

Population Genetics Models

Model the forces that produce and maintain genetic evolution within a population.

Mutation: the process by which one individual (gene) changes. Simulation wants to study the drift of the population: how the frequency of mutants in the total population evolves.

The Moran Process P. Moran: Random processes in genetics Cambridge Ph. Soc. 1958

- Start with *n* individuals. Randomly select one to mutate.
- Select randomly an individual x to replicate.
- Select randomly another y to die.
- Replace y by a clone of x.

Stochastic process. At time t the number mutants evolves in $\{-1, 0, +1\}$.



Evolutionary graph theory (EGT)

Lieberman, Hauert, Nowak: *Evolutionary dynamics on graphs* Nature 2005 (LHN)

EGT studies how the topology of interactions between the population affects evolution.

Graphs have two types of vertices: mutants and non-mutants.

The fitness r of an agent denotes its reproductive rate. Mutants have fitness $r \in \Theta(1)$, non-mutants have fitness 1.

Mutants and non-mutants extend by cloning one of their neighbors.

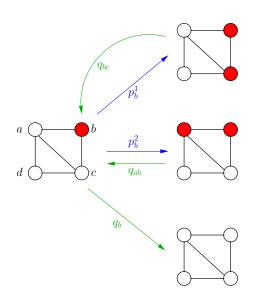
Moran process on Evolutionary Graphs

Given a graph G = (V, E), with |V| = n, and an r > 0, we start with all vertices non-mutant.

- at t=0 create uniformly at random a mutant in VAt any time t>0, assume we have k mutant and (n-k)non-mutant vertices. Define total fitness at time t by $W_t=kr+(n-k)$:
- Choose u with probability $\frac{r}{W_t}$ if u is mutant and $\frac{1}{W_t}$ otherwise,
- ullet choose uniformly at random a $v \in \mathcal{N}(u)$, and replace v with the clone of u

The process is Markovian, depending on r it tends to one of the two absorbing states: extinction or fixation.

Example of Moran process

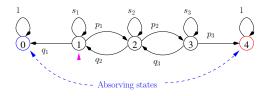


where:

$$\begin{split} p_b^1 &= \frac{r}{3+r} \cdot \frac{1}{2} \\ p_b^2 &= \frac{r}{3+r} \cdot \frac{1}{2} \\ q_{ab} &= \frac{1}{2+2r} \cdot \frac{5}{6} \\ q_{bc} &= \frac{1}{(n-1)+r} \cdot \frac{5}{6} \\ q_b &= \frac{2}{3+r} \cdot \frac{1}{3} \end{split}$$

Moran Process

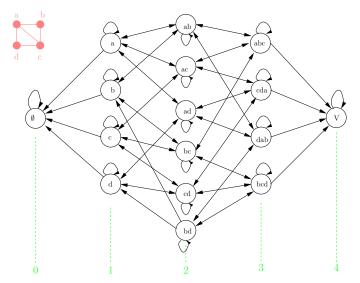
This random process defines discrete, transient Markov chain, on states $\{0, 1, ..., n-1, n\}$ with two absorbing states: n fixation (all mutant) and n0 extinction (all non-mutant).



The fixation probability $f_G(r)$ of G is the probability that a single mutant will takes over the whole G. The extinction probability of G is $1 - f_G(r)$.

The Markov chain of configurations

A configuration is a set $S \subseteq V$ of mutants.



Properties of $f_G(r)$

Given G = (V, E) connected and a fitness r > 0, for any $S \subset V$ let $f_{G,r}(S)$ denote the fixation probability, when starting with a set S of mutants.

Notice $f_G(r) = \sum_{v \in V} f_{G,r}(\{v\})$.

The case r = 1 is denoted neutral drift.

Shakarian, Ross, Johnson, Biosystems 2012 For any $r \ge 1$, $f_G(r) \ge f_G(1)$

Díaz, Goldberg, Mertzios, Richerby, Serna, Spirakis, SODA-2012 (DGMRSS)

For any undirected G = (V, E), $f_G(1) = \frac{1}{n}$.

Bounding $f_G(r)$

Let G = (V, E) be any undirected connected graph, with |V| = n.

(DGMRSS)

For any $r \ge 1$, $\frac{1}{n} \le f_G(r) \le 1 - \frac{1}{n+r}$, are bounds on the fixation probability for G.

