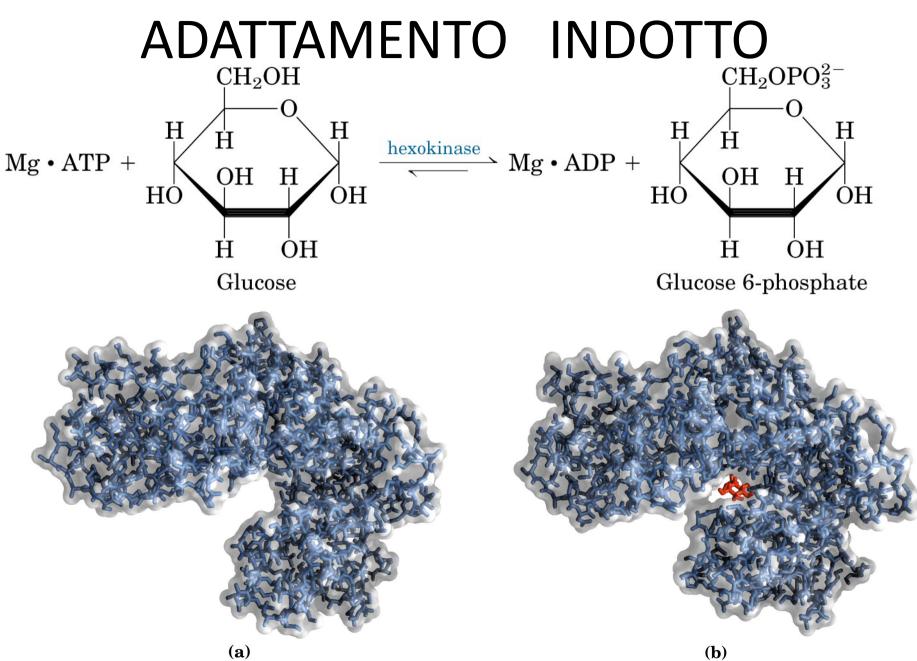
MECCANISMI DI CATALISI

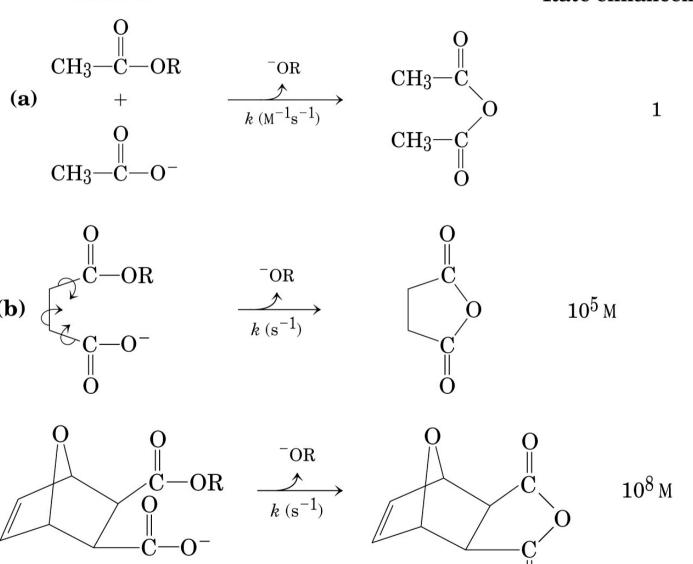
Fattori che influenzano ΔG^{\ddagger} :

- Minore libertà di movimento delle molecole
- Desolvatazione
- Distorsione del substrato
- Allineamento corretto gruppi funzionale dell'enzima



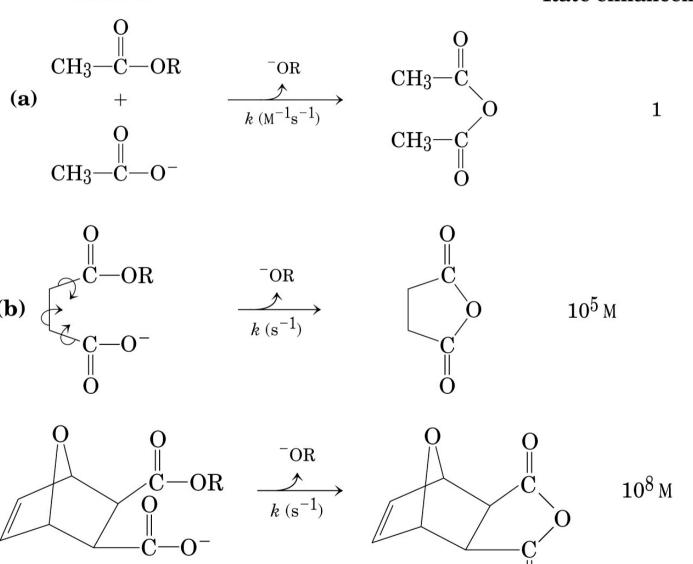
Catalisi per riduzione entropica

(c)



Catalisi per riduzione entropica

(c)



