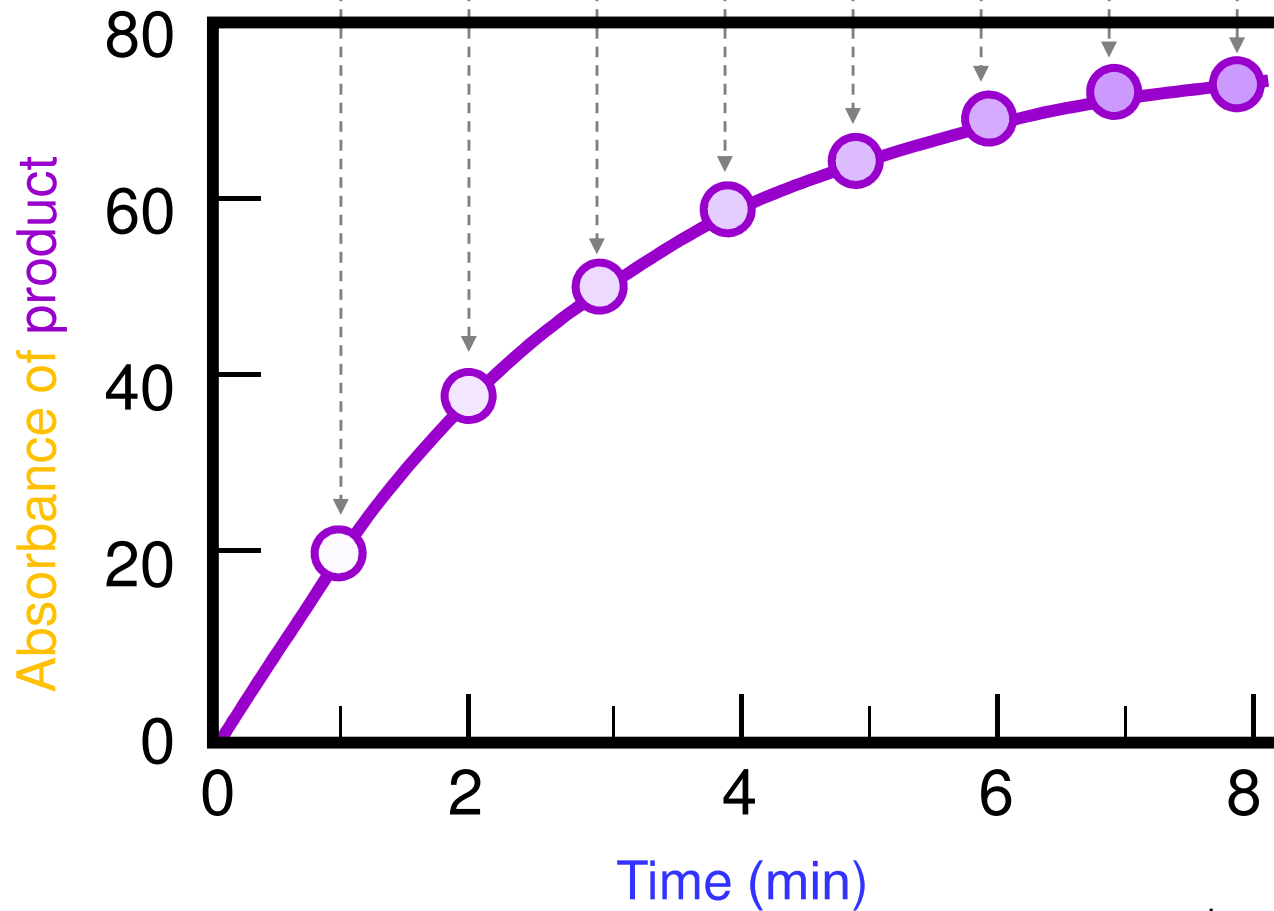
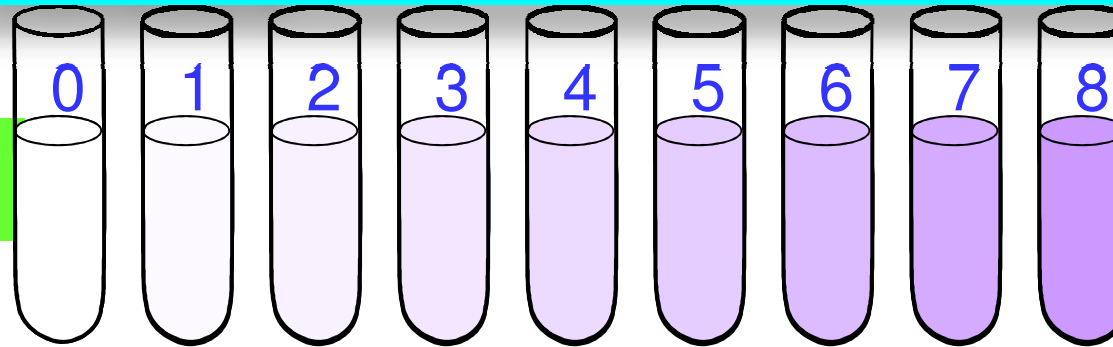
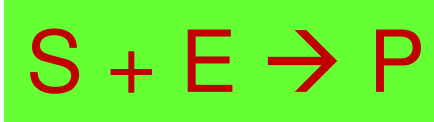
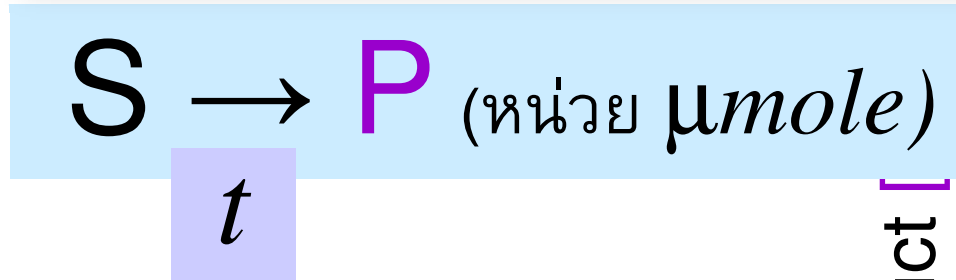


# Enzyme Kinetics



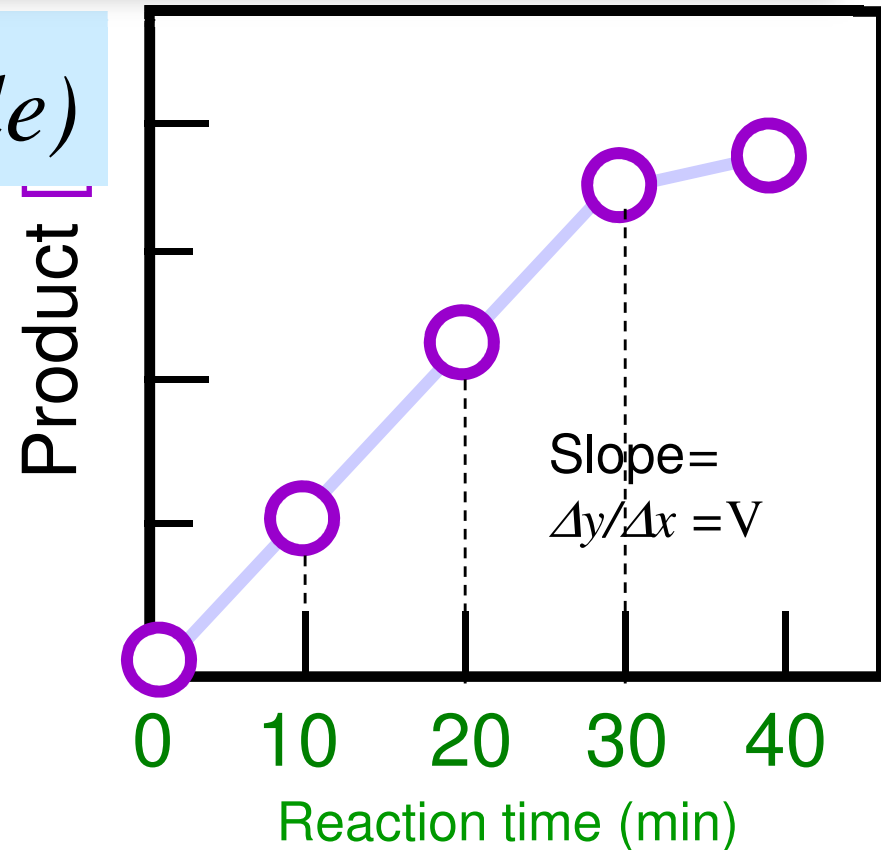
**Rate of reaction (V) คือ อัตราการเกิดผลิตภัณฑ์/หน่วยเวลา**