Merzios, Spirakis: ArXive-2014

For any $\epsilon > 0$,

$$f_G(r) \leq 1 - \frac{1}{n^{\frac{3}{4} + \epsilon}}.$$

Questions to study

Given a connected graph G = (V, E) (strongly connected is case of digraphs), and a fitness r:

1.- Is it possible to compute exactly the fixation probability $f_G(r)$?

Difficult for some graphs. For a given G the number of constrains and variables is equal to the number of possible configurations of mutants/non-mutants in $G \sim 2^n$.

2.- Given G, is it possible to compute the expected number of steps until arriving to absorption?



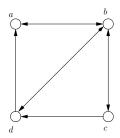
Isothermal graphs (LHN)

Given a directed $\vec{G} = (V, \vec{E}), \forall i \in V \text{ let } \deg^+(i)$ be its outgoing degree:

Define the stochastic matrix $W = [w_{ij}]$, where $w_{ij} = 1/\deg^+(i)$ if $(\vec{i}, \vec{i}) \in \vec{E}$ and $w_{ii} = 0$ otherwise.

The same definition of W applies to undirected G, with $w_{ii} = 1/\deg(i)$.

The temperature of $i \in V$ is $T_i = \sum_{i \in V} w_{ji}$ A graph \vec{G} is isothermal if $\forall i, j \in V$, $T_i = T_i$.

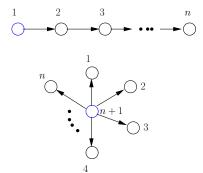


$$W = \left(\begin{array}{cccc} 0 & 1 & 0 & 0 \\ 1/3 & 0 & 1/3 & 1/3 \\ 0 & 1/2 & 0 & 1/2 \\ 1/2 & 1/2 & 0 & 0 \end{array}\right)$$

$$T_b = 2$$
 and $T_c = 1/3$

Computing the fixation probability

If \vec{G} is a digraph with a single source then $f_{\vec{G}}(r) = \frac{1}{n}$.



Isothermal Theorem (LHN)

For a strongly connected graph \vec{G} s.t. $\forall i,j \in V$ we have $T_i = T_j$ (i.e. W is bi-stochastic) then

$$f_{\vec{G}}(r) = \frac{1 - \frac{1}{r}}{1 - \frac{1}{r^n}} \equiv \rho$$

Undirected graphs

The isothermal theorem also applies to undirected graphs. Given G undirected and connected, then G is Δ -regular iff W is bi-stochastic.

If G is undirected and connected then $f_G(r) = \rho = \frac{1-1/r}{1-1/r^n}$ iff G is Δ -regular.

For example, if G is C_n or K_n then $f_G(r) = \rho$.

Notice:

- if r > 1 then $\lim_{n \to \infty} f_G(r) = 1 \frac{1}{r}$.
- if r < 1 then $f_G(r) = \frac{r^n r^{n-1}}{r^n 1} \to$ exponentially small.

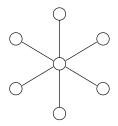
Amplifiers and suppressors

Given G (directed or undirected) and r, G is said to be an amplifier if $f_G(r) > \rho$. G is said to be a suppressor if $f_G(r) < \rho$.

The star

(LHN), (Broom, Rychtá. Proc.R. Soc. A 2008)

For
$$r > 1$$
 $f_G(r) = \frac{1 - \frac{1}{r^2}}{1 - \frac{1}{r^{2n}}} > \rho$



The star is an amplifier

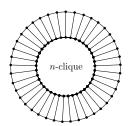
Suppressors

The directed line and the burst have fixation probability $\frac{1}{n} < \rho$, therefore they are examples of suppressors.

How about non-directed graphs as suppressors? Mertzios, Nikoletseas, Ratopoulos, Spirakis, TCS 2013 **The urchin**

For
$$< r < 4/3$$

 $\lim_{n \to \infty} f_G(r) = \frac{1}{2}(1 - \frac{1}{r}) < \rho$



The urchin is an undirected graph suppressor

Absorption time for undirected graphs

Given undirected connected G = (V, E), with |V| = n, run a Moran process $\{S_t\}_{t>0}$, where $\{S_t\}$ set of mutants at time t.