Catalisi acido base generale

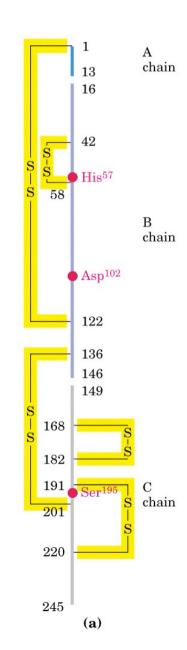
Amino acid residues	General acid form (proton donor)	General base form (proton acceptor)
Glu, Asp	R—COOH	R—COO-
Lys, Arg	${\rm R}\overset{H}{\overset{+}{{{{{{}{{}{{}{\overset$	$\overset{\cdots}{\mathrm{R-NH_2}}$
Cys	R—SH	$R-S^-$
His	R—C=CH HN NH	R—C=CH HN N:
Ser	R—OH	$R-O^-$
Tyr	R— OH	R — O^-

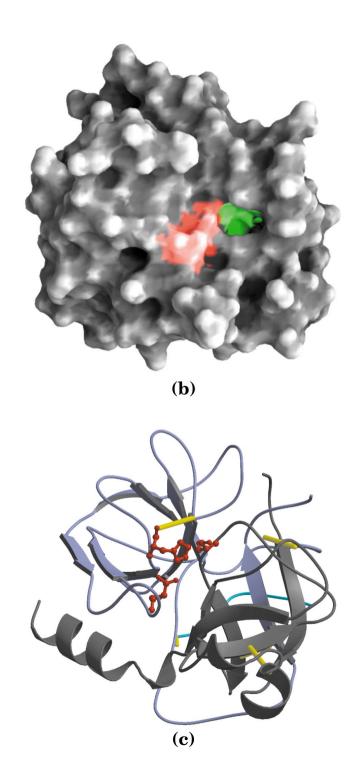
CATALISI COVALENTE

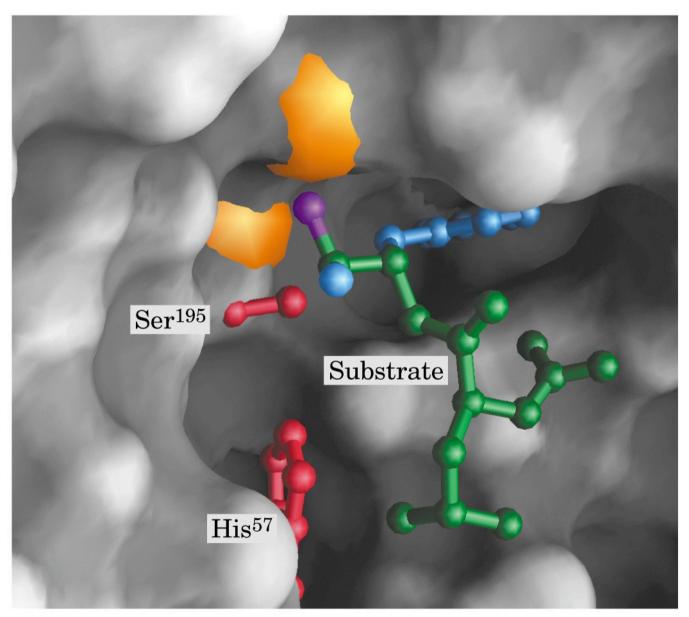
Si attua attraverso la formazione di un legame covalente transitorio tra una catena laterale amminoacidica dell' enzima o tra un cofattore enzimatico ed il substrato.

CHIMOTRIPSINA

PM 25.000 Da 3 CATENE 2 -S-S- INTER 3 -S-S-INTRA CATALISI: Covalente basica generale **AA CATALITICI** Ser 195 Asp 102 HIS 57 **FATTORE DI** ACCELERAZIONE: 10⁹







Nucleophiles

Electrophiles



Negatively charged oxygen (as in an unprotonated hydroxyl group or an ionized carboxylic acid)