$$V = [P] / \text{min}$$

ความเร็วของปฏิกิริยา

อัตราการเกิดผลิตภัณฑ์ ต่อนาที



**Enzyme activity** =  $\mu\text{mole} / \text{min}$

หน่วยของ กิจกรรมของเอนไซม์      ปริมาณสารตั้งต้น ที่ลดลง หรือ ปริมาณผลิตภัณฑ์ที่เกิดขึ้น ต่อนาที

**Enzyme activity คือ อัตราการเกิดผลิตภัณฑ์ หน่วยไมโครโมล/หน่วยเวลาเป็นนาที**

**Specific activity = enzyme activity** หน่วยเป็น **unit/ มิลลิกรัมของโปรตีน**

$$V = [P] / \text{min}$$

ความเร็วของปฏิกิริยา



อัตราการเกิดผลิตภัณฑ์ ต่อนาที

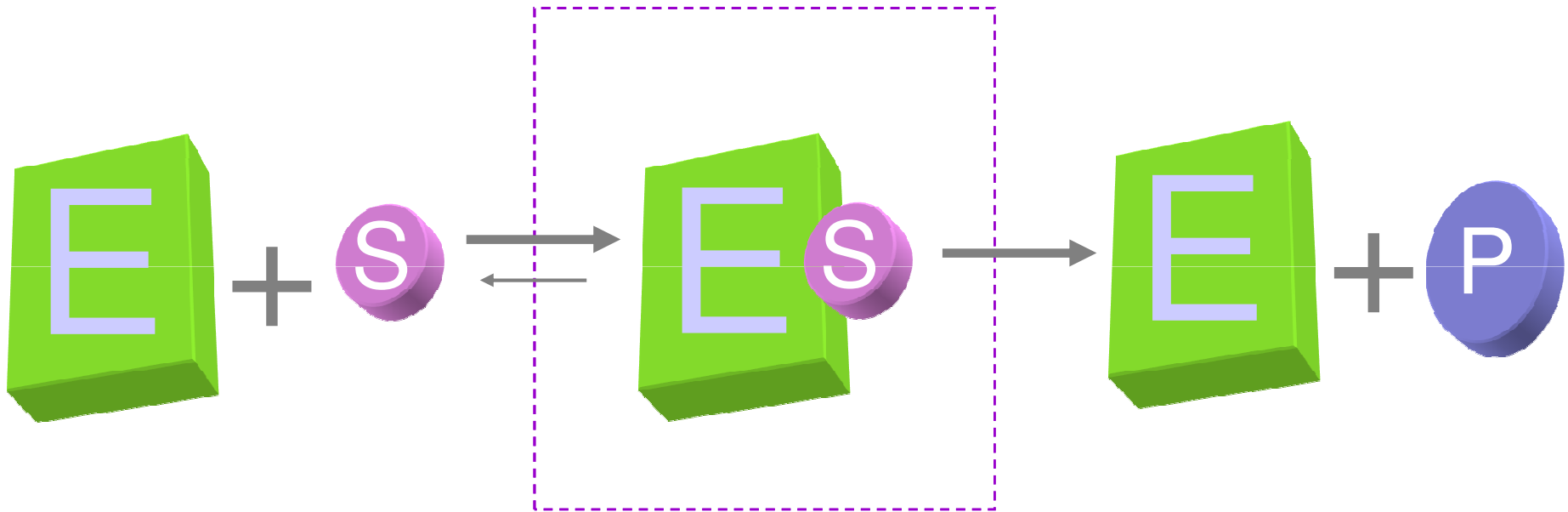
$$\text{Enz activity} = \mu\text{mole} / \text{min}$$

หน่วยของ กิจกรรมของเอนไซม์

ปริมาณสารตั้งต้น ที่ลดลง หรือ  
ปริมาณผลิตภัณฑ์ที่เกิดขึ้น ต่อนาที

$$\text{Specific Activity} = \frac{\text{Enz Activity Units}}{\text{Protein (mg)}}$$

