcal{X} = \mathcal{F}_{t_1}$, $\mathcal{E} = \mathcal{E}_u$ and $x = \Pr[\mathcal{F}_{t_1} \mid \mathcal{E}_u]$, we see that $\Pr[\mathcal{F}_{t_1} \mid \mathcal{E}_u] \geq (1 - \epsilon) \cdot \Pr[\mathcal{F}_{t_1}] \geq \frac{\Pr[\mathcal{F}_{t_1}]}{2} \geq \frac{1}{2n}$. Thus,

$$\Pr[|\mathcal{B} - \Pr[\mathcal{F}_{t_1} \mid \mathcal{E}_u]| \geq \epsilon/2 \mid \mathcal{E}_u] \leq \Pr[|\mathcal{B} - \Pr[\mathcal{F}_{t_1} \mid \mathcal{E}_u]| \geq \frac{\epsilon}{4n}] .$$

Since \mathcal{B} is the average of z independent simulations of a binary random variable, each of which are 1 with probability $\Pr[\mathcal{F}_{t_1} \mid \mathcal{E}_u] \geq \frac{1}{n}$ (by Lemma 14), we can apply the multiplicative Chernoff bound and obtain:

$$\Pr[|z\mathcal{B} - z\Pr[\mathcal{F}_{t_1} \mid \mathcal{E}_u]| \geq \frac{z\epsilon}{4n}] \leq 2e^{-2z \frac{\epsilon^2}{3 \cdot 4^{2n}}} = 2e^{-2} .$$

We have that $384^{-4} + 2e^{-2} < 1/3$. The desired result follows. \square

Lemma 18. *Algorithm Algo2 is correct and runs in time T_2 , for*

$$T_2 = \begin{cases} O(n \cdot \Delta \cdot \epsilon^{-2} \cdot \log \epsilon^{-1}) & \text{if } r \geq 2\Delta \\ O\left(\frac{n \cdot \Delta^2 \cdot \log \epsilon^{-1}}{\epsilon^2 \cdot \min(r-1, 1)}\right) & \text{if } 1 + \frac{1}{n \cdot \Delta} \leq r < 2\Delta \\ O(n^2 \cdot \Delta^3 \cdot \epsilon^{-2} \cdot \log \epsilon^{-1}) & \text{if } 1 < r < 1 + \frac{1}{n \cdot \Delta} \end{cases}$$

Proof. The proof is similar to Lemma 17, except applying multiplicative Chernoff Bound to

$$\Pr[|z\mathcal{B} - z\Pr[\mathcal{F}_{t_1} \mid \mathcal{E}_u]| \geq z\epsilon \Pr[\mathcal{F}_{t_1} \mid \mathcal{E}_u]/2] \leq \Pr[|\mathcal{B} - \Pr[\mathcal{F}_{t_1} \mid \mathcal{E}_u]| \geq \frac{\epsilon}{4}] ,$$

using that $\Pr[\mathcal{F}_{t_1} \mid \mathcal{E}_u] \geq 1 - \frac{1}{n} \geq \frac{1}{2}$ by Lemma 14. \square

Lemma 19. *Algorithm Algo3 is correct and runs in time T_3 , for*

$$T_3 = \begin{cases} O(n \cdot \Delta \cdot \epsilon^{-2} \cdot \log \epsilon^{-1}) & \text{if } r \geq 2\Delta \\ O\left(\frac{n \cdot \Delta^2 \cdot \log \epsilon^{-1}}{\epsilon^2 \cdot \min(r-1, 1)}\right) & \text{if } 1 + \frac{1}{n \cdot \Delta} \leq r < 2\Delta \\ O(n^2 \cdot \Delta^3 \cdot \epsilon^{-2} \cdot \log \epsilon^{-1}) & \text{if } 1 \leq r < 1 + \frac{1}{n \cdot \Delta} \end{cases}$$

Proof. The proof is somewhat similar to Lemma 17, but there are some differences and thus we include the proof.