Negatively charged sulfhydryl

Carbanion

Uncharged amine group

Imidazole

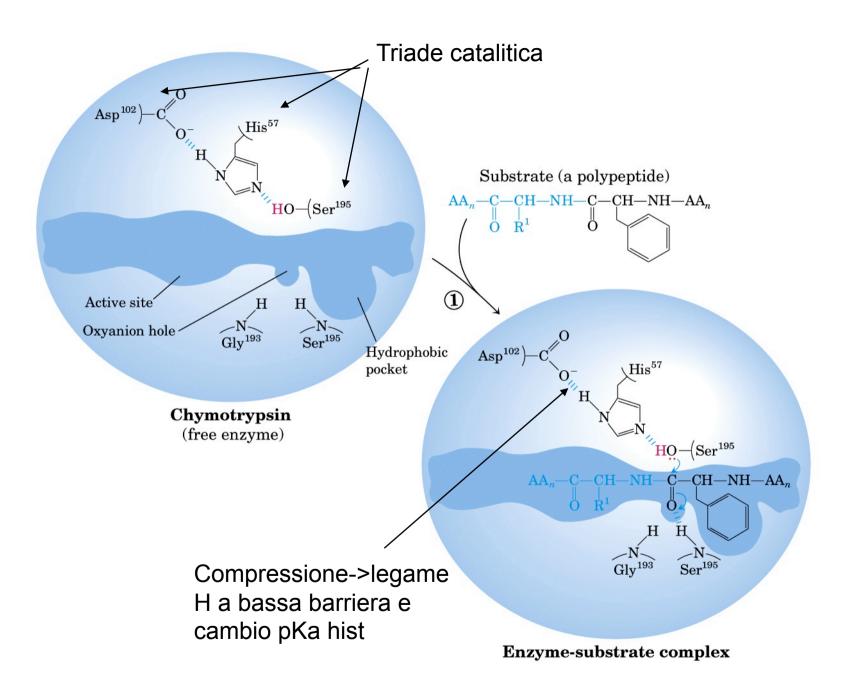
Hydroxide ion

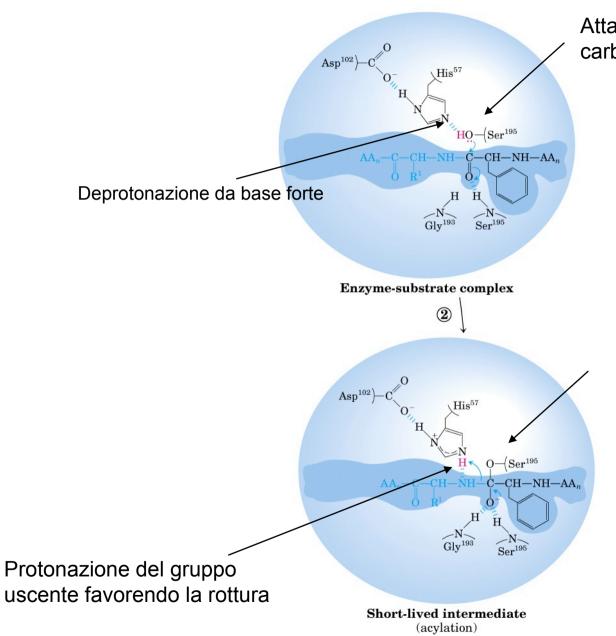
Carbon atom of a carbonyl group (the more electronegative oxygen of the carbonyl group pulls electrons away from the carbon)

Pronated imine group (activated for nucleophilic attack at the carbon by protonation of the imine)

Phosphorus of a phosphate group

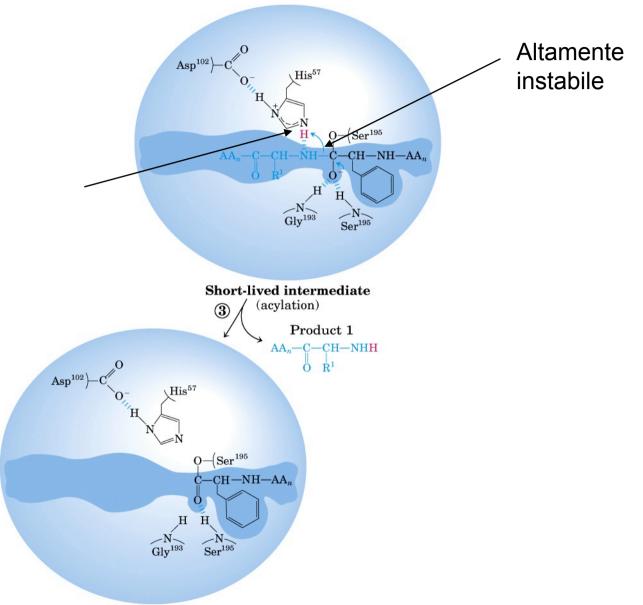
Proton





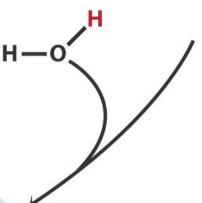
Attacco dello ione alcossido al carbonile

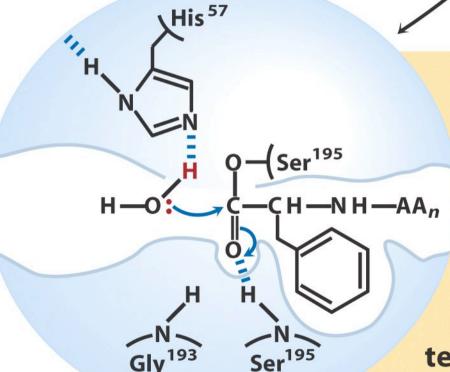
Formazione del l'intermedio tetraedrico con carica negativa stabilizzata da ponti H (i.e.energia legame per catalisi)