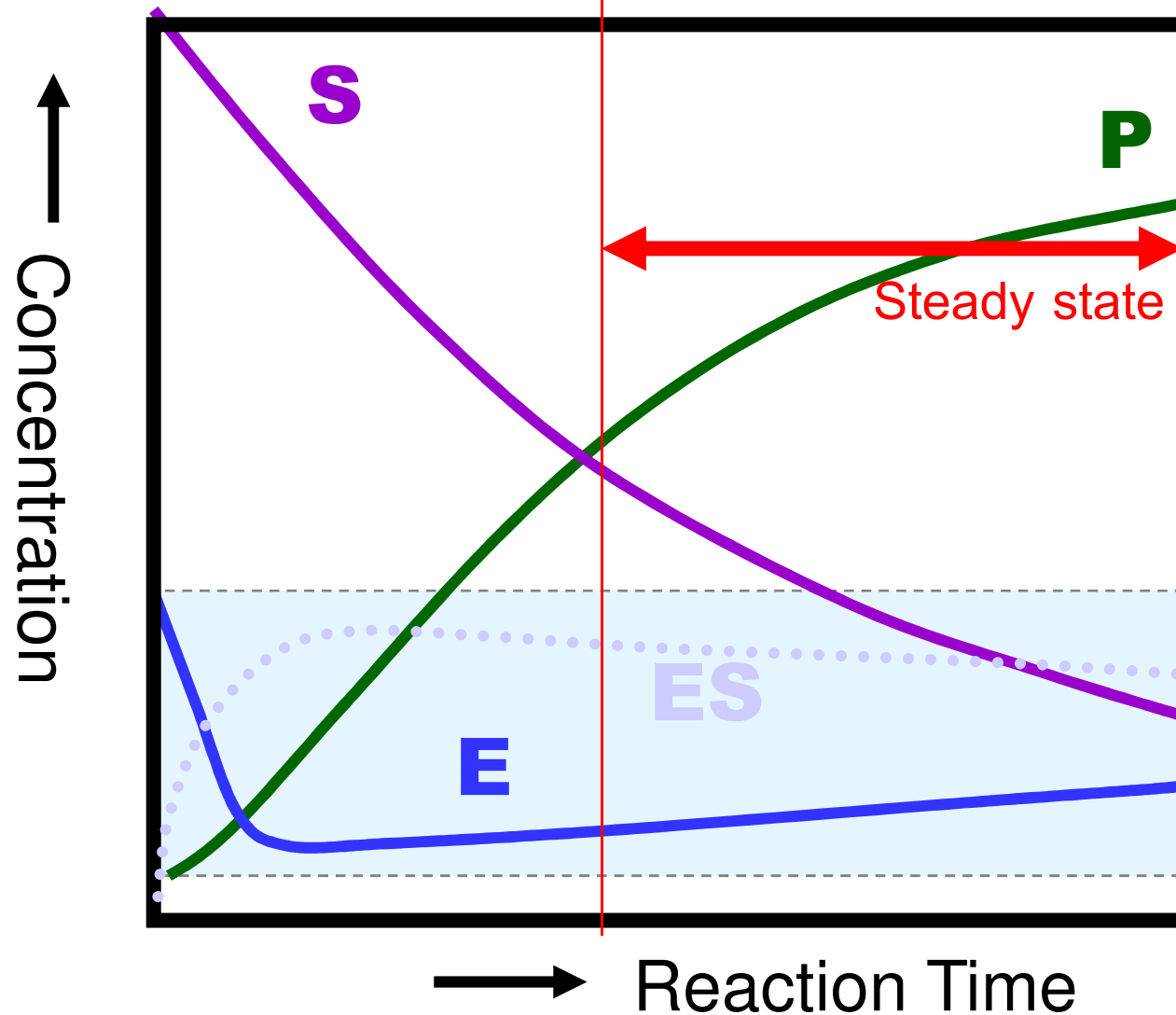
## Steady state



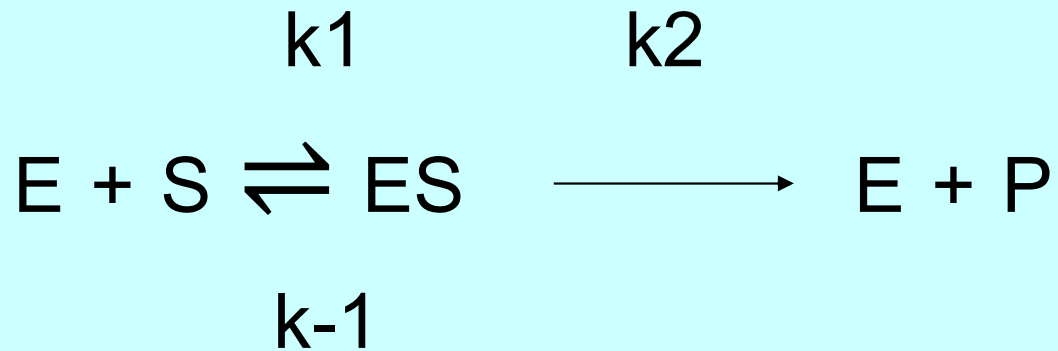
rate of production [ES] = rate of consumption [ES]  
So the concentration of [ES] constant.

# Michaelis-Menten Hypothesis: at Steady state

- 1) No free Enz (all enzymes are active)
- 2)  $[ES]$  const (according to steady state theory)
- 3)  $[P] \rightarrow V_{max}$



## How to derive equation for Km and Vmax



Rate of rx.

$$V_i = \frac{d[ES]}{dt}$$

$V_i$  = initial velocity

$$= k_1 [E] [S] - k_{-1} [ES] - k_2 [ES]$$

$$= k_1 \{ [E_t] - [ES] \} [S] - k_{-1} [ES] - k_2 [ES]$$

$$= k_1 [E_t] [S] - k_1 [ES] [S] - k_{-1} [ES] - k_2 [ES]$$

How to derive  
equation for  
 $K_m$  and  
 $V_{max}$  (cont.)

Hypothesis  $\rightarrow$   $[ES]$  constant

Thus 
$$v_i = \frac{d[ES]}{dt} = 0$$

$$k_1 [E_t] [S] - k_1 [ES] [S] - k_{-1} [ES] - k_2 [ES] = 0$$

$$k_1 [E_t] [S] = k_1 [ES] [S] + k_{-1} [ES] + k_2 [ES]$$

$$k_1 [E_t] [S] = [ES] \{ k_1 [S] + k_{-1} + k_2 \}$$

$$k_1 [E_t] [S] = [ES] \{k_1 [S] + k_{-1} + k_2\}$$

$$[ES] = \frac{k_1 [E_t] [S]}{k_1 [S] + k_{-1} + k_2}$$

$$[ES] = \frac{[E_t] [S]}{[S] + \frac{k_{-1} + k_2}{k_1}} K_m$$



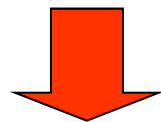
$$k_2 [ES] = V_i$$

$$k_2 [E_t] = V_{max}$$

$$k_2 [ES] = \frac{k_2 [E_t] [S]}{[S] + K_m}$$

$V_{max}$  and  $K_m$  have meaning  
→ application

$$[S] + K_m$$



$$V_i = \frac{V_{max} [S]}{[S] + K_m}$$

$K_m$  = Michaelis-Menten const.

**Michaelis – Menten equation**

$$[S] + K_m$$



Relationship between initial velocity ( $V$ )  
and substrate concentration  $[S]$

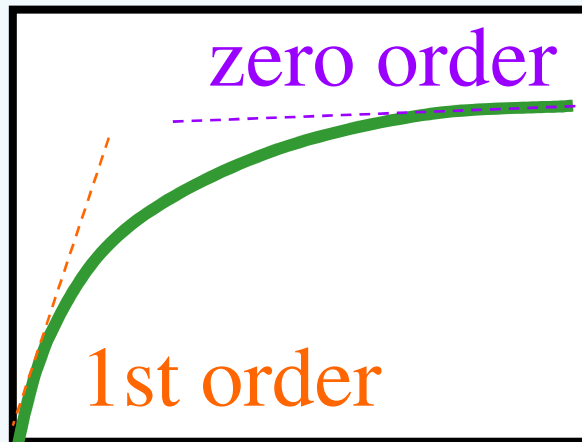
**V<sub>max</sub>: Enzyme activity → 1 Enz unit = ...μmol/min**

V<sub>max</sub> บอกทางอ้อมถึงปริมาณเอนไซม์ ถ้า V<sub>max</sub> มาก คาดว่าปริมาณเอนไซม์มาก