The output of `Algo3` is `MetaSimulation(t, z, uz, r, D)`, for $z = 6/\epsilon^2$ and \mathcal{D} the distribution assigning probability 1 to f . We see that

$$\begin{aligned} \Pr[\mathcal{F}_t] &= \Pr[\mathcal{E}_u] \cdot \Pr[\mathcal{F}_t \mid \mathcal{E}_u] + (1 - \Pr[\mathcal{E}_u]) \cdot \Pr[\mathcal{F}_t \mid \neg\mathcal{E}_u] \geq \Pr[\mathcal{F}_t \mid \mathcal{E}_u] \cdot \Pr[\mathcal{F}_t \mid \mathcal{E}_u] \\ &\geq (1 - z^{-5}) \cdot \Pr[\mathcal{F}_t \mid \mathcal{E}_u] \geq \Pr[\mathcal{F}_t \mid \mathcal{E}_u] - z^{-5} \geq \Pr[\mathcal{F}_t \mid \mathcal{E}_u] - \epsilon/2, \end{aligned}$$

using Lemma 16. Also,

$$\Pr[\mathcal{F}_t] = \Pr[\mathcal{E}_u] \cdot \Pr[\mathcal{F}_t \mid \mathcal{E}_u] + (1 - \Pr[\mathcal{E}_u]) \cdot \Pr[\mathcal{F}_t \mid \neg\mathcal{E}_u] \leq \Pr[\mathcal{F}_t \mid \mathcal{E}_u] + z^{-5} \leq \Pr[\mathcal{F}_t \mid \mathcal{E}_u] + \epsilon/2.$$

Thus, we just need to ensure that we output a number x in $[\Pr[\mathcal{F}_t \mid \mathcal{E}_u] - \epsilon/2, \Pr[\mathcal{F}_t \mid \mathcal{E}_u] + \epsilon/2]$ with probability at least $\frac{3}{4}$.

We output a number with probability at least $1 - z^{-4} > 1 - 24^{-4}$ according to Lemma 16. In case we output a number, we can apply Hoeffding's inequality, since the output is the average of z trials, and obtain:

$$\Pr[|\mathcal{B} - \Pr[\mathcal{F}_t \mid \mathcal{E}_u]| \geq \frac{\epsilon}{2}] \leq 2e^{-2z\frac{\epsilon^2}{2^2}} = 2e^{-3}.$$

We see that we thus fail to output a number in $[\Pr[\mathcal{F}_t] - \epsilon, \Pr[\mathcal{F}_t] + \epsilon]$ with probability at most $2e^{-3} + 24^{-4} < 1/10$. This completes the proof. \square

Theorem 11 follows from Lemmas 17, 18, 19.

11 Algorithm for extinction of t_2

In this section we present the algorithm for extinction of t_2 . First we present an approximation of the fixation probability.

11.1 Approximating the fixation probability

We will next show that whenever r is sufficiently big, $\frac{1}{r}$ is a good approximation of the solution of the extinction problem for t_2 . To do so, we first argue that with probability at most $\frac{1}{r+1}$ does t_1 go extinct in the first effective step, when there is only a single random node which is a member of t_1 .

Lemma 20. *The probability that t_1 goes extinct in the first effective step is at most $\frac{1}{r+1}$, when initially a single random node is of type t_1 .*

Proof. The proof is similar to the proof of [4, Lemma 4], except that instead of lower bounding, we try to upper bound. For each node $v \in V$, let $Q(v) = \sum_{(u,v) \in E} \frac{1}{\deg u}$. Observe that $\sum_{v \in V} Q(v) = n$. Let p be the probability that t_1 does not go extinct in the first effective step, when initially a single random node is of type t_1 . From the proof of [4, Lemma 4], we see that $p = \frac{r}{n} \sum_{v \in V} \frac{1}{r+Q(v)}$. We want to find the minimum p such that $p = \frac{r}{n} \sum_{i=1}^n \frac{1}{r+q_i}$, subject to the constraint that $q_i > 0$ and $\sum_{i=1}^n q_i = n$.