Define the absorption time $\tau = \min\{t \mid S_t = \emptyset \lor S_t = V\}.$

Theorem DGMRSS

Given G undirected, for the Moran process $\{S_t\}$ starting with $|S_1|=1$:

- 1. If r < 1, then $\mathbf{E}[\tau] \le \frac{r}{r-1} n^3$,
- 2. if r > 1, then $\mathbf{E}[\tau] \le \frac{r}{r-1} n^4$,
- 3. if r = 1, then **E** $[\tau] \le n^6$.

Sketch of the proof

We bound $\mathbf{E}[\tau]$ using a potential function that decreases in expectation until absorption.

Define the potential function $\phi(S) = \sum_{v \in S} \frac{1}{\deg(v)}$ Notice $\phi(\lbrace v \rbrace) \geq 1/n$ and $0 \leq \phi(S_\tau) \leq n$

Use the following result from MC (Hajek, Adv Appl. Prob. 1983)

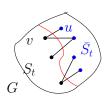
If $\{X_t\}_{t\geq 0}$ is a MC with state space Ω and there exist constants $k_1, k_2 > 0$ and a $\phi: \Omega \to \mathbb{R}^+ \cup \{0\}$ s.t.

- (1) $\phi(S) = 0$, $\exists S \in \Omega$,
- (2) $\phi(S) \leq k_1$,
- (3) $\mathbf{E} [\phi(X_t) \phi(X_{t+1}) \mid X_t = S] \ge k_2, \forall t \ge 0 \text{ s.t. } \phi(S) > 0,$ then $\mathbf{E} [\tau] \le k_1/k_2$, where $\tau = \min\{t \mid \phi(S) = 0\}.$

Sketch of the proof

To compute evolution of $\mathbf{E} [\phi(S_{t+1}) - \phi(S_t)].$

To show that the potential decreases (increases) monotonically for r < 1 (r > 1), consider the contribution of each (u,v) in the cut for $S_{t+1} = S_t \cup \{v\}$ and to $S_{t+1} = S_t \setminus \{v\}$.



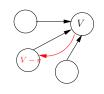
- 1. For r < 1, $\mathbf{E} \left[\phi(S_{t+1}) \phi(S_t) \right] < \frac{r-1}{n^3} < 0$.
- 2. For r > 1, $\mathbf{E}\left[\phi(S_{t+1}) \phi(S_t)\right] \ge (1 \frac{1}{r})\frac{1}{n^3}$.
- 3. For r = 1, $\mathbf{E} \left[\phi(S_{t+1}) \phi(S_t) \right] = 0$.

Domination argument for r < 1

For any fixed initial $S \subset V$:

Let $\{Y_i\}_{i\geq 0}$ be a stochastic process as Moran's, except if it arrives to state V, u.a.r. choose v and exit to state $V\setminus\{v\}$.

Let
$$\tau' = \min\{i \mid Y_i = \emptyset\}$$



Then,

$$\mathbf{E}[\tau | X_0 = S] \le \mathbf{E}[\tau' | Y_0 = S] \le \frac{1}{1-r} n^3 \phi(S)$$

$$\Rightarrow \mathbf{E}[\tau] \leq \frac{1}{1-r} n^3$$
.

Domination argument for r > 1

For any fixed initial $S \subset V$: Define a process $\{Y_i\}_{i \geq 0}$ as in Moran's, except if arrives to state \emptyset , u.a.r. choose v and exit to state $\{v\}$.



Let
$$\tau' = \min\{i \mid Y_i = V\}$$

Then,

$$\mathbf{E}\left[\tau|X_0=S\right] \leq \mathbf{E}\left[\tau'|Y_0=S\right] \leq \frac{rn^3}{r-1}(\phi(G)-\phi(S))$$

$$\Rightarrow \mathbf{E}[\tau] \leq \frac{r}{r-1}n^4$$
.

Proof for r = 1

For undirected G = (V, E) with r = 1,

$$\mathbf{E}\left[\tau\right] \leq \phi(V)^2 n^4 \leq n^6.$$

In this case $\mathbf{E} \left[\phi(S_t) - \phi(S_{t-1}) \right]$ does not change \Rightarrow Use a martingale argument

At each t, the probability that ϕ changes is $\geq 1/n^2$, and it changes by $\leq 1/n$.