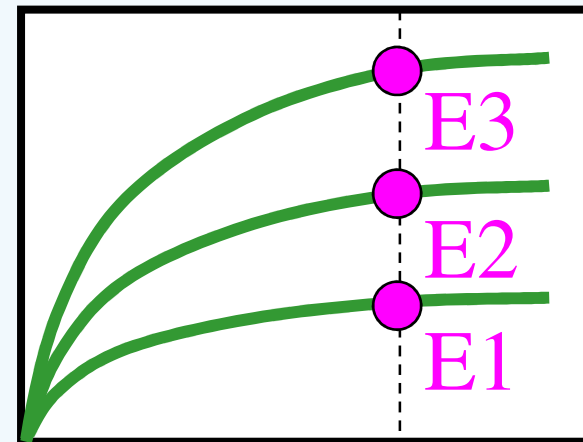
V<sub>max</sub> : enzyme activity

Amount of enzyme = .....μmol/min = 1 Enzyme unit

$$V_i = \frac{V_{max} [S]}{K_m + [S]}$$



[S] = **Low** → **High**



Proportional to  
enzyme concentration

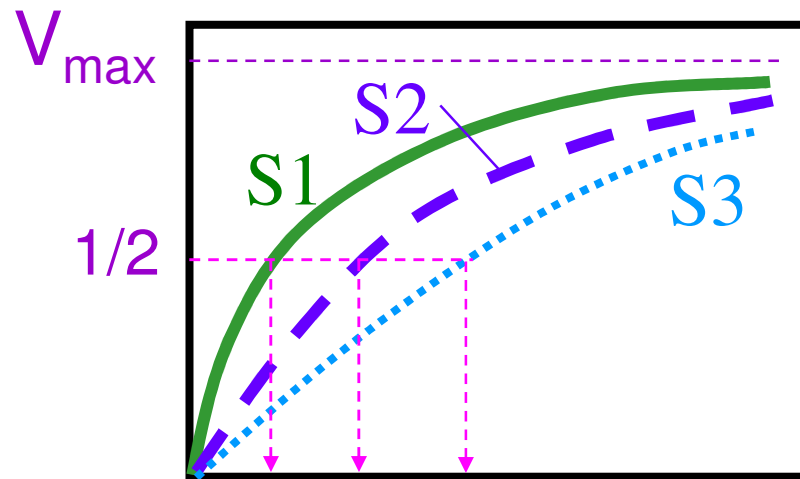
[S] = Fixed concentration

$K_m$ : Affinity with substrate  $\rightarrow$   $[S]$  about 2-5  $K_m$

$K_m$  บอกถึงความชอบพอของเอนไซม์กับซับสเตรท  $K_m$  กับซับสเตรทใ้ดน้อย คาดว่าเอนไซม์ชอบซับสเตรทนั้นมาก

$$K_m = \frac{V_{max}}{2}$$

When using different substrate



S1 S2 S3

$K_m$  Affinity changes

- $K_m$  is a constant
- Small  $K_m$  means tight binding
- High  $K_m$  means weak binding
- Useful to compare  $K_m$  for different substrates for one enzyme

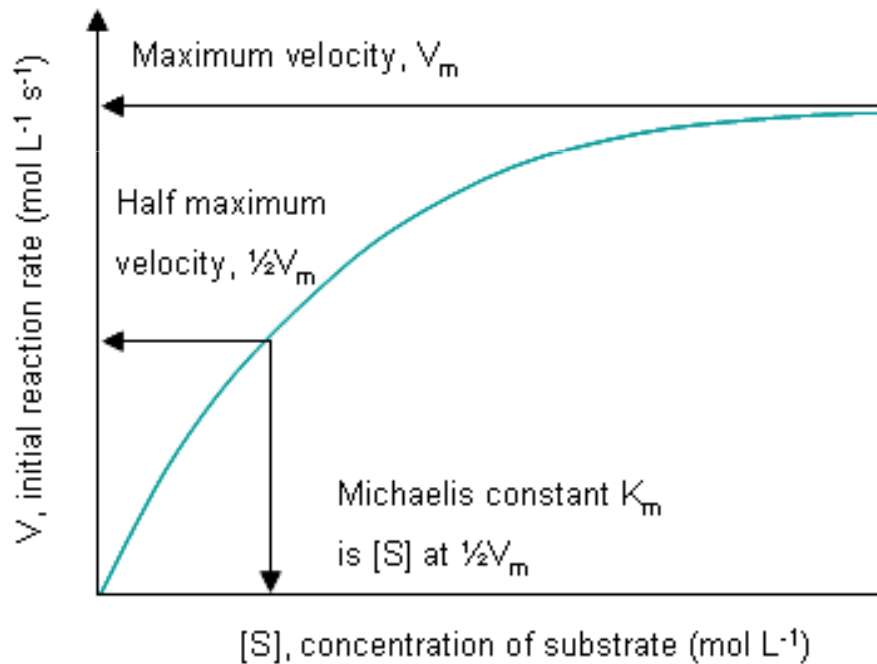
Hexokinase :

D-fructose – 1.5 mM

D-glucose – 0.15 mM

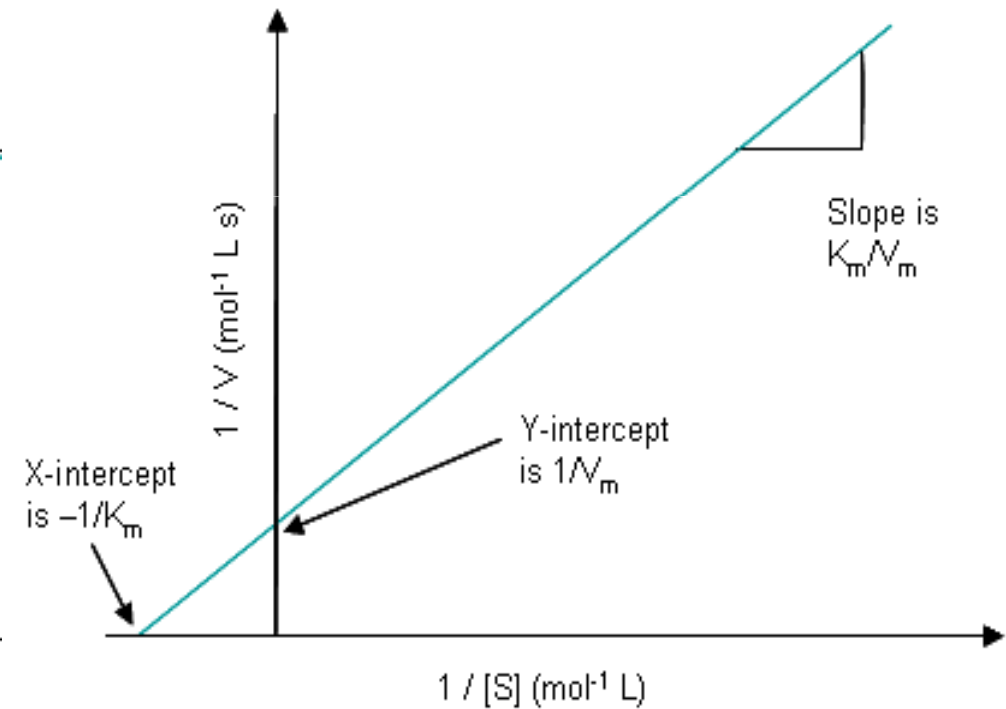
# Direct plot & Lineweaver-Burk plot

## Direct plot



[www.steve.gb.com/science/enzymes.html](http://www.steve.gb.com/science/enzymes.html)

## Lineweaver-Burk plot (Double reciprocal plot)



[www.steve.gb.com/science/enzymes.html](http://www.steve.gb.com/science/enzymes.html)

