

### 13. Selection: How do populations stop growing?

#### Modelling replication

We have seen that replication is represented by the model:

$$\dot{x} = r x$$

where  $x$  is the size of a population and  $r$  is the specific growth rate of that population. This model generates the exponential growth story, for which we can formulate an exact model:

$$x(t) = x_0 e^{rt}, \text{ with doubling time } T_2 = \ln(2)/r.$$

- ? A bacteria population has  $r = 0.035 \text{ min}^{-1}$ . Calculate the population's doubling time.
- ? How many minutes are in a day? How many cells does 1 bacterium generate in 3 days?

This number is enormous. In fact, it is so enormous that it cannot be true! *There is no such thing as exponential growth in real life.* Rather, limited resources cause the population growth rate to drop as the population gets bigger. This is modelled by the logistic model:

$$\dot{x} = rx (1 - x/K)$$

Here,  $r$  is the specific replication rate of the population only when  $x$  is much smaller than the resource limitation (carrying capacity)  $K$ . If  $x \rightarrow 0$ , or if  $x \rightarrow K$ ,  $\dot{x} \rightarrow 0$ , so the population has an *unstable* fixed point at  $x^* = 0$ , but grows from any initial value  $x_0 > 0$  towards the *stable* fixed point at  $x^* = K$ . (A superscript asterisk denotes a fixed-point value.)

#### Modelling selection

Suppose we have two exponential populations  $x$  and  $y$  that reproduce at different rates  $r$  and  $s$ . Suppose they have initial conditions  $x(0) = x_0$ ,  $y(0) = y_0$ , then:

$$\begin{cases} \dot{x} = r x \\ \dot{y} = s y \end{cases} \Rightarrow \begin{cases} x(t) = x_0 e^{rt} \\ y(t) = y_0 e^{st} \end{cases}$$

Both  $x$  and  $y$  grow exponentially.  $x$  has doubling time  $\ln 2/r$  and  $y$  has doubling time  $\ln 2/s$ , so if  $r > s$ ,  $x$  will grow faster than  $y$ . Eventually, there will be more  $x$ 's than  $y$ 's.

- ? Define  $\rho(t) \equiv \frac{x(t)}{y(t)}$ . Use the quotient rule to prove that  $\dot{\rho} = (r - s)\rho$ .

The solution of this equation is  $\rho(t) = \rho_0 e^{(r-s)t}$ , so if  $r > s$ ,  $\rho$  will grow toward infinity, and  $x$  *outcompetes*  $y$ . If in addition we assume resource are limited, the total population  $x + y$  will remain constant, so if  $x$  gets infinitely bigger than  $y$ , this must mean that  $y \rightarrow 0$ .

This is *selection*: where the growth of  $x$  drives  $y$  to extinction. For selection to happen, we need different rates of growth of the populations  $x$  and  $y$ , *plus* resource limitation.

To study selection situations, we often use two simple modelling tricks:

- We think of  $x$  and  $y$  not as populations, but as *frequencies*. That is, we assume the sum of both population types is 1 ( $x + y = 1$ ), so that  $x$  describes *what proportion* of the combined population are  $x$ -individuals, and  $y$  describes what proportion are  $y$ .
- In addition, we think of the growth rates  $r$  and  $s$  as *fitness* values:  $r$  describes how fit the type  $x$  is, in terms of how effectively it grows by comparison with  $y$ .
- ? We want to make sure that the sum  $x + y = 1$  of the two frequencies stays constant. To do this, we reduce the growth rates of  $x$  and  $y$  by equal amounts  $R$  in the selection equations:  $\dot{x} = (r - R)x$  and  $\dot{y} = (s - R)y$ . Prove that this is only possible if  $R$  is the average fitness of the two population types:  $R = rx + sy$ .

- ? One advantage of this selection model is that  $y$  depends upon  $x$ :  $y = 1 - x$ . Show how we can eliminate  $y$  from the two selection equations, so that we only need to solve the single equation:  $\dot{x} = (r - s)x(1 - x)$ .

We know this equation: it is the logistic equation with specific growth rate  $(r - s)$  and carrying capacity 1. We also know how the logistic story evolves over time – it has two equilibria at 0 and 1:

- If  $r > s$ ,  $x \rightarrow 1$ , so  $y \rightarrow 0$ , and type  $x$  is selected over type  $y$ ;
- If  $s > r$ ,  $x \rightarrow 0$ , so  $y \rightarrow 1$ , and type  $y$  is selected over type  $x$ ;

Martin Nowak calls this situation “*Survival of the Fitter*”.

### Survival of the fittest

We can extend this 2-type model to selection between  $n$  different types in a population. If we name the individual type frequencies  $x_i(t)$  (where  $i = 1, \dots, n$ ), the structure describing all  $n$  types is a vector:  $\mathbf{x} \equiv (x_1, x_2, \dots, x_n)$ . Now define  $r_i \geq 0$  as the fitness of type  $i$ , then the average fitness of the entire population of  $n$  types is:

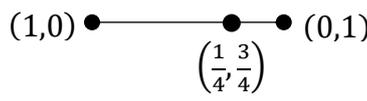
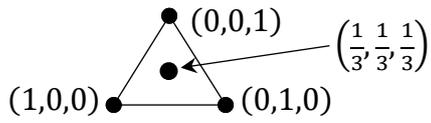
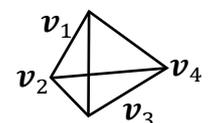
$$R = \sum_{i=1}^n x_i r_i = \mathbf{x} \cdot \mathbf{r}$$

We can then write the selection dynamics model as:

$$\dot{x}_i = x_i(r_i - R) \quad \text{(Linear selection model)}$$

The frequency  $x_i$  of type  $i$  *increases* if its fitness  $r_i$  is higher than the population average  $R$ ; otherwise  $x_i$  *decreases*. However, the total population stays constant:  $\sum_{i=1}^n x_i = 1$  and  $\sum_{i=1}^n \dot{x}_i = 0$ . This is useful if we want to study the rise and fall of types within a population.