Dominate by process $Z_t(\phi_t)$, which increases in expectation until stopping time, when the process absorbs.

Then $\mathbf{E}[Z_{\tau}] \geq \mathbf{E}[Z_0]$ and we get a bound for $\mathbf{E}[\tau]$.

Approximating $f_G(r)$

A FPRAS for a function f: A randomized algorithm A such that, given a $0 \le \epsilon \le 1$, for any input x,

$$\Pr\left[(1-\epsilon)f(x) \le A(x) \le (1+\epsilon)f(x)\right] \ge \frac{3}{4},$$

with a running time $\leq \text{poly}(|x|, 1/\epsilon)$.

Corollary to absorption bounds

- ▶ There is an FPRAS for computing the fixation probability, for any fixed $r \ge 1$.
- ▶ There is an FPRAS for computing the extinction probability, for any fixed r < 1.

Absorption time Δ -regular graphs, r > 1

Díaz, Goldberg, Richerby, Serna. ArXive 2014 Recall the upper bound for absorption time undirected G is $\frac{r}{r-1}n^4$.

Theorem If G = (V, E) is a connected Δ -regular graph with |V| = n, the upper bound to the expected absorption time is

$$\mathbf{E}\left[\tau\right] \leq \frac{r}{r-1} n^2 \Delta.$$

Sketch of proof For any $\emptyset \subseteq S \subseteq V$, use $\phi(S) = \sum_{v \in S} \frac{1}{\deg(v)} = \frac{|S|}{\Delta}$ and $\phi(V) = \frac{n}{\Delta}$.

$$\mathbf{E}\left[\phi(S_{t+1}) - \phi(S_t)\right] = \frac{r-1}{W_{t+1}} \frac{1}{\deg(u)\deg(v)} = \Theta(\frac{1}{\Delta^2 n})$$

△-regular digraphs

 Δ -regular digraph: $\forall v, \deg^-(v) = \deg^+(v) = \Delta$.

Recall for regular digraphs:

- Fixation probability is ρ , independent of the particular topology of the graph.
- As $n \to \infty$, $\rho \to 1 \frac{1}{r}$, therefore the expected number of active steps $\to n(1 \frac{1}{r})$, independently of the graph.



Expected absorption time for regular digraphs, r > 1

The expected absorption time does depend on the graph.

Theorem Let G be a strongly connected Δ -regular n-vertex digraph. Then the expected absorption time is

$$\left(\frac{r-1}{r^2}\right)nH_{n-1}\leq \mathbf{E}\left[\tau\right]\leq n^2\Delta,$$

where H_n is the nth. Harmonic number.

Corollaries

- For K_n ($\Delta = n 1$) \Rightarrow **E**[τ] = $\Omega(n \log n)$ and **E**[τ] = $O(n^3)$.
- For $C_n \Rightarrow \mathbf{E}[\tau] = \Omega(n \log n)$ and $\mathbf{E}[\tau] = O(n^2)$.

Undirected Δ -regular and isoperimetric inequality

Given an undirected graph G = (V, E), the isoperimetric number (Harper, J. Comb. Theory 1966) is defined as

$$i(G) = \min_{S} \left\{ \frac{|\delta S|}{S} \mid S \subset V, 0 < |S| \le |V|/2 \right\},$$

where δS is the set of edges in the cut between S and $V \setminus S$.

Proposition If G is Δ -regular undirected (good expander)

$$\mathbf{E}\left[\tau\right] \leq \frac{2\Delta n H_n}{i(G)}.$$

For some Δ -reg. G the isoperimetric bound improves the general theorem.

Applications of the isoperimetric result

- The K_n has $i(G) = \Theta(1/\sqrt{n}) \Rightarrow$ $\mathbf{E}[\tau] = \Theta(n \log n)$ $(\mathbf{E}[\tau] = O(n^3)).$
- The $\sqrt{n} \times \sqrt{n}$ -grid has $i(G) = \Theta(1/\sqrt{n}) \Rightarrow$ $\mathbf{E}[\tau] = O(n^{3/2} \log n)$ ($\mathbf{E}[\tau] = O(n^2)$).
- The C_n has $i(G) = 4/n \Rightarrow$ $\mathbf{E}[\tau] = O(n^2 \log n)$ ($\mathbf{E}[\tau] = O(n^2)$).