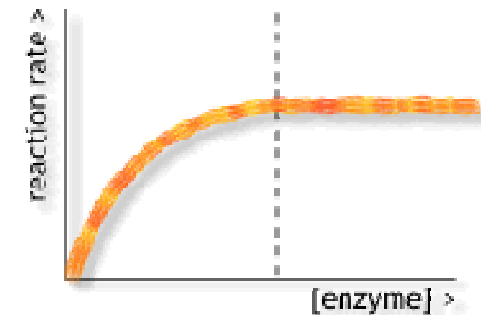
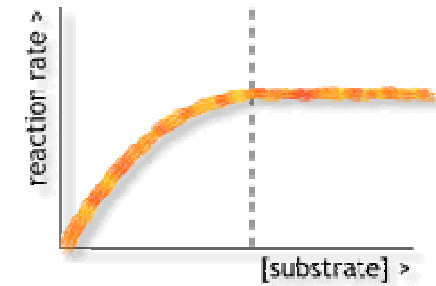
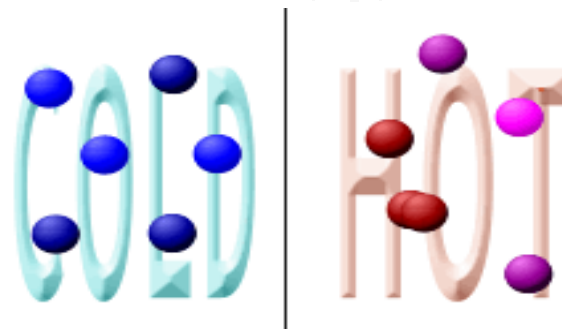
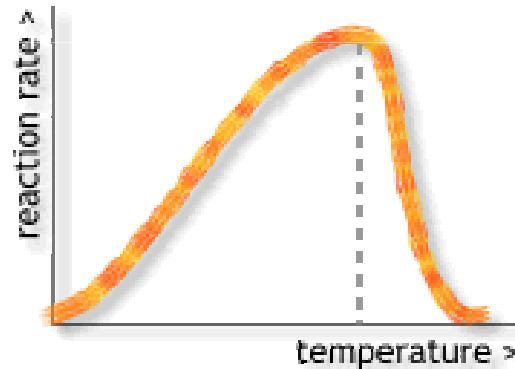
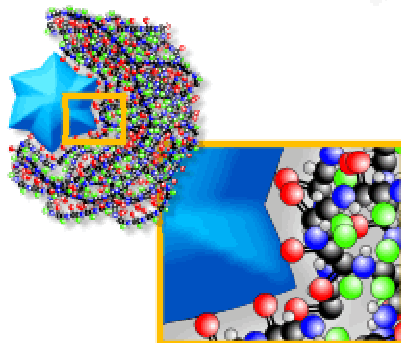
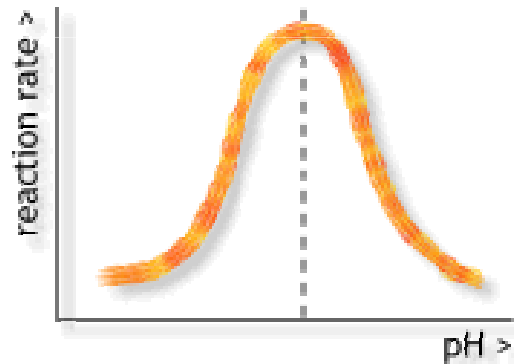
# Turn Over Numbers of Enzymes

| Enzymes               | Substrate              | $k_{\text{cat}}$ ( $\text{s}^{-1}$ ) |
|-----------------------|------------------------|--------------------------------------|
| Catalase              | $\text{H}_2\text{O}_2$ | 40,000,000                           |
| Carbonic anhydrase    | $\text{HCO}_3^-$       | 400,000                              |
| Acetylcholinesterase  | Acetylcholine          | 140,000                              |
| $\beta$ -Lactamase    | Benzylopenicillin      | 2,000                                |
| Fumarase              | Fumarate               | 800                                  |
| RecA protein (ATPase) | ATP                    | 0.4                                  |

The number of product transformed from substrate by one enzyme molecule in one second

# Factors affecting enzyme action

- pH
- Temperature
- Substrate concentration
- Enzyme concentration



# Enzyme inhibition

## Irreversible inhibition

Involve with formation of  
breaking covalent bond

## Reversible inhibition

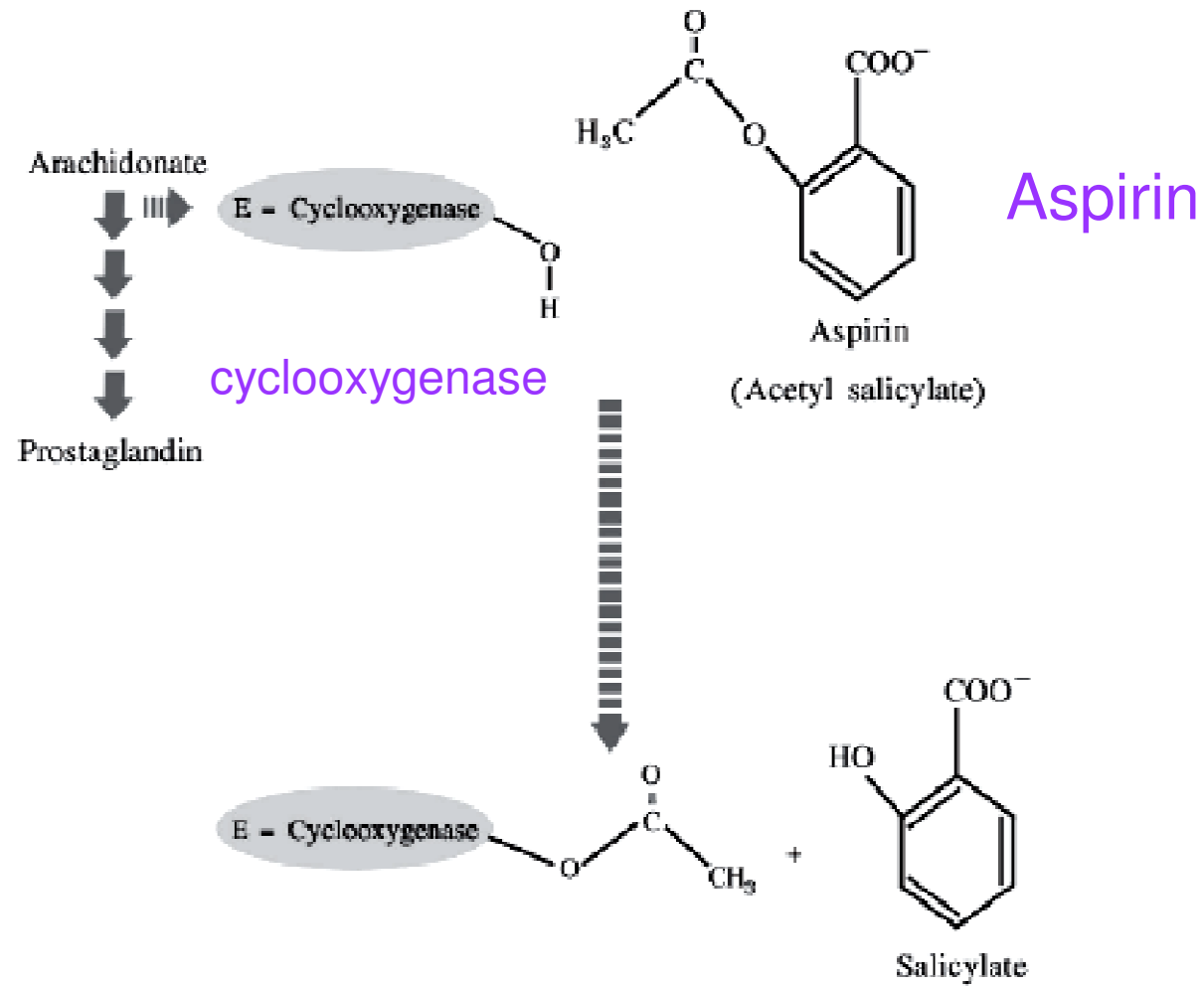
1. Competitive
2. Noncompetitive
3. Uncompetitive

Involve with formation of  
non-covalent bond

Enzyme inhibitors are important because they can be used to.....

