## 1 Stability of equilibrium points & bifurcations

## 1.1 Simple population model

The population model has in general two solutions (and hence two fixed points) for  $\dot{N}=0$  , namely

$$N_1 = 0$$
 and  $N_2 = K \frac{\alpha - \beta}{\alpha}$ .

The stability of these fixed points in function of  $\alpha$  and  $\beta$  can be summarised as follows:

Parameter region	Fixed points
$\alpha < \beta$	$N_1=0$ : stable $N_2<0$ : unstable
$\alpha = \beta$	$N_1=N_2=0$ : half-stable (unstable for $N<0$ , stable for $N>0$ )
$\alpha > \beta$	$N_1=0$ : unstable $N_2>0$ : stable

The system thus undergoes a transcritical bifurcation at  $\alpha=\beta$ . Note that the fixed point  $N_2<0$  is not meaningful in this model, as N represents a non-negative population count.

For the given parameter values,  $\alpha>\beta$ . Using the above results, we therefore find an unstable fixed point  $N_1=0$ , and a stable fixed point  $N_2=K(\alpha-\beta)/\alpha=4\,023\,913$ . As the population starts at N>0, it will evolve towards  $N_2$ . The difference between N(t) and  $N(\infty)=N_2$  decays exponentially, as a Taylor approximation of  $\dot{N}$  around  $N_2$  can show.

## 1.2 Gene control model

For r = 0, the system equations become decoupled:

$$\dot{x} = \frac{\alpha_1}{2} - x$$

$$\dot{y} = \frac{\alpha_2}{2} - y.$$

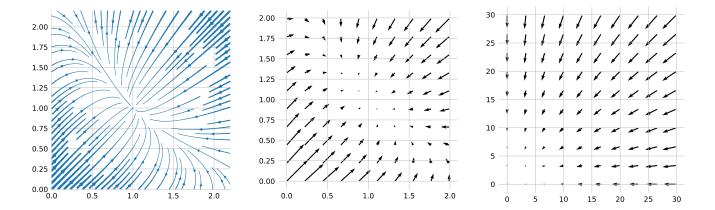


Figure 1: There is only one fixed point for  $0 \le r \le 2$ . Phase space plots of the gene control model, for r=1 and  $\alpha_1=\alpha_2=2$ . Left: some (partial) trajectories in phase space. Thicker lines represent a higher local speed. Note the attractor at (1,1). Middle: local velocities, evaluated on a grid. Right: same as middle, but for a larger region in phase space.

We can therefore analyse them separately. It is clear that there is one fixed point, at  $x^* = \alpha_1/2$  and  $y^* = \alpha_2/2$ . It is a globally stable attractor, as  $\forall x < x^*, \ \dot{x} > 0$  and  $\forall x > x^*, \ \dot{x} < 0$  (and analogously for  $\dot{y}$  and  $y^*$ ). The fixed point is thus an attracting star.

For  $r \ge 0$  and  $\alpha_1 = \alpha_2 = 2$ , the equilibrium equations become

$$x(1+y^r) = 2$$
$$y(1+x^r) = 2.$$

It is easily verified that (1,1) is a solution and hence a fixed point. We have already shown that it is the only fixed point for r=0. Plotting the gradient in phase space for different  $r\in(0,2)$  strongly suggests that it is also the only fixed point for nonzero r<2 (at least for  $x\geq 0$  and  $y\geq 0$ ). See e.g. fig. 1 for r=1. As a final piece of evidence, different trajectories simulated back in time all either tend towards  $(\infty,\infty)$  or cross into forbidden  $x<0,\ y<0$  territory.

To analyse the stability of the (1,1) fixed point, we approximate  $(\dot{x},\dot{y})$  as a linear system around this point. The Jacobian of  $(\dot{x},\dot{y})$  evaluated in (1,1) is:

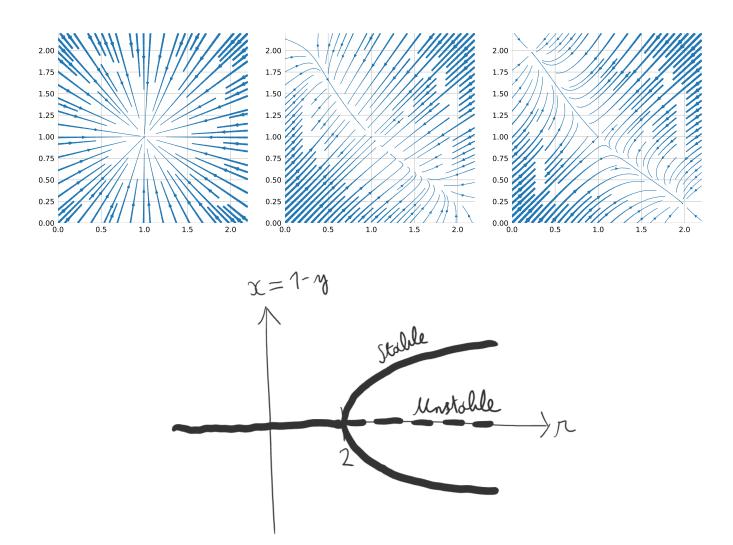
$$\begin{pmatrix} -1 & -\frac{r}{2} \\ -\frac{r}{2} & 1 \end{pmatrix}.$$

It has two distinct eigenvalue-eigenvector pairs:

$$\lambda_1 = -\frac{r}{2} - 1, \quad \mathbf{v}_1 = (1, 1)$$

$$\lambda_2 = \frac{r}{2} - 1, \quad \mathbf{v}_2 = (-1, 1).$$

For  $0 \le r < 2$ ,  $\lambda_1 \in (-2, -1]$  and  $\lambda_2 \in [-1, 0)$ . Both eigenvalues are negative, and (1, 1) is therefore a stable node. Because  $\lambda_1 < \lambda_2$ ,  $\mathbf{v}_1 = (1, 1)$  is the fast eigendirection



and  $\mathbf{v}_2 = (-1, 1)$  is the slow eigendirection, as can be seen in fig. 1. (For r = 0, both eigenvalues are equal; i.e. (1,1) is then a star node).

Based on phase space plots for different values of r, it seems that a supercritical pitchfork bifurcation occurs at r=2. This is a plausible bifurcation given the system that is modelled: when the mutual repression rate r is high, a slight abundance of one gene (say x) over the other will create a positive feedback loop: x represses y more than y represses x, resulting in more x and less y. This again results in even more x and even less y, etcetera. Vice versa for an initial slight abundance of y. There is thus a precarious balance when x=y (the unstable fixed point at (1,1) for r>2). On the other hand, when the mutual repression rate r is low, both genes can be expressed in equal amounts. (For large r, this system might be called a winner-takes-all model, or a competitive model).

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