

1 Calculation of the maximum probability of survival of a single cell

2 Consider an individual C cell placed in a location with no resource. We want to estimate $\Pr(\varepsilon)$, the
3 minimal probability that the entire population goes extinct when its release rate is very high. In general,
4 this will be the probability that the cell does not divide and goes extinct, plus the probability that the
5 cell divides once and both cells go extinct, etc. Let b be the number of births before population
6 extinction, then

$$\Pr(\varepsilon) = \Pr(\varepsilon|b=0) + \Pr(\varepsilon|b=1) + \dots + \Pr(\varepsilon|b=n). \quad (1)$$

7 As each term becomes less and less likely to occur, we calculate a lower bound on this probability by
8 only considering the first two terms.

9 In the first time step Δt , the cell will see no resource, and $\Pr(\varepsilon)$ is simply the probability of death, p_d . If
10 this cell survives, suppose that it has a probability p_b of giving birth in the next time step. Then in the
11 next time step, the probability of death before cell division becomes $(1-p_d)(1-p_b)p_d$. Continuing, we get

$$\Pr(\varepsilon|b=0) = p_d \left[1 + [(1-p_d)(1-p_b)]^1 + [(1-p_d)(1-p_b)]^2 + \dots + [(1-p_d)(1-p_b)]^i \right]. \quad (2)$$

12 Rearranging gives

$$\Pr(\varepsilon|b=0) = p_d \sum_0^{\infty} [(1-p_d)(1-p_b)]^i = \frac{p_d}{p_d + p_b - p_b p_d}. \quad (3)$$

13 The maximum birth rate was 0.45 hr^{-1} , the death rate was 0.1 hr^{-1} , and the time step was 0.005. Thus, p_b
14 = 0.00225, $p_d = 5 \times 10^{-4}$, and $\Pr(\varepsilon|b=0) = 0.1819$. Similarly, we can estimate a lower bound for $\Pr(\varepsilon|b=1)$
15 using the results for $\Pr(\varepsilon|b=0)$. Let $x = \Pr(\varepsilon|b=0)$, then

$$\Pr(\varepsilon|b=1) > (1-p_d) p_b x^2 \left[1 + [(1-p_b)(1-p_d)]^1 + \dots + [(1-p_b)(1-p_d)]^i \right]. \quad (4)$$

16 Therefore,

$$\Pr(\varepsilon|b=1) > (1-p_d) p_b x^2 \sum_{i=0}^{\infty} [(1-p_b)(1-p_d)]^i = \frac{(1-p_d) p_b x^2}{p_d + p_b - p_b p_d}. \quad (5)$$

17 Thus, $\Pr(\varepsilon|b=1) > 0.027$ and $\Pr(\varepsilon) > 0.2089$. The fit in Fig. 2B predicts an ε of ~ 0.23 . Five additional
 18 experiments were run with 1024 initial cells and a very high release rate of 10^5 units/hr. The number of
 19 extinctions were 226, 229, 223, 234, and 219, for a mean extinction frequency of 0.22 ± 0.005 (2x
 20 SEM).