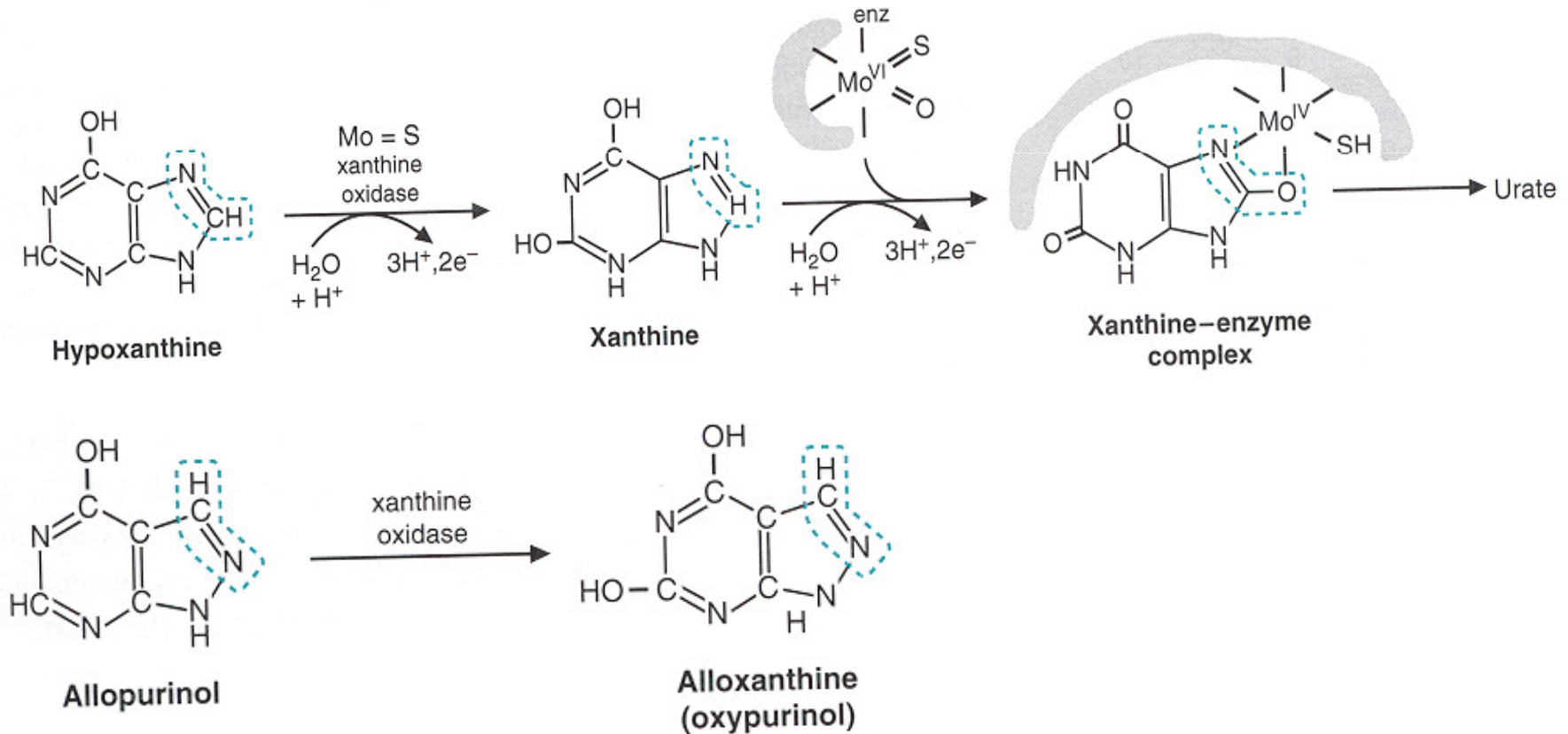
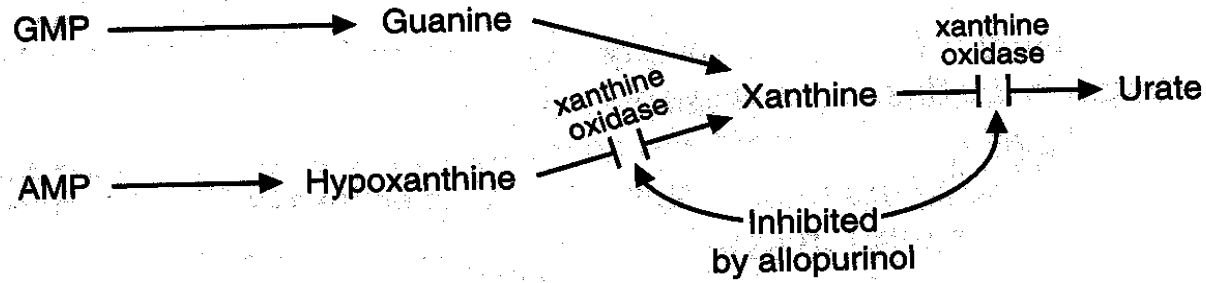
- 1) gain information about
  - the shape on the enzyme active site and the amino acid residues in the active site
  - the chemical mechanism
  - the regulation or control of a metabolic pathway
- 2) design new drug

# Example of irreversible inhibition - **aspirin**





# Example of irreversible inhibition - **allopurinol**



# Reversible inhibition mechanism

|                          | ▶ Competitive   | ▣ Non-competitive  | ◩ Uncompetitive   |
|--------------------------|---|--|---|
| Cartoon Guide            | <p>Substrate</p> <p>Inhibitor</p> <p>Compete for active site</p>                                      | <p>Different site</p>  |   |
| Equation and Description | $E + S \rightleftharpoons ES \rightarrow E + P$ $+ I$ $\updownarrow$ $EI$                             | $E + S \rightleftharpoons ES \rightarrow E + P$ $+ I$ $\updownarrow$ $EI + S \rightleftharpoons EIS$ | $E + S \rightleftharpoons ES \rightarrow E + P$ $+ I$ $\updownarrow$ $EIS$          |
|                          | <p>[I] binds to free [E] only, and competes with [S]; increasing [S] overcomes inhibition by [I].</p> | <p>[I] binds to free [E] or [ES] complex; Increasing [S] can not overcome [I] inhibition.</p>        | <p>[I] binds to [ES] complex only, increasing [S] favors the inhibition by [I].</p> |

# Enzyme inhibition plots

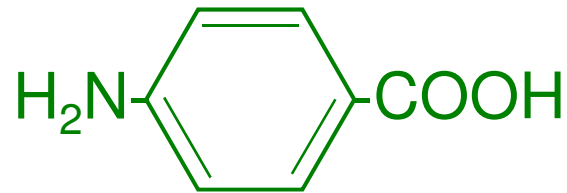
|                   | ▶ Competitive   | ▣ Non-competitive   | ◀ Uncompetitive  |
|-------------------|---|---|--|
| Direct Plots      | <p><math>V</math> vs <math>[S], \text{mM}</math>. <math>V_{\max}</math> unchanged, <math>K_m</math> increased to <math>K_m'</math>.</p> | <p><math>V</math> vs <math>[S], \text{mM}</math>. <math>V_{\max}</math> decreased to <math>V_{\max}'</math>, <math>K_m = K_m'</math>.</p> | <p><math>V</math> vs <math>[S], \text{mM}</math>. Both <math>V_{\max}</math> and <math>K_m</math> decreased to <math>V_{\max}'</math> and <math>K_m'</math>.</p> |
| Double Reciprocal | <p>Intersect at Y axis. <math>1/V_{\max}</math> unchanged, <math>1/K_m</math> increased.</p>  | <p>Intersect at X axis. <math>1/V_{\max}</math> decreased, <math>1/K_m</math> unchanged.</p>  | <p>Two parallel lines. <math>1/V_{\max}</math> and <math>1/K_m</math> both decreased.</p>  |

# Example of reversible inhibition – sulfa drug



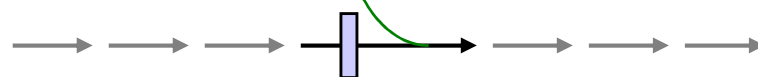
Domagk (1939)

Para-aminobenzoic acid (PABA)



Bacteria needs PABA for the biosynthesis of folic acid

Precursor



Folic acid

Tetrahydro-folic acid



Sulfa drugs has similar structure with PABA, and inhibit bacteria growth.

