

## KINETIC MODEL

### SIMPLEST KINETIC MODEL

Consider the different state probabilities for a specific virus particle:

- $P_0(t)$ : the CD4-attached complex that will serve as the initial condition. This state is achieved by exposing the host cells to HIV for a long time while maintaining a low temperature.
- $P_1(t)$ : the activated CD4-attached complex. Once temperature is raised, the CD4-bound state transitions into an activated state that has not yet bound to CCR5 but that is receptive to subsequent CCR5 binding.
- $P_2(t)$ : the CD4-CCR5 complex. Once the state described by  $P_1(t)$  binds CCR5, it is described by  $P_2(t)$ .
- $P_3(t)$ : the activated CD4-CCR5 complex. Once CCR5 has bound, the system transitions into the prehairpin complex.
- $Q$ : the membrane-fused state. The prehairpin complex can then initiate membrane fusion and viral entry.  $Q(t)$  represents the probability that the virus has entered the host cell.

The kinetic equations for the probabilities can be written as

$$\begin{aligned}\frac{dP_0(t)}{dt} &= -k_0P_0 + q_1P_1 \\ \frac{dP_1(t)}{dt} &= k_0P_0 - (k_1 + q_1)P_1 + q_2P_2 \\ \frac{dP_2(t)}{dt} &= k_1P_1 - (k_2 + q_2)P_2 + q_3P_3 \\ \frac{dP_3(t)}{dt} &= k_2P_2 - (k_f + q_3)P_3 \\ \frac{dQ(t)}{dt} &= k_fP_3\end{aligned}\tag{1}$$

In the above,  $k_0, k_1, k_2$ , and  $k_f$  represent rates at which the CD4-bound complex is activated, the rate at which CCR5 binds, the rate at which the CD4-CCR5 complex transitions

into the prehairpin complex, and the rate of membrane fusion, respectively. The CCR5 binding rate  $k_1$  would be a function of the available CCR5 or CCR5 binding site concentration. The parameters  $q_i$  are simply the associated backward rates. For example,  $q_2$  is the rate of CCR5 detachment.

When agonists and fusion inhibitors are administered, the rates will change. For example, depleting CCR5 will decrease  $k_1$ , while administration of T20 would decrease  $k_2$  and/or  $k_f$ .

Solutions to Eq. 1 can be explicitly found when  $q_i = 0$  and the viral entry process is irreversible. We can Laplace transform Eqs. 1 and find

$$\tilde{Q}(s) = \frac{k_0 k_2 k_2}{s(s + k_0)(s + k_1)(s + k_2)(s + k_f)}. \quad (2)$$

For constant rates (no inhibitor added), the solution for the fraction of the initial CD4-bound virus in the special irreversible case is

$$Q(t) = 1 + k_f k_0 k_1 k_2 \left[ \frac{e^{-k_f t}}{k_f(k_f - k_0)(k_f - k_1)(k_f - k_2)} + \frac{e^{-k_0 t}}{k_0(k_0 - k_f)(k_0 - k_1)(k_0 - k_2)} \right. \\ \left. + \frac{e^{-k_1 t}}{k_1(k_1 - k_f)(k_1 - k_0)(k_1 - k_2)} + \frac{e^{-k_2 t}}{k_2(k_2 - k_f)(k_2 - k_0)(k_2 - k_1)} \right]. \quad (3)$$

where  $t = 0$  corresponds to the time at which temperature is instantly raised.

The fraction of initial CD4-bound virus that enter up to time  $T_m \approx 48\text{hrs}$  is given by

$$F = Q(T) + k_f \int_T^{T_m - T} P_3(t) dt \approx Q(T) + k_f \int_T^\infty P_3(t) dt, \quad (4)$$

where  $P_3(t > T)$  is determined by the solution to  $P_3(t = T)$  before  $k_1$  is set to zero and the last approximation assumes  $T_m$  is much longer than any other timescale in the problem. We now consider three possible mechanisms of action for timed viral entry inhibitor experiments.

### CCR5 antagonist: $k_1 = 0$ at time $T$

Suppose Maraviroc is administered at time  $t = T$ . We assume its effective mechanism is to bind to and “sequester” the available CCR5. Thus, for a perfect and instantaneous depletion of CCR5, we set  $k_1 = 0$  starting at time  $T$ . If we assume irreversibility and the limit  $T_m \rightarrow \infty$ , the expression 4 for the fraction of viruses entered becomes

$$F \approx Q(T) + P_2(T) + P_3(T) = 1 - \left( \frac{k_1 e^{-k_0 T} - k_0 e^{-k_1 T}}{k_1 - k_0} \right). \quad (5)$$

Thus, for experiments where only Maraviroc is administered at time  $T$ , we fit the relative infection to the two-parameter result for  $F$ . However, since Eq. 5 is symmetric in  $k_0, k_1$ , both  $k_0$  and  $k_1$  cannot be independently obtained. Thus, we need an independent estimate for one of  $k_0, k_1$ .

A simple parameter consistency check is to compare the peak of the derivative  $dF/dT$  to that found from the data. The time  $T_d$  defined by  $d^2F/dT^2|_{T=T_d} = 0$  is

$$T_d = \frac{1}{k_0 - k_1} \ln \left( \frac{k_0}{k_1} \right), \quad (6)$$

which provides a symmetric relationship between  $k_0$  and  $k_1$  given an observed  $T_d$ .

**Late stage T20:  $k_f = 0$  at time  $T$**  Now, consider experiments in which T20 is applied and assume that it acts by preventing the final fusion step. At and after time  $T$ , we set  $k_f = 0$ . The total fraction entered in this case is predicted to be  $F = Q(T_m) \approx Q(\infty) = Q(T)$  given by Eq. 3 above evaluated at  $t = T$ .

If  $k_0$  and  $k_1$  can be estimated for a specific strain by the Maraviroc experiment, these parameter values can be used in Eq. 3 and  $F$  can be fit to T20 data to estimate  $k_2$  and  $k_f$ .

**Early stage T20:  $k_2 = 0$  at time  $T$**

If T20 prevents the transition described rate  $k_2$ , we set  $k_2 = 0$  after  $t = T$ , Eq. 4 becomes

$$F = Q(T) + k_f P_3(T) \int_T^{T_m-T} e^{-(k_f+q_3)(t-T)} dt \approx Q(T) + \frac{k_f P_3(T)}{k_f + q_3}, \quad (7)$$

where  $P_3(T)$  is determined by solving Eqs. 3 before  $k_2$  is set to zero. It will depend on all the rates  $k_i, q_i$ .

Again, in the irreversible case  $q_i = 0$ , we find

$$F \approx Q(T) + P_3(T) = 1 - \frac{k_0 k_2 e^{-k_1 T}}{(k_1 - k_0)(k_1 - k_2)} - \frac{k_1 k_2 e^{-k_0 T}}{(k_0 - k_1)(k_0 - k_2)} - \frac{k_0 k_1 e^{-k_2 T}}{(k_2 - k_0)(k_2 - k_1)}, \quad (8)$$

which is independent of  $k_f$  and adds the additional parameter  $k_2$  into the fitting of T20 data.