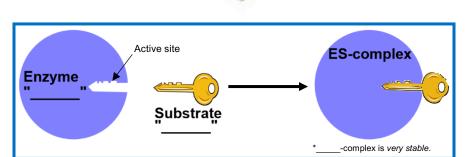
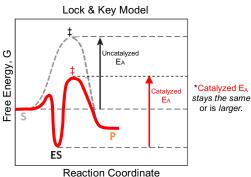
## **CONCEPT: LOCK & KEY VS INDUCED FIT MODELS**

- •There are \_\_\_\_\_ main models for enzyme-substrate specificity.
- 1) Lock & Key Model: active site shape is rigid & complementary to the \_\_\_\_\_\_ shape for a perfect "puzzle-piece" fit.
  - □ \_\_\_\_\_\_ likely model for enzyme catalysis since ES-complex is stabilized & E<sub>A</sub> stays the same or increases.

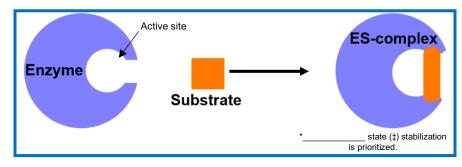
**EXAMPLE:** Lock & Key Model.

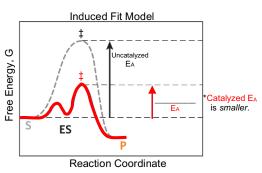




- 2) Induced Fit Model: active site shape is adjustable & more complementary to the \_\_\_\_\_\_ state (‡) to stabilize it.
  - □ Conformational changes *induced* in active site & substrate.
  - □ \_\_\_\_\_\_ likely model for enzyme catalysis since ES-complex is \_\_\_\_\_ stabilized & E<sub>A</sub> decreases.

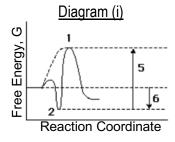
**EXAMPLE:** Induced Fit Model.

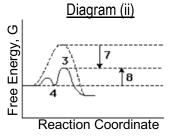




**PRACTICE:** Compare the two enzyme-catalyzed reaction diagrams below (i & ii) to determine which of the following is true.

- a) The ES-complex in diagram (i) is #2 and in diagram (ii) is #3...
- b) Catalyzed E<sub>A</sub> in diagram (i) is arrow #5 and in diagram (ii) is arrow #7.
- c) Binding energy in diagram (i) is arrow #5 and in diagram (ii) is arrow #7.
- d) Diagram (i) describes a "lock & key" model while (ii) describes more of an "induced fit" model.





## **CONCEPT:** LOCK & KEY VS INDUCED FIT MODELS

PRACTICE: What is a potential disadvantage for an enzyme having too high of an affinity for its substrate?

- a) Enzyme catalysis may not occur.
- b) The enzyme would catalyze the reaction but remain covalently & permanently linked to the product.
- c) The ES complex could end up in a large energy trough with a consequentially large E<sub>A</sub> to the transition state.
- d) There's no potential disadvantage to strong affinity. The stronger the affinity, the faster/better the catalysis.
- e) a, b and c.
- f) a and c.

**PRACTICE:** Select the best option that fills in the blanks appropriately in the order of their appearance.

The left graph depicts an energy diagram for the "\_\_\_\_\_\_" model of enzyme-substrate specificity, whereas the right diagram depicts an energy diagram for the modern "\_\_\_\_\_\_" model. In the lock & key model, the enzyme binds tightly and precisely to the "\_\_\_\_\_\_". In the induced fit model, the enzyme binds weakly to the "\_\_\_\_\_\_" and then changes conformation to bind tightly to the "\_\_\_\_\_\_."

- a) Lock & Key; Induced fit; transition state; substrate; product.
- b) Lock & Key; Induced fit; substrate, transition state; product.
- c) Lock & Key; Induced fit; substrate; substrate; transition state.
- d) Induced fit; Lock & Key; transition state; product; substrate.
- e) Induced fit; Lock & Key; product; transition state; substrate.
- f) Induced fit; Lock & Key; substrate; product; transition state.