Claim 21. *The number p is minimized for $q_i = 1$ for all i .*

Proof. We will argue that p is minimized for $q_i = 1$ for all i , as follows: Our argument is by contradiction. For a vector q of length n , let $p^q = \frac{r}{n} \sum_{i=1}^n \frac{1}{r+q_i}$. Assume that $q^* = (q_i^*)_{i \in [1, \dots, n]}$ is a vector of q_i 's that minimizes p^q and that $q_j^* \neq q_k^*$ for some j, k (this is the case if not all are 1). Let $2z = q_j^* + q_k^*$. Consider the vector q' such that $q'_i = q_i^*$ for all i , except that $q'_j = q'_k = z$. We will argue that $p^{q'} < p^{q^*}$, contradicting that q^* minimizes p^q . We have that

$$\frac{1}{r+q_j^*} + \frac{1}{r+q_k^*} = \frac{r+q_k^*+r+q_j^*}{(r+q_j^*)(r+q_k^*)} = \frac{2r+2z}{r^2+2rz+q_j^*q_k^*} > \frac{2r+2z}{r^2+2rz+z^2} = \frac{1}{r+z} + \frac{1}{r+z} ,$$

since, $q_j^*q_k^* < z^2$ and thus, since all other terms of $p^{q'}$ is equal to p^{q^*} , we have that $p^{q'} < p^{q^*}$. Thus, in the worst case $q_i = 1$ for all i . \square

Hence, $p \geq \frac{r}{r+1} = 1 - \frac{1}{r+1}$. This completes the proof. \square

Theorem 12. Let $0 < \epsilon < \frac{1}{2}$ be given. If

$$r \geq \max(\Delta^2, n)/\epsilon ,$$

then

$$\frac{1}{r} \in [(1-\epsilon) \cdot \rho, (1+\epsilon) \cdot \rho] ,$$

where ρ is the solution of the extinction problem for t_2 .

Proof. Fix $0 < \epsilon < \frac{1}{2}$ and $r \geq \max(\Delta^2, n)/\epsilon$. Let ρ be the solution of the extinction problem for t_2 . We consider two cases.

$\frac{1}{r}$ is below $(1+\epsilon) \cdot \rho$ By [4, Lemma 4] (recalled as Lemma 15), we have that $\rho \geq \frac{1}{n+r}$. Thus, $(1+\epsilon) \cdot \rho \geq \frac{1+\epsilon}{n+r} > \frac{1+\epsilon}{(1+\epsilon)r} = \frac{1}{r}$, as desired, since $n/\epsilon \leq r$.

$\frac{1}{r}$ is above $(1-\epsilon) \cdot \rho$ The probability that after the first effective step there are two nodes of type t_1 is at least $1 - \frac{1}{r+1}$, by Lemma 20. In every subsequent step before fixation, we have probability at most $\frac{1}{r/\Delta+1}$, by Lemma 3, of decreasing the number of nodes of type t_1 by 1 and otherwise we increase the number by 1.

Modeling as Markov chain Consider a biased random walk on a line with absorbing boundaries (this problem is also sometimes called Gambler's Ruin). More precisely, let M be the following Markov chain, with states $\{0, \dots, n\}$, and whenever 0 or n is reached, they will never be left (i.e., the states are absorbing). Also, in each other state $i \in \{1, \dots, n-1\}$, there is a probability of $p = \frac{1}{r/\Delta+1}$ of going to $i-1$ and otherwise, the next state is $i+1$. The fixation probability for t_2 , if fixation did not happen in the first step, is at most

the absorption probability in state 0 in M starting in state 2. The absorption probability in state 0 of M , starting in state 2, is well-known, and it is

$$\frac{\left(\frac{1-p}{p}\right)^{n-2} - 1}{\left(\frac{1-p}{p}\right)^n - 1} = \frac{\left(\frac{r}{\Delta}\right)^{n-2} - 1}{\left(\frac{r}{\Delta}\right)^n - 1} < \frac{\left(\frac{r}{\Delta}\right)^{n-2}}{\left(\frac{r}{\Delta}\right)^n} = \Delta^2/r^2 \leq \epsilon/r$$

using that $\frac{1-p}{p} = \frac{r}{\Delta}$ in the first equality, $\frac{a-1}{b-1} < \frac{a}{b}$ for all $1 < a < b$ in the first inequality and $\Delta^2/\epsilon \leq r$ in the last inequality. Hence, the probability to fixate for t_2 is below ϵ/r if t_2 does not fixate in the first step. Thus, the probability ρ of fixation for t_2 is at most $(1 + \epsilon) \cdot \frac{1}{r}$. Observe that

$$(1 - \epsilon)\rho \leq (1 - \epsilon)(1 + \epsilon) \cdot \frac{1}{r} < \frac{1}{r}, \quad \text{since } (1 - \epsilon)(1 + \epsilon) = 1 - \epsilon^2 < 1 .$$