Sulfanilamide

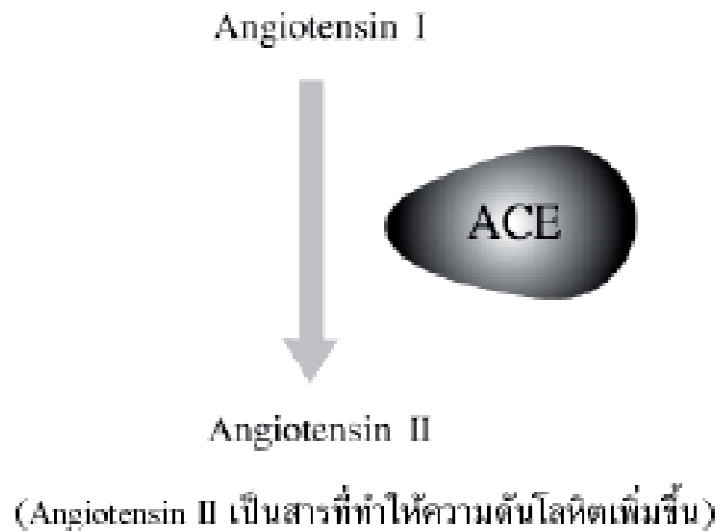
Sulfa drug (anti-inflammation)

## Example of reversible inhibition

Enz = ACE (angiotensin converting enzyme)

Substrate = angiotensin I

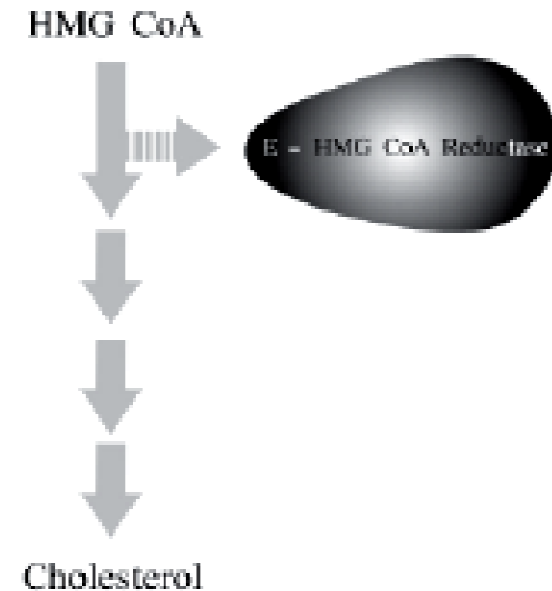
Competitive inhibitor = Captopril and enalapril (blood pressure ↓)



