Chem 452 - Lecture 5
Catalytic Strategies
Part 3

Question of the Day: Transition states in enzyme catalyzed reactions are usually very unstable and therefore hard to observe. What was the trick used by investigators to observe the transition state for the the hydrolyis of ATP by the myosin motor domain ATPase?

Introduction

- Enzymes exhibit both catalytic power and specificity
- We will consider closely, four examples.

Introduction

- Some Basic Catalytic Principles
  - Covalent Catalysis
  - General Acid/Base Catalysis
  - Catalysis by Approximation
  - Metal Ion Catalysis
  - Transition State Stabilization

Introduction

- Chymotrypsin (1gct) 3.4.21.1
  - A Hydrolase, which cleaves peptide bonds in proteins
- Carbonic anhydrase (1ca2) 4.2.1.1
  - A Lyase, which adds water to CO_2.
- EcoRV (1rvb) 3.1.21.4
  - A Hydrolase, which cleave phosphodiester bonds in DNA
- Myosin motor domain ATPase (1fmv & 1fmw) 3.6.4.1 (Chapter 9)
  - An enzyme that couples the hydrolysis of ATP to the mechanical motion.

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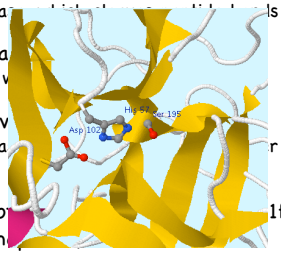

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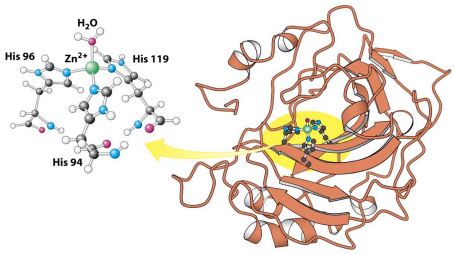

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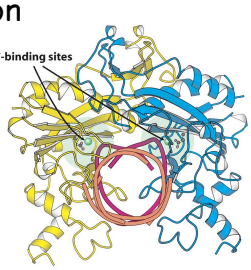

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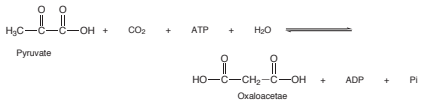
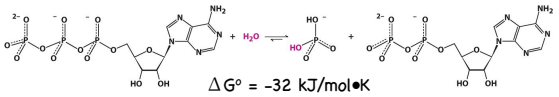
Myosin ATPase

- ATP is considered the energy currency of the cell.
  - The hydrolysis of ATP is used to drive many unfavorable processes
    - Reactions (ligases)
    - Transport across membranes
    - Mechanical motion
- For example, pyruvate carboxylase
  - Couples the hydrolysis of ATP to the formation of a carbon-carbon bond.



Myosin ATPase

- Myosin ATPase couples the hydrolysis of ATP to mechanical motion.
  - The reaction is expected to have a pentavalent transition state intermediate



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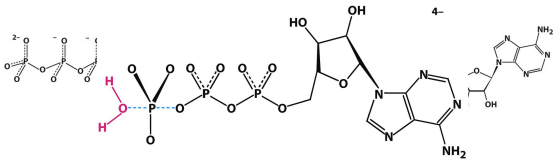

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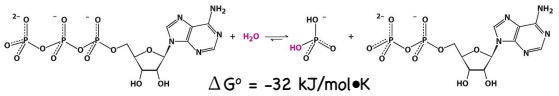

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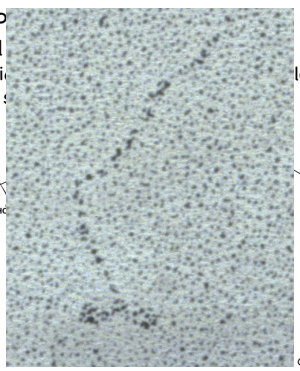
Myosin ATPase

- Myosin ATP ATP to mechanical
  - The reacti lent transition



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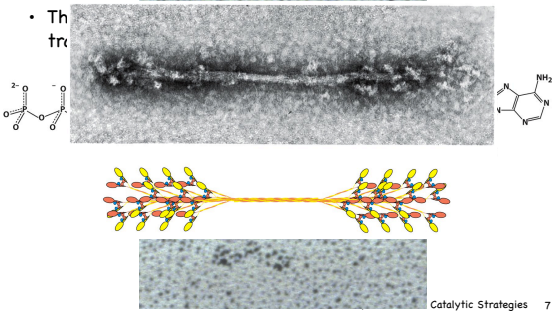