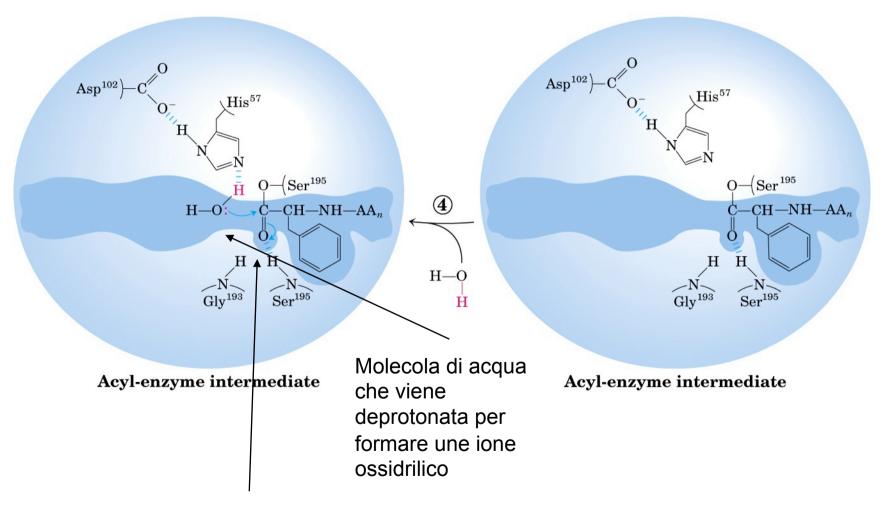
Acyl-enzyme intermediate







An incoming water molecule is deprotonated by general base catalysis, generating a strongly nucleophilic hydroxide ion. Attack of hydroxide on the ester linkage of the acyl-enzyme generates a second tetrahedral intermediate, with oxygen in the oxyanion hole again taking on a negative charge.

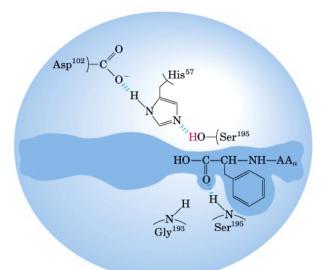


Presenza forte nucleofilo porta nuovamente a carica negativa

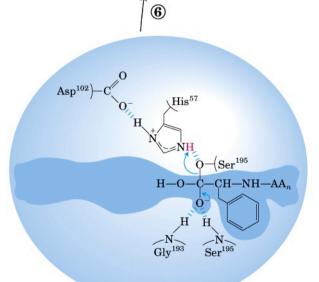
Short-lived intermediate (deacylation)

His 57 Ser 195 $-CH-NH-AA_n$

Collapse of the tetrahedral Gly 193 Se intermediate forms the second product, a carbohydrate anion, and displaces Ser 195.

