

# MECHANISM OF SERINE PROTEASE

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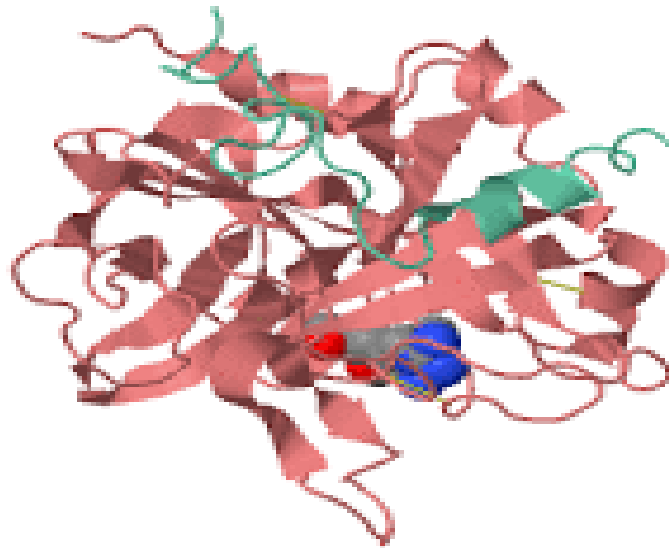
# INTRODUCTION

- They are a large family of proteolytic enzymes
- All such have serine residue at their active site which plays a crucial part in the enzymatic activity.

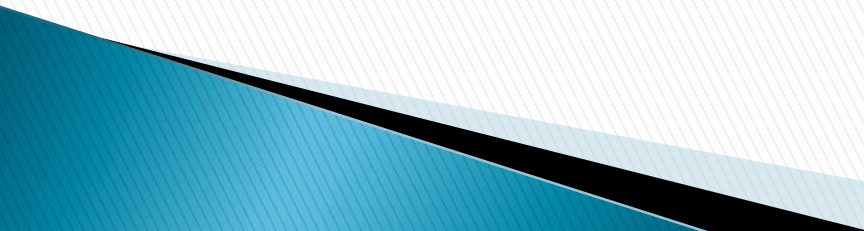
**Serine proteases** are enzymes that cleave peptide bonds in proteins, in which **serine** serves as the nucleophilic amino acid at the (enzyme's) active site. They are found ubiquitously in both eukaryotes and prokaryotes.

All the mammalian serine proteases have a common 3D structure. They all have the same catalytic mechanism.

# Three Dimensional Structure



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- Serine proteases include:
  - the pancreatic proteases: trypsin, chymotrypsin and elastase,
  - *various tissue/intracellular* proteases such as leukocyte elastase
  - enzymes of the blood clotting cascade
  - some enzymes of complement system
  - Many serine proteases are synthesized as inactive precursors (zymogens) which are activated by proteolysis
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# Classification

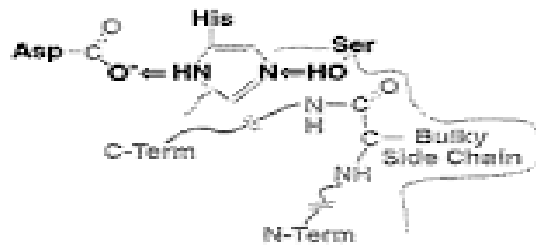
- Trypsin like serine peptidase – if it prefers to cleave peptide bonds of lys & arg
- Chymotrypsin like serine peptidase – if it prefers to cleave aromatic amino acids.
- Elastase like serine peptidase – if it prefers to cleave amino acids with small side chain groups like ala.

# Mechanism of serine protease catalysis.

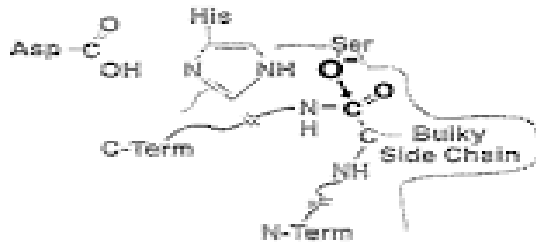
- ▶ The three key residues in serine proteases are collectively known as the catalytic triad. Aspartate is the H-bond acceptor, holding histidine in place. Histidine is a general acid/base catalyst, acting on serine, water, and the addition-elimination leaving groups. Serine is the nucleophile which initiates attack using its oxygen atom.

- ▶ The reaction proceeds as two addition–elimination steps. The first forms a covalent acyl intermediate, while the second deacylates the serine using water, freeing up the active site. All proton transfers are accounted for using histidine as an acid and a base, and the oxyanion hole stabilises the TI.

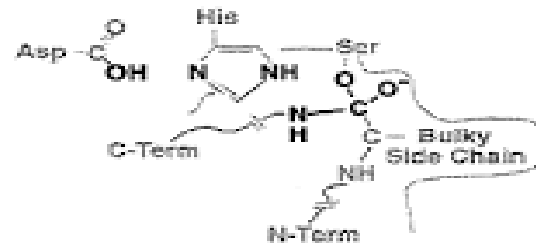
**Step 1: H<sup>+</sup> shift generates Ser-O<sup>-</sup>**



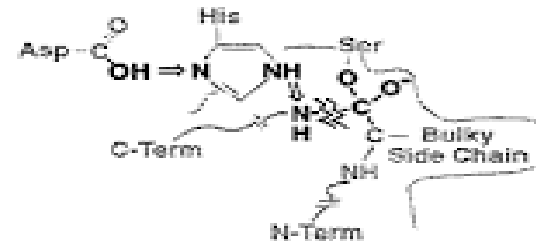
**Step 2: Ser-O<sup>-</sup> binds to C=O**



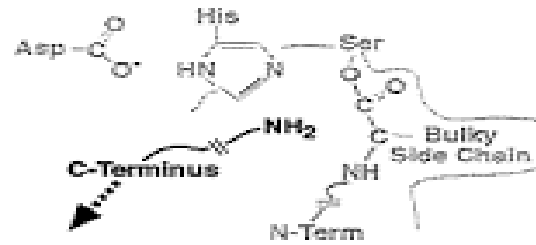
**Step 3: Transition State I**



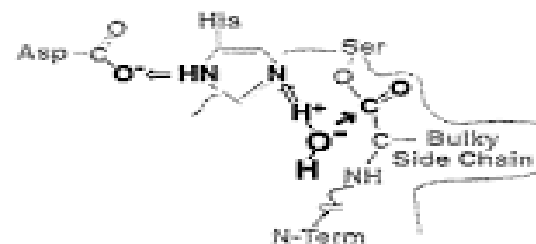
**Step 4: Peptide bond breaks**



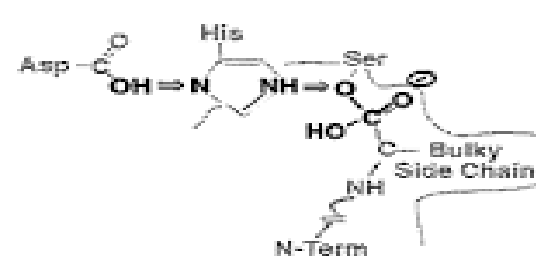
**Step 5: C-Terminal peptide leaves**



**Step 6: Ionization of water**



**Step 7: Transition State II**



**Step 8: N-terminal peptide leaves**