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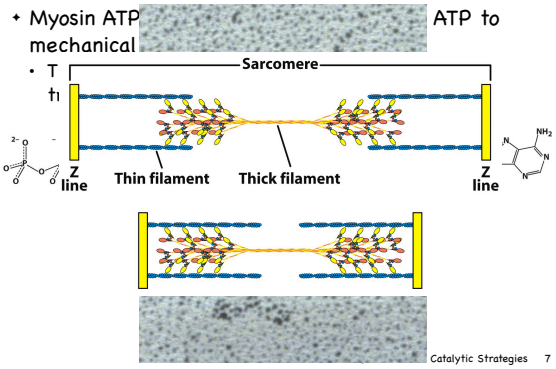

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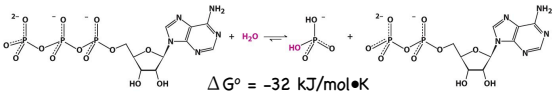

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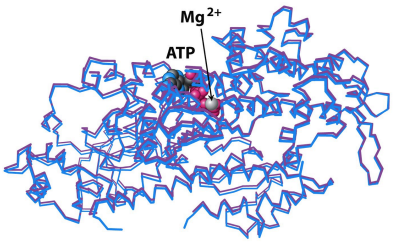
Myosin ATPase

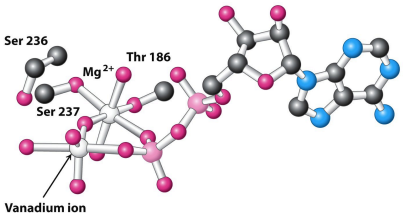
- An X-ray crystal structure of myosin II ATPase with ATP bound did not reveal how the γ-phosphate would be attacked by water.

An overlay with and without ATP



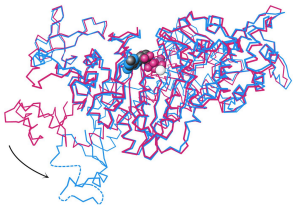
Myosin ATPase

- An X-ray crystal structure of myosin II ATPase with a transition state analogue for ATP revealed a mechanism
  - VO_4^{3-} + ADP was substituted for ATP.



Myosin ATPase

- In order to stabilize the transition state the Myosin II ATPase must undergo a marked conformational change



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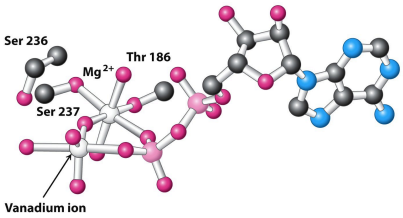

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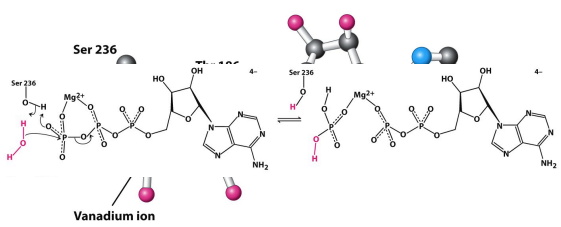

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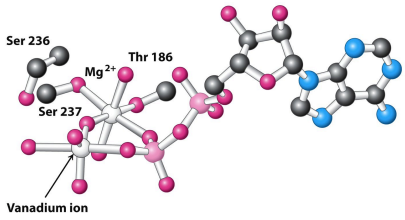
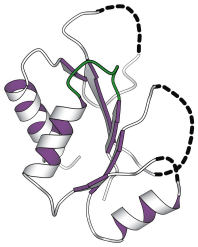


Myosin ATPase

- Many NPTase share a similar binding domain for the nucleotide, and contains a peptide look called the "P-loop"



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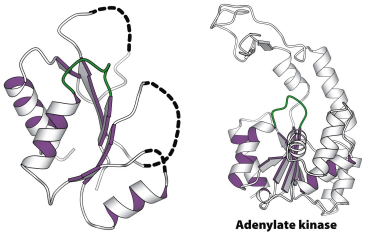

Adenylate kinase

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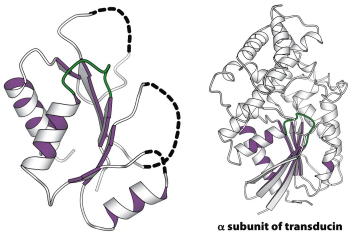

α subunit of transducin

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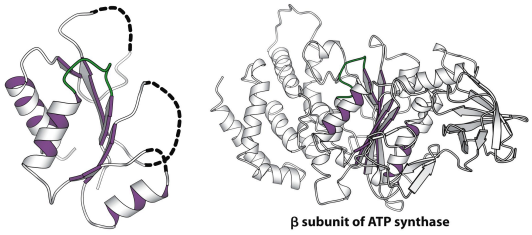

β subunit of ATP synthase

Next up

- Regulatory Strategies (Chapter 10)