

substrate to form an [ES] complex, we can use the equilibrium coefficient to express [ES] in terms of [S].

The equilibrium constant is

$$K'_m = \frac{k_{-1}}{k_1} = \frac{[E][S]}{[ES]} \quad (3.5)$$

Since $[E] = [E_0] - [ES]$ if enzyme is conserved, then

$$[ES] = \frac{[E_0][S]}{(k_{-1}/k_1) + [S]} \quad (3.6)$$

$$[ES] = \frac{[E_0][S]}{K'_m + [S]} \quad (3.7)$$

where $K'_m = k_{-1}/k_1$, which is the dissociation constant of the ES complex. Substituting eq. 3.7 into eq. 3.2 yields

$$v = \frac{d[P]}{dt} = k_2 \frac{[E_0][S]}{K'_m + [S]} = \frac{V_m[S]}{K'_m + [S]} \quad (3.8)$$

where $V_m = k_2[E_0]$.

In this case, the maximum forward velocity of the reaction is V_m . V_m changes if more enzyme is added, but the addition of more substrate has no influence on V_m . K'_m is often called the Michaelis–Menten constant, and the prime reminds us that it was derived by assuming rapid equilibrium in the first step. A low value of K'_m suggests that the enzyme has a high affinity for the substrate. Also, K'_m corresponds to the substrate concentration, giving the half-maximal reaction velocity.

An equation of exactly the same form as eq. 3.8 can be derived with a different, more general assumption applied to the reaction scheme in eq. 3.1.

3.3.2.2. The quasi-steady-state assumption. In many cases the assumption of rapid equilibrium following mass-action kinetics is not valid, although the enzyme–substrate reaction still shows saturation-type kinetics.

G. E. Briggs and J. B. S. Haldane first proposed using the quasi-steady-state assumption. In most experimental systems a closed system (batch reactor) is used in which the initial substrate concentration greatly exceeds the initial enzyme concentration. They suggest that since $[E_0]$ was small, $d[ES]/dt \approx 0$. (This logic is flawed. Do you see why?) Computer simulations of the actual time course represented by eqs. 3.2, 3.3, and 3.4 have shown that *in a closed system the quasi-steady-state hypothesis holds* after a brief transient *if* $[S_0] \gg [E_0]$ (for example, 100×). Figure 3.4 displays one such time course.

By applying the quasi-steady-state assumption to eq. 3.3, we find

$$[ES] = \frac{k_1[E][S]}{k_{-1} + k_2} \quad (3.9)$$

Substituting the enzyme conservation eq. 3.4 in eq. 3.9 yields

$$[ES] = \frac{k_1([E_0] - [ES])[S]}{k_{-1} + k_2} \quad (3.10)$$