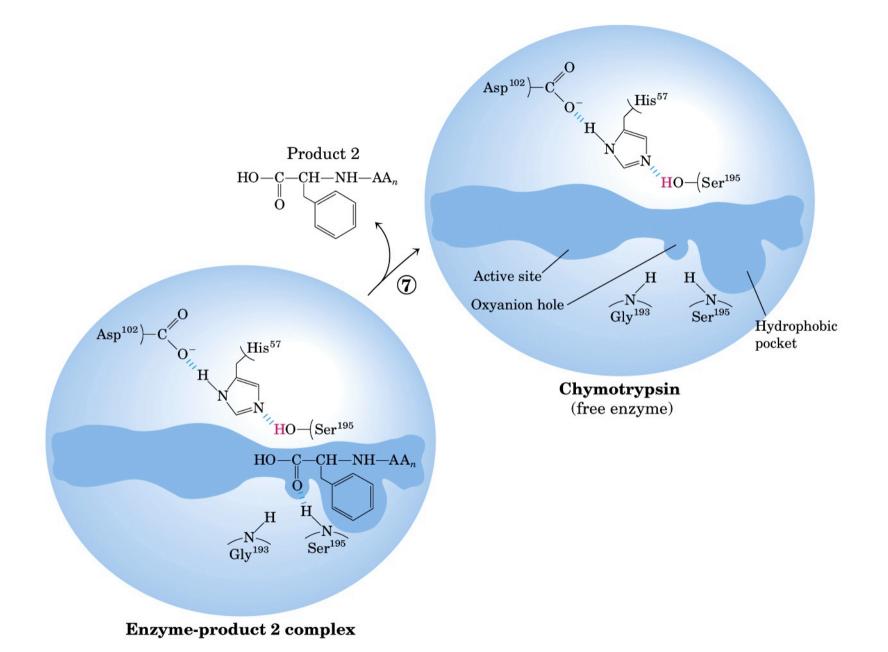


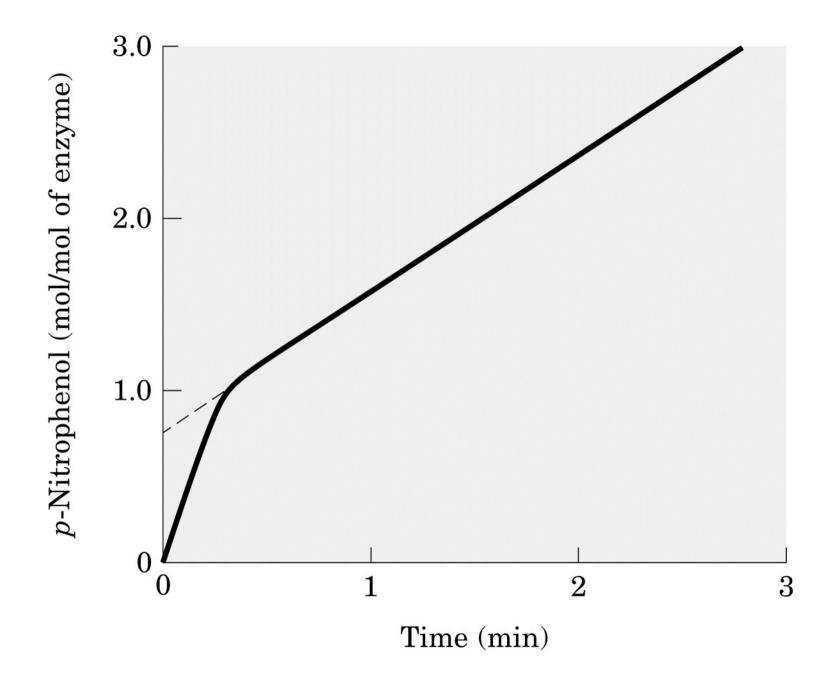
Enzyme-product 2 complex

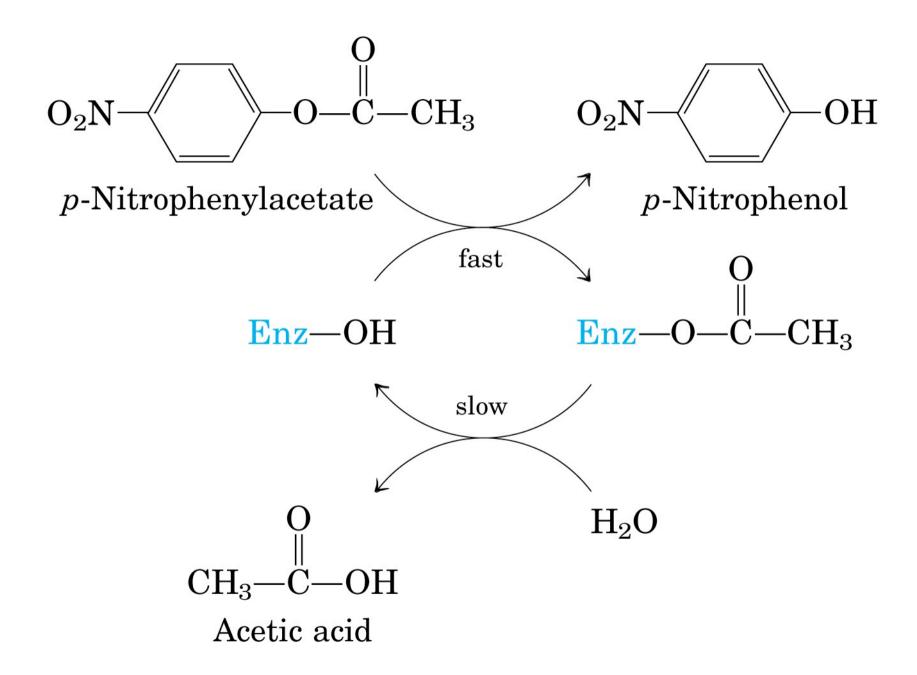


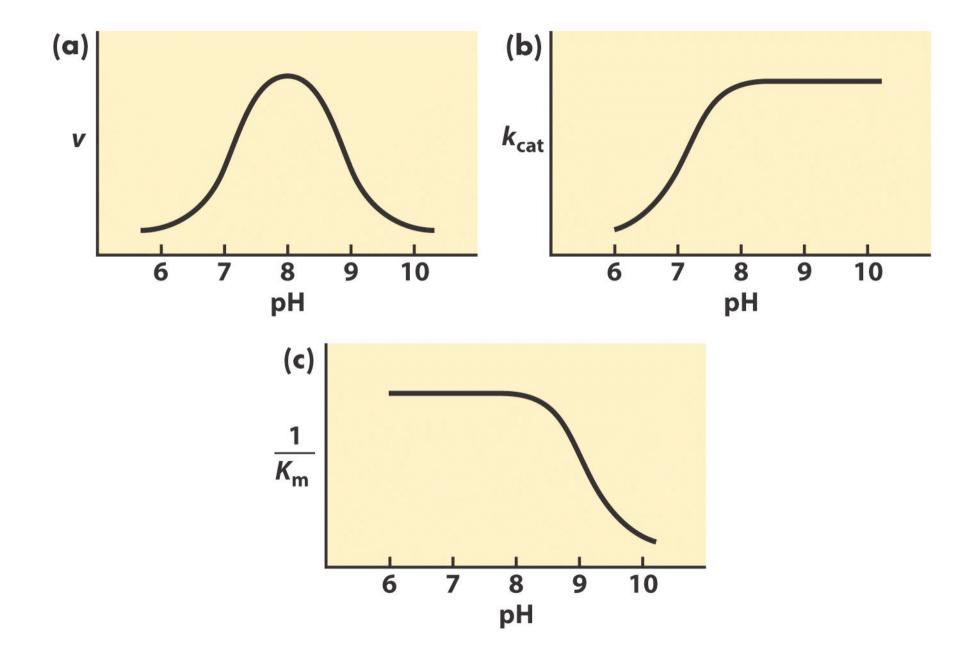
Ripetizione del primo ciclo di deprotonazione

Short-lived intermediate (deacylation)