Bolobás, Eur. J. Comb. 1988: For $\Delta \geq 3$ there is a number $0 < \nu < 1$ such that, as $n \to \infty$, for almost all undirected Δ -regular G, $i(G) = \nu \Delta/2$.

• Bollobás result \Rightarrow for almost all undirected Δ -regular G, $\mathbf{E}[\tau] = O(n \log n)$.

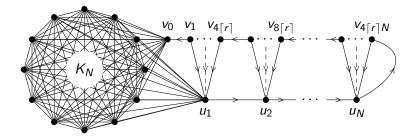


Worst absorption time for directed graphs

Recall the absorption time of undirected graphs $\mathbf{E}[\tau] \leq O(n^4)$.

Theorem There is an infinite family of strongly connected digraphs such that the expected absorption time for an *n* vertex graph is

$$\mathbf{E}\left[\tau\right]=2^{\Omega(n)}.$$



Domination

Given a Moran's process $\{X_t\}$ on G, intuition says that for any S and any $S' \subset S$, $f_S(r) > f_{S'}(r)$ and $\tau(S) < \tau(S')$.

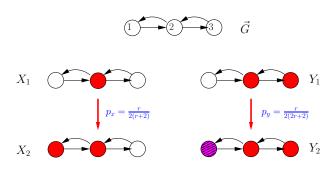
 \therefore To analyze $\{X_t\}$, we can couple it with a process $\{Y_t\}$, which is easier to analyze (for instance by allowing transitions that create new mutants but forbidding some of the transitions removing mutants).

Then we must ensure that for every t > 1, if $X_1 \subseteq Y_1 \Rightarrow X_t \subseteq Y_t$.

NOT ALWAYS TRUE for discrete Moran's processes



Counterexample



Coupling $\{X_i\}$ and $\{Y_i\}$ fails as for r > 1,

$$\Pr\left[X_2 \not\subseteq Y_2\right] > 0$$

Continuous time process

To use domination for the discrete processes $\{X_i\}$ and $\{Y_i\}$, consider the continuous versions $\tilde{X}[t]$ and $\tilde{Y}[t]$, where vertex v with fitness $r_v \in \{1, r\}$ waits an amount of time which follows an exponential distribution with parameter r_v .

The discrete Moran process is recovered by taking the sequence of configurations each time a vertex reproduces.

Notice: in continuous time, each v reproduces at a rate given by r_v , independently of the other vertices, while in discrete time the population "coordinates" before deciding who is next to reproduce.

Coupling Lemma and consequences

Coupling Lemma For $\vec{G}=(V,\vec{E})$, let $X\subseteq Y$ and $1\leq r\leq r'$. Let $\tilde{X}[t]$ and $\tilde{Y}[t]$ $(t\geq 0)$ be the continuous-time Moran process on G with mutant fitness r and r', and with $\tilde{X}[0]=X$ and $\tilde{Y}[0]=Y$. There is a coupling between the two processes s. t. $\tilde{X}[t]\subseteq \tilde{Y}[t]$, $\forall t\geq 0$.

Theorem For any \vec{G} , if $0 < r \le r'$ and $S \subseteq S'$ then

$$f_{\vec{G},r}(S) \leq f_{\vec{G},r'}(S').$$

Corollary (Monotonicity)

For any \vec{G} and $0 < r \le r'$ then, $f_{\vec{G}}(r) \le f_{\vec{G}}(r')$.

Corollary (Subset domination)

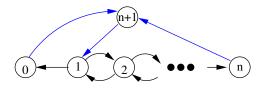
For any \vec{G} and 0 < r then, if $S \subseteq S'$ then $f_{\vec{G,r}}(S) \le f_{\vec{G,r'}}(S')$.



Glimpse of proof for

$$\left(\frac{r-1}{r^2}\right)nH_{n-1}\leq \mathbf{E}\left[\tau\right]\leq n^2\Delta,$$

Dominate the process by a Markov chain:



Solve difference equation to find the expected number of active steps going from state j to state n+1.

Compute bound on the time you spend in each state j.

Thank you for your attention