The set of all values  $x_i > 0$  obeying the property that  $\sum_{i=1}^n x_i = 1$  is called a *simplex* (denoted  $S_n$ ). The useful thing about simplexes is that we can represent them graphically:

$n$	Simplex $S_n$	Geometrical visualisation	
1	Point		
2	Line segment		
3	Triangle		
4	Tetrahedron		<p>If <math>\mathbf{v}_i</math> (<math>i = 1, 2, 3, 4</math>) are four vertex position vectors, the general point of <math>S_4</math> is the <i>convex combination</i>: <math>\mathbf{x} \equiv x_1 \mathbf{v}_1 + x_2 \mathbf{v}_2 + x_3 \mathbf{v}_3 + x_4 \mathbf{v}_4</math></p>

For example, consider the 3-simplex (or triangle)  $S_3$ . Here, we interpret the top point  $(0,0,1)$  as representing the situation in which only population type 3 is present, and the other two are not. On the other hand, we interpret the centre point  $(\frac{1}{3}, \frac{1}{3}, \frac{1}{3})$  as the situation where all three types are present in equal quantities.

- ? Which point would represent the situation in which type 2 is absent, and types 1 and 3 are present in equal quantities?
- ? In the linear selection model above, imagine that type  $k \in \{1, 2, \dots, n\}$  has greater fitness than any other type:  $r_k > r_i, \forall i \neq k$ . What does this mean for the value of the factor  $(r_i - R)$ ? What does this mean for the growth rate  $\dot{x}_k$  of type  $k$  whenever other types are

present? What will be the frequency of the types after a long time? What will happen to any interior point of the simplex  $S_n$  over time?

You have demonstrated that the exponential selection model only ever has one outcome: total competitive exclusion. This is the meaning of the phrase “*Survival of the Fittest*”.

### Exercise project (1 week)

In this project we will build a slightly more general model of selection:

$$\dot{x}_i = r_i x_i^c - R x_i; \quad R = \sum_{i=1}^n r_i x_i^c; \quad c < 1 \quad \text{(Sublinear selection model)}$$

$$\dot{x}_i = r_i x_i^c - R x_i; \quad R = \sum_{i=1}^n r_i x_i^c; \quad c > 1 \quad \text{(Superlinear selection model)}$$

1. Notice that when  $c = 1$ , these equations reduce to the exponentially growing linear selection model. If  $c < 1$ , the population growth is slower than exponential (*subexponential*), and if  $c > 1$ , growth is faster than exponential (*superexponential*). An extreme example of subexponential growth is immigration at a constant rate. An example of superexponential growth is sexual reproduction, where two organisms (perhaps male and female) must meet in order to replicate.
2. Let's take the simple case  $n = 3$ . Show that in this case, if the population lies in the simplex  $S_3$  (so  $x_1 + x_2 + x_3 = 1$ ), then the rate of change ( $\dot{x}_1 + \dot{x}_2 + \dot{x}_3$ ) of the entire population is equal to zero. What does this imply for evolution in relation to  $S_3$ ?
3. Design a Matlab class `Selection` that uses RK2 to simulate the evolution of a population of three types. Your client function should use the class constructor to set the values of  $c$  and the three specific growth rates, then call the method `simulate([x0 y0 z0], T)` to evolve the population over a time  $T$ , starting from the initial frequencies  $[x0 \ y0 \ z0]$ , and plot this evolution graphically within the triangular simplex  $S_3$ :

```
sel = Selection(1.2, [0.2, 0.3, 0.4]); sel.simulate([0.3, 0.3, 0.4], 20);
```

4. Use your selection class to show that  $c < 1$  leads to *Survival of All*, while  $c > 1$  leads to *Survival of the First*, and present this work in a poster.

### Summary

- Charles Darwin and Alfred Russell Wallace realised in 1858 that all resources are limited, which *necessarily* leads to selection and prevents exponential growth.
- The *linear selection* model is  $\dot{x}_i = x_i(r_i - R)$ , where  $x_i$  and  $r_i$  are the *frequency* and *specific replication rate*, or *fitness*, of population type  $i$ ;  $R = \sum_{i=1}^n x_i r_i = \mathbf{x} \cdot \mathbf{r}$  is the average fitness of the population; and  $\sum_{i=1}^n x_i = 1$ .
- The condition  $\sum_{i=1}^n x_i = 1$  means that a population in the linear selection model is represented by a point moving over time within a *simplex*  $S_n$  whose  $k$ -th vertex represents the presence of only the single population type  $k \in \{1, 2, \dots, n\}$ .
- Linear selection always leads to *Survival of the Fittest*: the movement of the population from any interior point of  $S_n$  to the vertex  $k$  whose fitness is highest.
- *Sublinear* selection ( $\dot{x}_i = r_i x_i^c - R x_i$ , where  $R = \sum_{i=1}^n r_i x_i^c$  and  $c < 1$ ) models subexponential growth such as immigration; it leads to *Survival of All*.
- *Superlinear* selection ( $\dot{x}_i = r_i x_i^c - R x_i$ , where  $R = \sum_{i=1}^n r_i x_i^c$  and  $c > 1$ ) models superexponential growth such as sexual replication; it leads to *Survival of the First*.