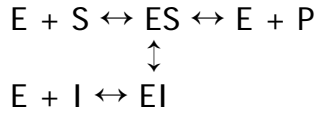


Inhibitors of Enzymes

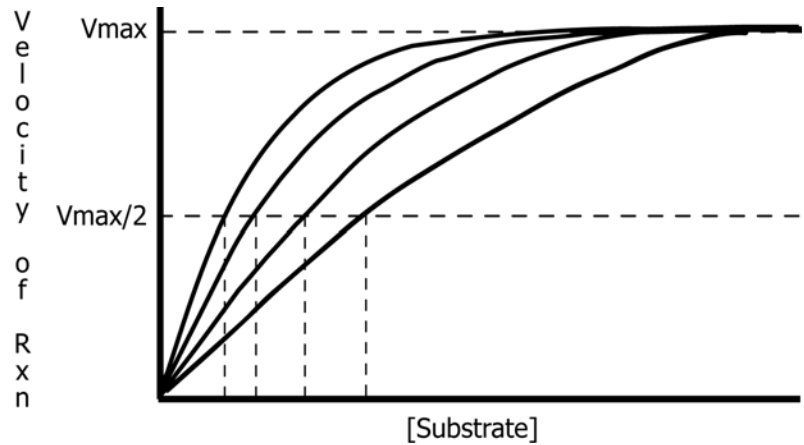
Inhibitors which mimic the transition state bind the most strongly.

There are two classes of inhibitors:

Competitive: These bind to the active site.

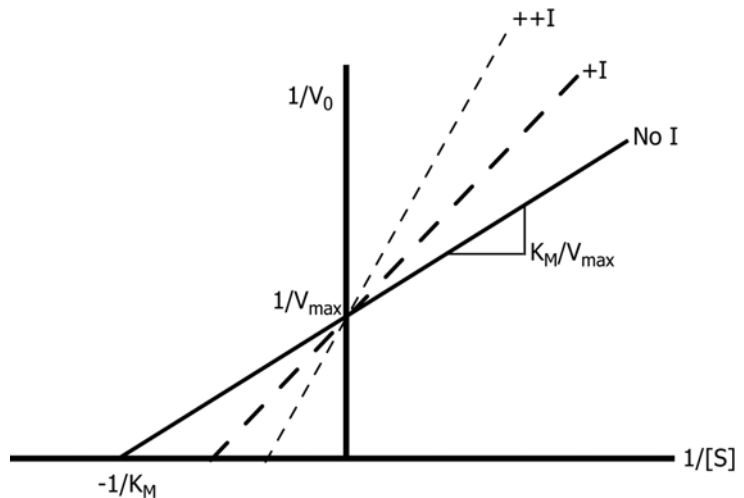


With enough substrate present, you can displace the inhibitor from the active sites on the enzymes; thus, V_{max} does not change. However, since now you need more substrate to bind the enzyme, the apparent K_m goes up. (Michaelis-Menten plot - Top graph)



$$K_M \rightarrow K_{M_{\text{apparent}}} = K_M \left(1 + \frac{[I]}{K_I} \right)$$

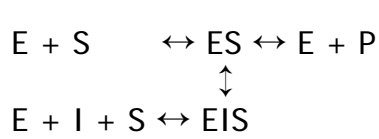
(Lineweaver-Burke plot - Bottom graph)



Inhibitors of Enzymes

Noncompetitive:

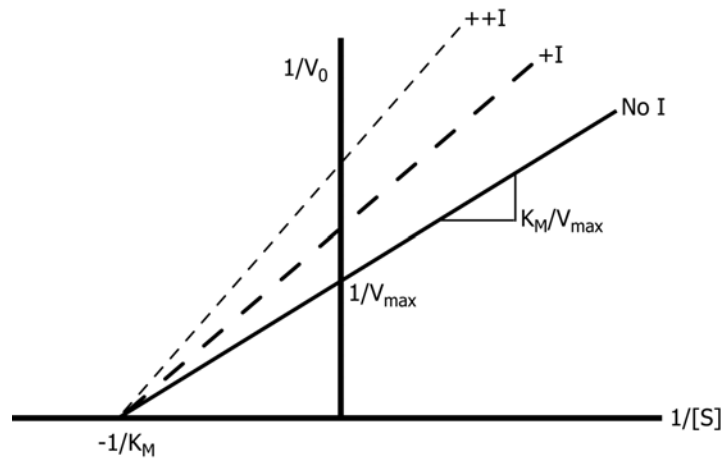
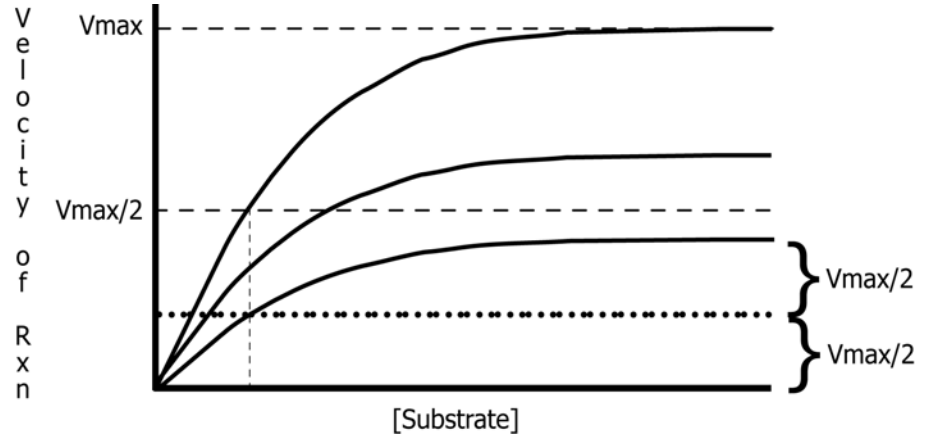
These bind to any site other than the active site.



No matter how much substrate is present, you can never displace the inhibitor. This effectively lowers the number of active enzymes, which therefore lowers the V_{max} .

Since you do not change how well the substrate can bind to the enzyme, the apparent K_M does not change. (Michaelis-Menten - Top graph)

(Lineweaver-Burke - Middle graph)



The other type of noncompetitive inhibition you need to know is **Allostery**.

This follows a sigmoidal binding curve. At low substrate concentrations, the enzyme does not work well, while at high concentrations it works extremely well. There is a non-linear jump in between which is characteristic of the curve. (Bottom graph)