Hence, $\frac{1}{r}$ is in $[(1 - \epsilon) \cdot \rho, (1 + \epsilon) \cdot \rho]$ as desired. \square

11.2 Algorithm

Algorithm Algo4 We consider the extinction problem for t_2 . Algorithm Algo4 is as follows:

1. If $r \geq \max(\Delta^2, n)/\epsilon$ return $\frac{1}{r}$.
2. Let \mathcal{D} be the uniform distribution over the n type functions where exactly one node is t_1 .
3. Return $\text{MetaSimulation}(t_2, z, u_{z,r}, \mathcal{D})$, for $z = 24 \frac{(n+r)^2}{\epsilon^2}$.

We next prove the following theorem.

Theorem 22. *Let G be a connected undirected graph of n nodes with highest degree Δ , divided into two types of nodes t_1, t_2 , such that $r = w(t_1) > w(t_2) = 1$. Let $\frac{1}{2} > \epsilon > 0$.*

- **Extinction problem given t_2** *Let $w = \max(\Delta^2, n)/\epsilon$ and let T be*

$$T = \begin{cases} O(1) & \text{if } r \geq w \\ O(r^2 \cdot n \cdot \Delta \cdot \epsilon^{-2} \cdot \max(\log r, \log \epsilon^{-1})) & \text{if } n \leq r < w \\ O(n^3 \cdot \Delta \cdot \epsilon^{-2} \cdot \max(\log n, \log \epsilon^{-1})) & \text{if } 2\Delta \leq r < n \\ O\left(\frac{n^3 \cdot \Delta^2}{\epsilon^2 \cdot \min(r-1, 1)} \cdot \max(\log n, \log \epsilon^{-1})\right) & \text{if } 1 + \frac{1}{n \cdot \Delta} \leq r < 2\Delta \\ O(n^4 \cdot \Delta^3 \cdot \epsilon^{-2} \cdot \max(\log n, \log \epsilon^{-1})) & \text{if } 1 < r < 1 + \frac{1}{n \cdot \Delta} . \end{cases}$$

Algorithm Algo4 is an FPRAS algorithm, with running time T , that with probability atleast $\frac{3}{4}$ outputs a number in $[(1 - \epsilon) \cdot \rho, (1 + \epsilon) \cdot \rho]$, where ρ is the solution of the extinction problem given t_2 .

| | | All steps | Effective steps |
|---------------------------------|---------|--|---|
| #steps in expectation | $r > 1$ | $O(n^2 \Delta^2)$ | $O(n\Delta)$ |
| | $r = 1$ | $O(n^3 \Delta^3)$ | $O(n^2 \Delta^2)$ |
| | $r < 1$ | $O(n\Delta^2)$ | $O(\Delta)$ |
| Concentration bounds | $r > 1$ | $\Pr[\tau \geq \frac{n^2 \Delta^2 r x}{r-1}] \leq 1/x$ | $\Pr[\tau \geq \frac{6n\Delta x}{\min(r-1,1)}] \leq 2^{-x}$ |
| | $r = 1$ | $\Pr[\tau \geq n^3 \Delta^3 x] \leq 1/x$ | $\Pr[\tau \geq 4n^2 \Delta^2 x] \leq 2^{-x}$ |
| | $r < 1$ | $\Pr[\tau \geq \frac{n\Delta^2 x}{1-r}] \leq 1/x$ | $\Pr[\tau \geq \frac{6n\Delta x}{1-r}] \leq 2^{-x}$ |
| Sampling a step | | $O(1)$ | $O(\Delta)$ |
| Fixation for t_1 | | $O(n^6 \Delta^2 \epsilon^{-4})$ | $O(n^2 \Delta^2 \epsilon^{-2} (\log n + \log \epsilon^{-1}))$ |
| Extinction for t_1 | | $O(n^6 \Delta^2 \epsilon^{-4})$ | $O(n\Delta^2 \epsilon^{-2} \log \epsilon^{-1})$ |
| Extinction for t_2 | | $O(n^5 \Delta^2 \epsilon^{-4})$ | $O(n^3 \Delta^2 \epsilon^{-2} (\log n + \log \epsilon^{-1}))$ |
| Generalized fixation (additive) | | – | $O(n\Delta^2 \epsilon^{-2} \log \epsilon^{-1})$ |

