

Recursive relation leading to first passage times of BD process.

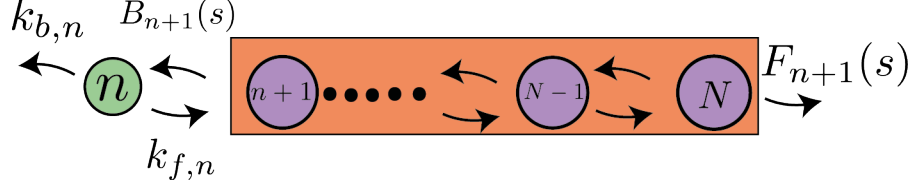


FIG. 1: If you are currently at state n , the (moment generating function of) the event that you eventually make it to $n-1$ is given by $B_n(s)$. This, in turn, is set by the probability to walk from n to $n+1$ and $B_{n+1}(s)$.

The RGN is described as either being unbound, bound to the PAM (in case of CRISPR systems), having formed an R-loop of length $n = 1, \dots, N$ or having cleaved its target substrate. Let us label these states as $n \in [-1, N+1]$, with N being the total length of the guide (target) sequence. Each state $n \in [0, N]$ has rates $k_{f,n}$ and $k_{b,n}$ associated with it for transitioning to $n+1$ and $n-1$ respectively. Originally, we calculated the probability to cleave a sequence, starting from the PAM bound state. To do so, we wrote down a recursive relation for $P_{clv,n}$, the probability to cleave (visit $N+1$ before visiting $n=-1$) starting from state n .

Here, I attempted to take the same approach, but now using moment generating functions in order to (hopefully) get the expected time of the event:

Let $F_n(s)$ / $B_n(s)$ represent the Laplace transform of the first passage time distribution to make it to $n-1/N+1$ starting from state n (see figure). I think the following properly denotes it the way Frank would:

$$F_n(s) \equiv \mathcal{L}\{P(n_t = n-1 | n_0 = n)\} \equiv \mathbb{E}[e^{sT_{n-1}} | T_{n-1} < T_{N+1}] \quad (1)$$

$$B_n(s) \equiv \mathcal{L}\{P(n_t = N+1 | n_0 = n)\} \equiv \mathbb{E}[e^{sT_{N+1}} | T_{N+1} < T_{n-1}] \quad (2)$$

Eventually, we are seeking $B_0(s)$, as this gives us all events that started at state 0 (PAM bound) and eventually made it to state -1 (solution state/ dissociated) before making it to state $N+1$ (post cleavage). More precisely, $B_0(s)$ is the moment generating function. Initially, we seek the first moment:

$$\begin{aligned} T_{ub} &= \frac{\int P(n_t = -1 | n_0 = 0) \times t dt}{\int P(n_t = -1 | n_0 = 0) dt} \\ &= -\frac{1}{B_0(s=0)} \left. \frac{\partial B_0}{\partial s} \right|_{s=0} \\ &= -\left. \frac{\partial \log(B_0(s))}{\partial s} \right|_{s=0} \end{aligned} \quad (3)$$

Additionally, we seek $F_0(s)$, to obtain the time needed to cleave, knowing you did not dissociate. By counting all paths that start at state n and make it to $n-1$ for the first time before it has visited $N+1$ (eventually walk **B**ackwards):

$$B_n(s) = \sum_{m=0}^{\infty} \left(\frac{k_{f,n}}{k_{f,n} + k_{b,n} + s} \right)^m \frac{k_{b,n}}{k_{f,n} + k_{b,n} + s} \quad (4)$$

Similarly, counting any path that ends up in $N+1$ in stead (eventually walk **F**orward):

$$F_n(s) = \sum_{m=0}^{\infty} \left(\frac{k_{f,n}}{k_{f,n} + k_{b,n} + s} \right)^m \frac{k_{f,n}}{k_{f,n} + k_{b,n} + s} \times F_{n+1}(s) \quad (5)$$

Recognizing the geometric series:

$$B_n = \frac{k_{b,n}}{s + k_{b,n} + k_{f,n}(1 - B_{n+1}(s))} \quad (6)$$

$$F_n = \frac{k_{f,n}F_{n+1}(s)}{s + k_{b,n} + k_{f,n}(1 - B_{n+1}(s))} \quad (7)$$

If $\gamma_n = k_{b,n}/k_{f,n}$, then we also see:

$$\frac{1}{\gamma_n}B_n(s) = \frac{F_n(s)}{F_{n+1}(s)} \quad (8)$$

this might help.

BOUNDARY CONDITIONS

$$B_N(s) = \frac{k_{b,N}}{k_{f,n} + k_{b,n} + s} \quad (9)$$

$$F_N(s) = \frac{k_{f,N}}{k_{f,n} + k_{b,n} + s} \quad (10)$$

Typical guidelengths (N) are 20-30 (the most known system, spCas9, has a 20nt guide sequence).