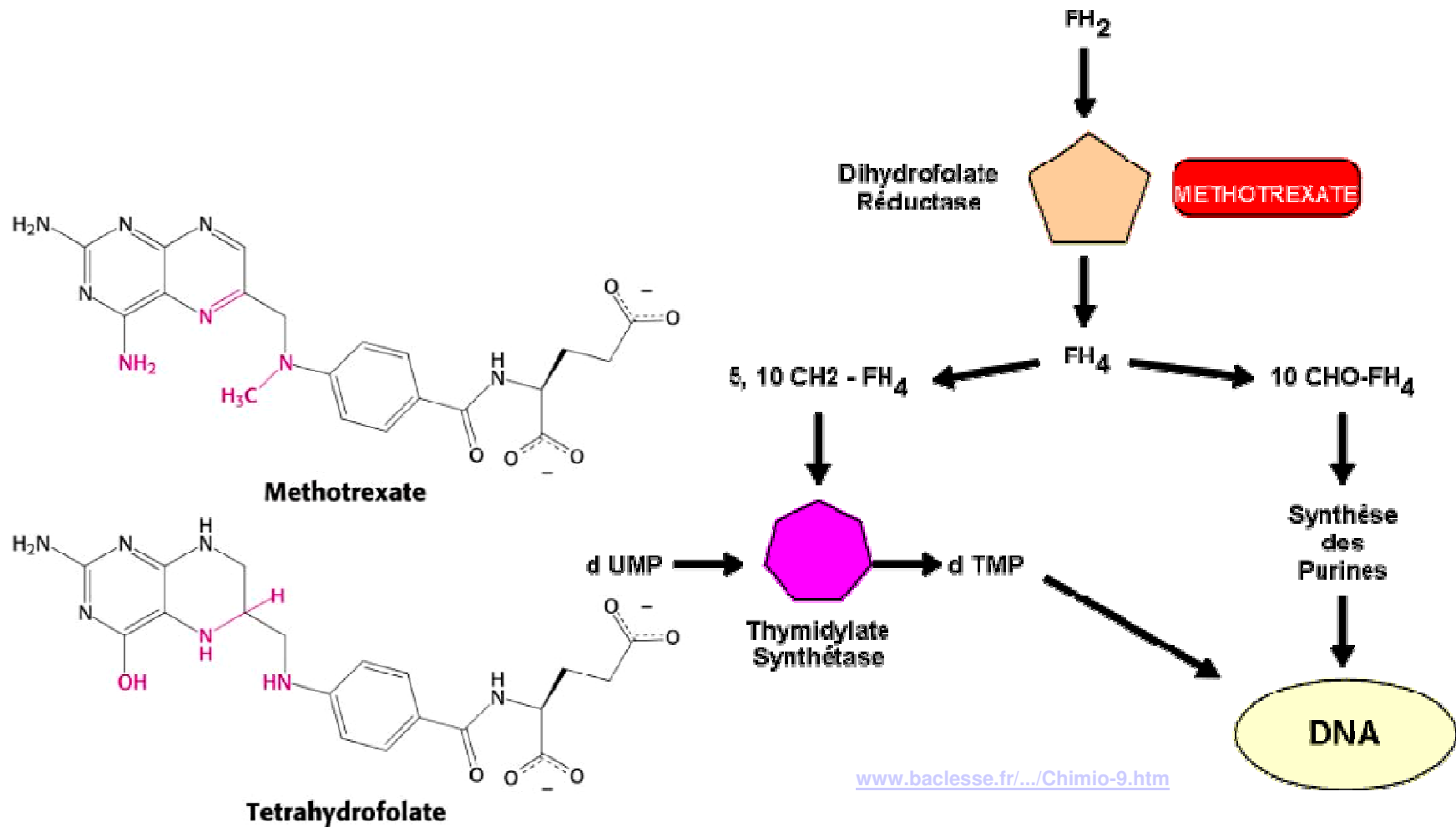
Enz = HMG CoA reductase

Substrate = HMG CoA

Competitive inhibitor = lovastatin and mevilonin

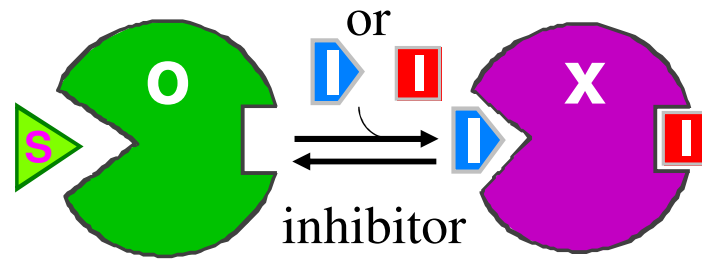


# Example of reversible inhibition – cancer treatment

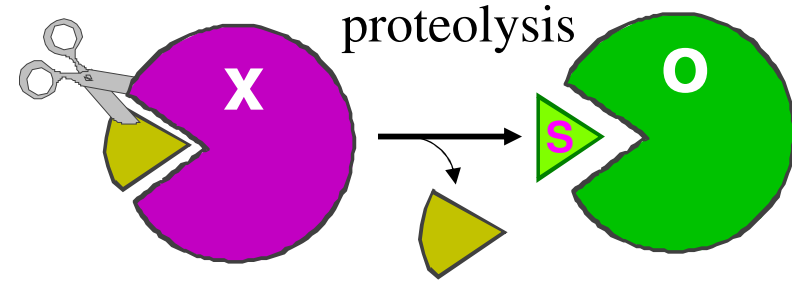


# Enzyme Regulation

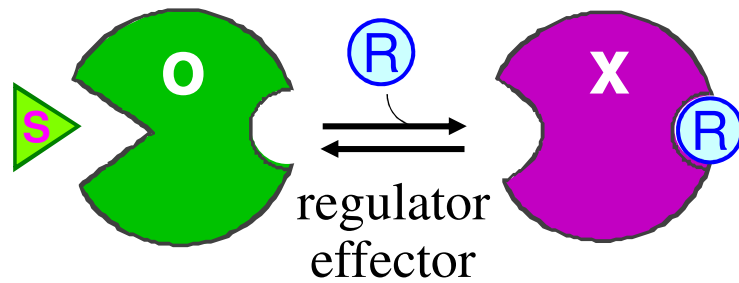
## Inhibitor



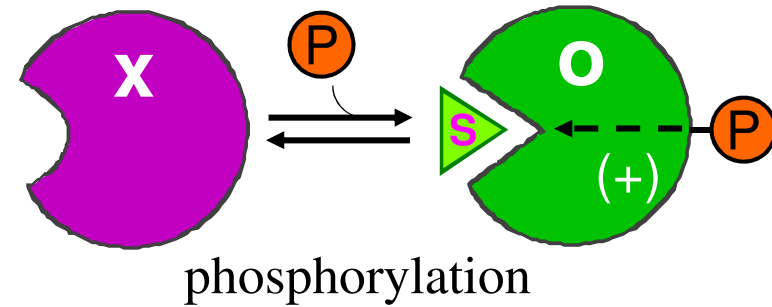
## Proteolysis



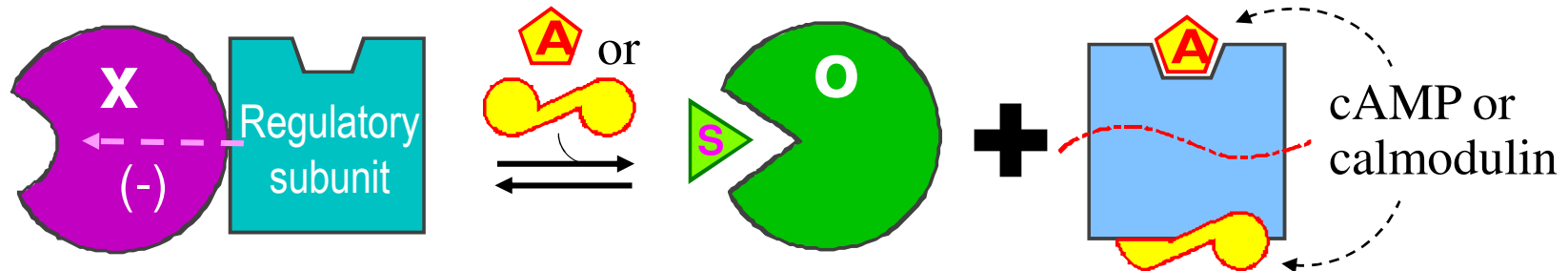
## Feedback regulation



## Covalent modification(Phosphorylation)



## Signal transduction



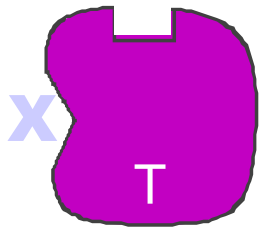
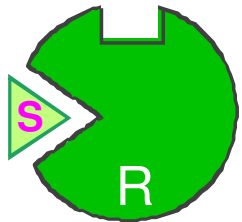
# Allosteric enzyme

## Kinetics

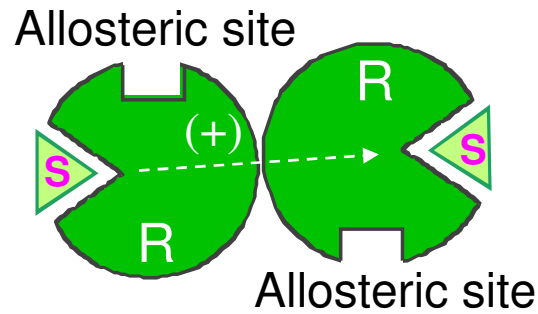
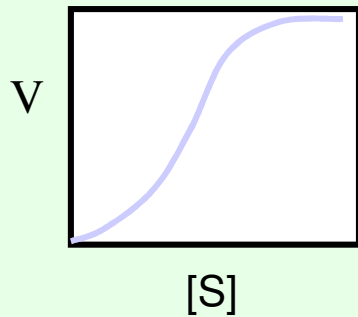
## Models

## Cooperation

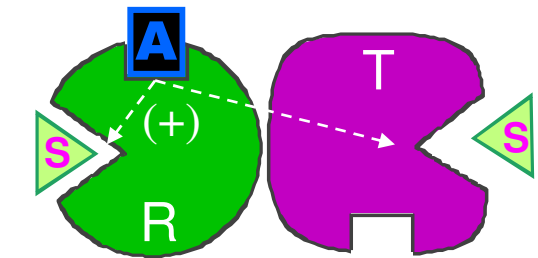
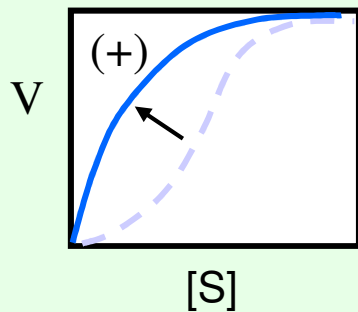
R = Relax  
(active)



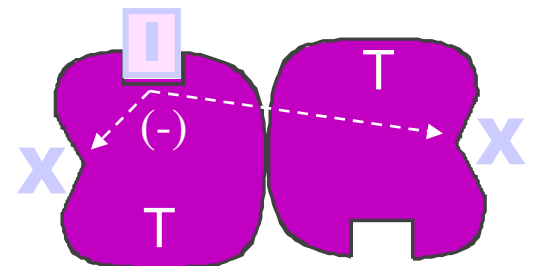
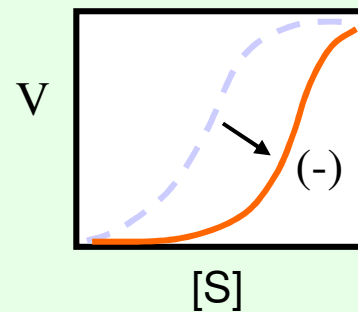
T = Tense  
(inactive)



Homotropic  
(+)  
Concerted



Heterotropic  
(+)  
Sequential



Heterotropic  
(-)  
Concerted



# Summary

Enzyme is a biological catalyst. It decreases  $E_a$  but not  $\Delta G$ .

Enzyme can be classified into 6 classes. Isoenzymes are different but catalyze the same rx.

$K_m \rightarrow$  affinity for substrate,  $V_{max} \rightarrow$  enzyme activity

Enzyme inhibition consists of irreversible and reversible inhibition. Competitive inhibition is advantage to drug design.

There are many ways to regulate enzymes. Allosteric enzyme is one type of enzyme regulation.