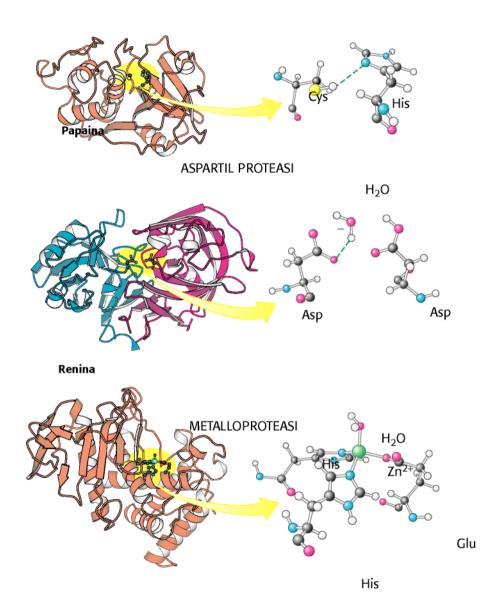






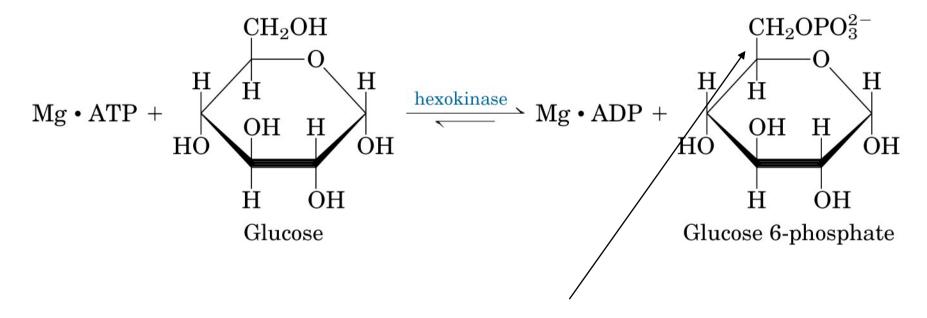
- Cistein-proteasi
- Aspartil-proteasi
- Metalloproteasi

CISTEINA PROTEASI



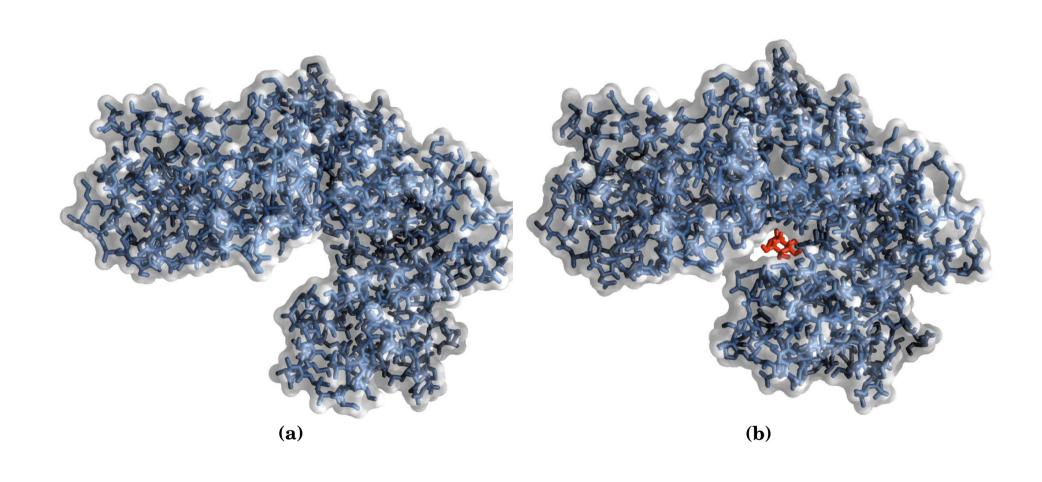
Termolisina

ESOCHINASI

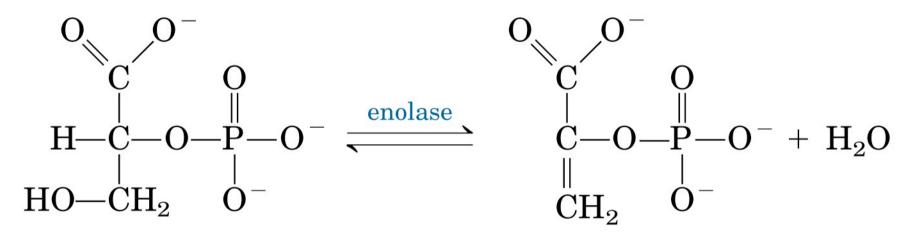


Gruppo ossidrilico sul C6 ha reattività simile all'acqua, ma c'e specificità di 10E6