Table 2: Comparison with previous work, for constant r . We denote by n , Δ , τ , and ϵ , the number of nodes, the maximum degree, the random variable for the fixation time, and the approximation factor, respectively. The results in the column “All steps” is from [4], except that we present their dependency on Δ which was considered as n . The entry marked – is done so, since [4] did not provide an algorithm for the generalized fixation problem. The results of the column “Effective steps” is the results of this paper

Remark 6. A detailed comparison of our results and previous results of [4] is presented in Table 2. Note that we do not present approximation of the fixation problem for t_2 (i.e., fixation for the type with the smallest fitness). This is because in this case the fixation probabilities can vary greatly, while being exponentially small, even for constant r : It is close to $\frac{1-r^{-2}}{1-r^{-2n}}$ for star graphs [11] and precisely $\frac{1-r^{-1}}{1-r^{-n}}$ for complete graphs [11]. For, for instance, $r = 1/2$ it is thus close to

$$\frac{1 - 2^2}{1 - 2^{2n}} = \frac{2^2 - 1}{2^{2n} - 1} \approx 2^{-2n+2}$$

for star graphs and

$$\frac{1 - 2}{1 - 2^n} = \frac{2 - 1}{2^n - 1} = \frac{1}{2^n - 1}$$

for complete graphs. Observe that those two numbers differs by an exponentially large factor. For this case [4] also do not present any polynomial-time algorithm.

Variant of Theorem 12 for directed graphs A variant of Theorem 12 can also be proven for strongly connected (i.e. graphs where there is a path from any node to any node) directed graphs. However, the variant requires $O(m)$ computation.

In essence, the proof of Theorem 12 is as follows: The probability that t_2 fixates in 1 step is always nearly $\frac{1}{r}$. If t_2 does not fixate in 1 step (and there are thus at least 2 nodes of type t_1), then t_2 has probability at most ϵ/r of fixating.

This suggests a simple scheme for the directed case: (1) Find the probability ρ_1 that t_2 fixates in 1 step. (2) Show that fixating if there are two states of type t_2 is small compared to ρ_1 .

It is easy to find the probability that t_2 fixates in 1 step in any (even directed) graph, by simply consider each start configuration and computing the probability that fixation happens in 1 step. For a fixed configuration f with one node v of type t_1 , this takes time equal to the in- and out-degree of v (assuming that the out- and in-degree of all nodes are known). Overall this thus takes $O(m)$ operations. The probability of fixation in 1 step for a configuration is in $[\frac{1}{\Delta r+1}, \frac{n-1}{r+(n-1)}]$, recalling that Δ is the max out-degree. (This bound is easy to establish: Let v be the node of type t_1 . The bounds follow from that at least one node u of type t_2 is such that v is a successor of u (resp. u is a successor of v), because the graph is connected. On the other hand, all nodes of type t_2 might be a successor of v (or the other way around)) It seems reasonably to consider that operations on that small numbers can be done in constant time. Therefore, we get a $O(m)$ running time.

To then show that fixating for t_2 is small (compared to ρ_1) one needs a variant of Lemma 3 for directed graphs. This can easily be done, but just shows that the probability to increase the number of members of t_2 is at most $p := \frac{1}{r/(n\Delta)+1}$ (i.e. it is a factor of n worse as compared to undirected graphs). The remainder of the proof is then similar (i.e. we consider the Markov chain M on $0, 1, \dots, n$, starting in state 2, that has pr. p of decrementing and otherwise increments). We get that the pr. of fixating for t_2 if it did not do it in the first step is at most $\epsilon/(2\Delta r) < \epsilon/(\Delta r + 1)$ if $r \geq 2\Delta^3 n^2/\epsilon$. Hence, $\rho_1 \in [(1 - \epsilon) \cdot \rho, (1 + \epsilon) \cdot \rho]$ for such r . This leads to the following theorem.

Theorem 23. *Let $\epsilon > 0$ be given. Let G be a directed graph and let $r \geq 2\Delta^3 n^2/\epsilon$.*

Consider the extinction problem for t_2 . Let ρ be the fixation pr. and let ρ_1 be the pr. of fixation in 1 effective step. Then

$$\rho_1 \in [(1 - \epsilon) \cdot \rho, (1 + \epsilon) \cdot \rho] .$$