ADATTAMENTO INDOTTO



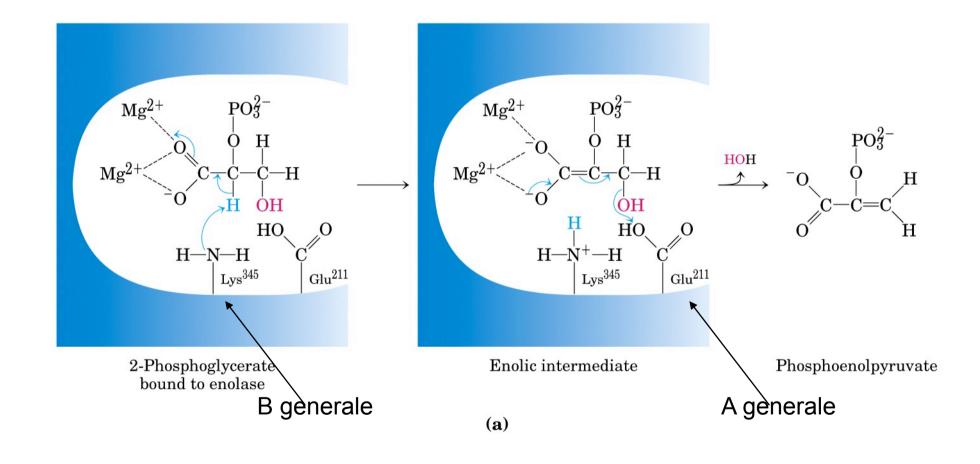
ENOLASI



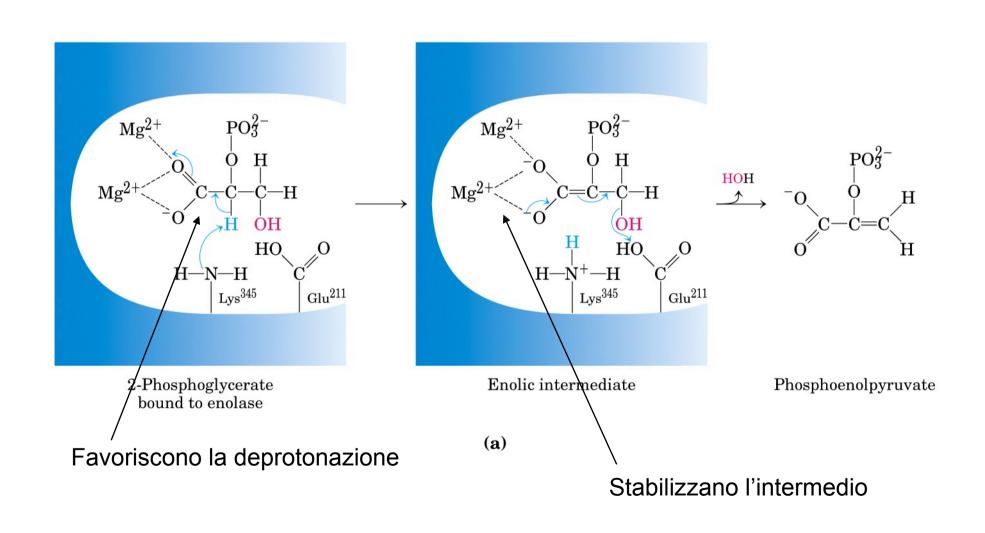
2-Phosphoglycerate

Phosphoenolpyruvate

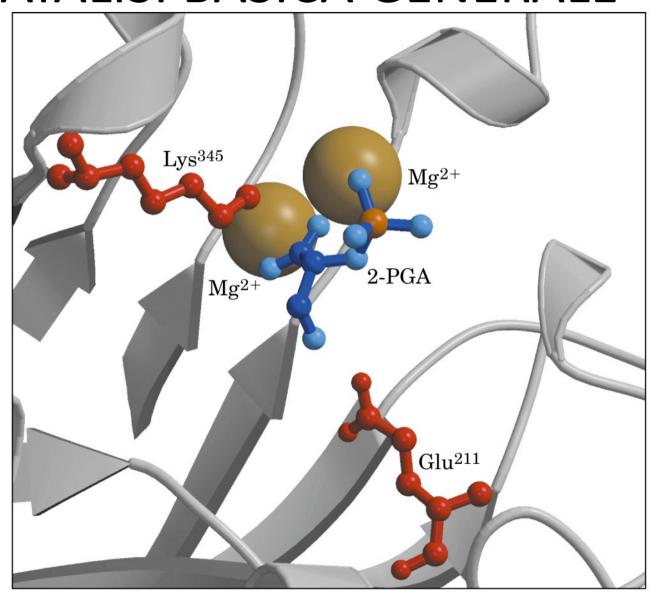
Catalisi da ioni metallici Acido-base generale Stabilizzazione stato transizione



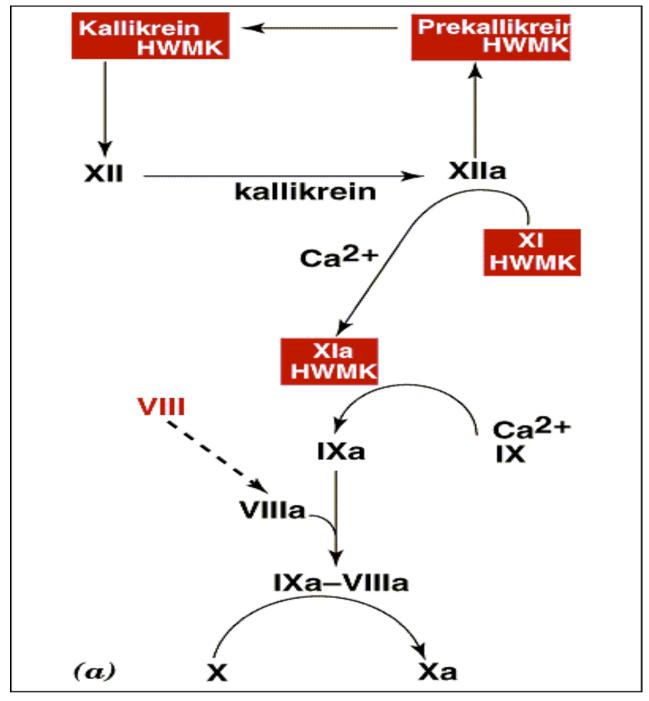
CATALISI DA IONI METALLICI



CATALISI BASICA GENERALE

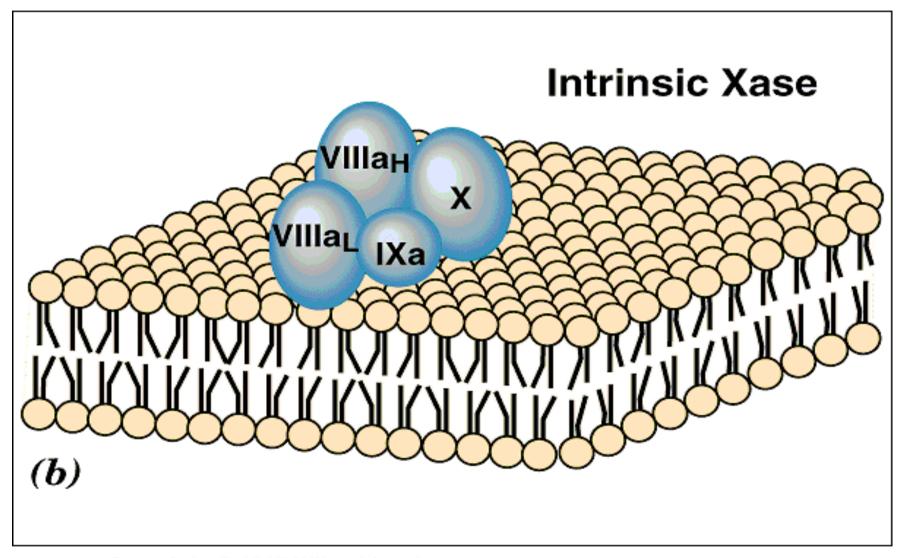


VIA INTRINSECA DELLA COAGULAZIONE

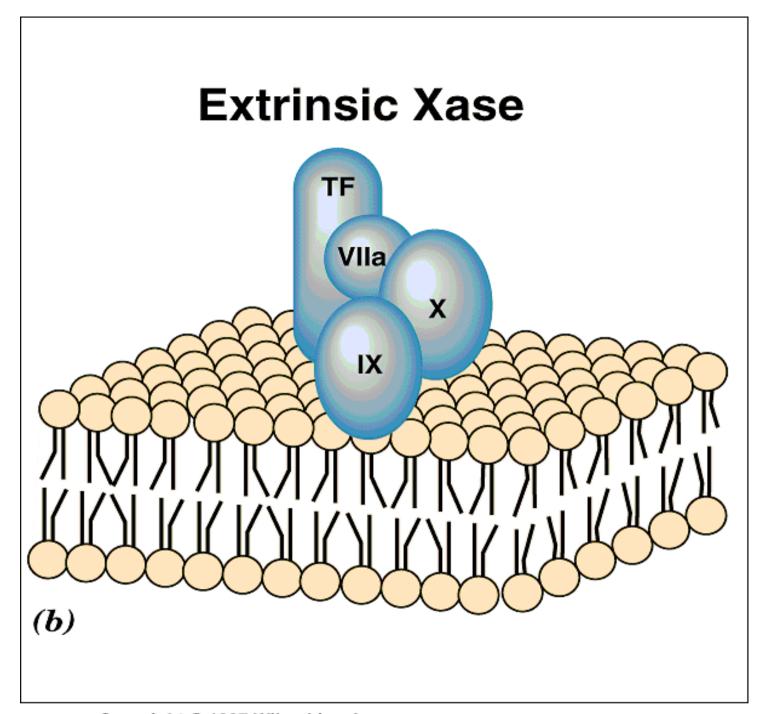


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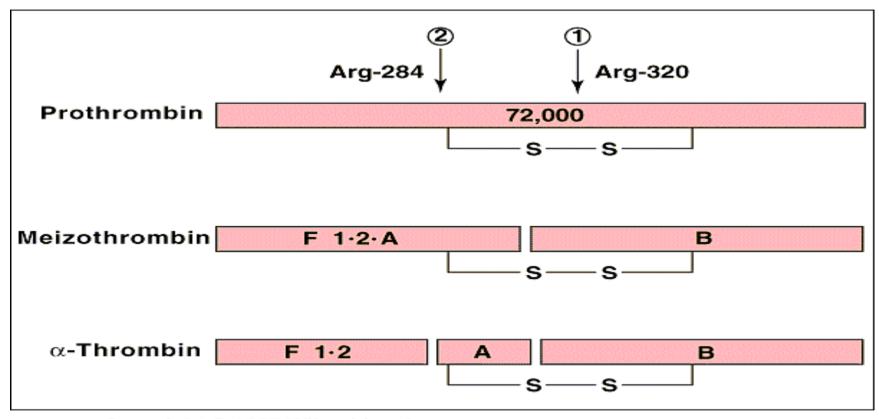
ATTIVAZIONE del FATTOR X



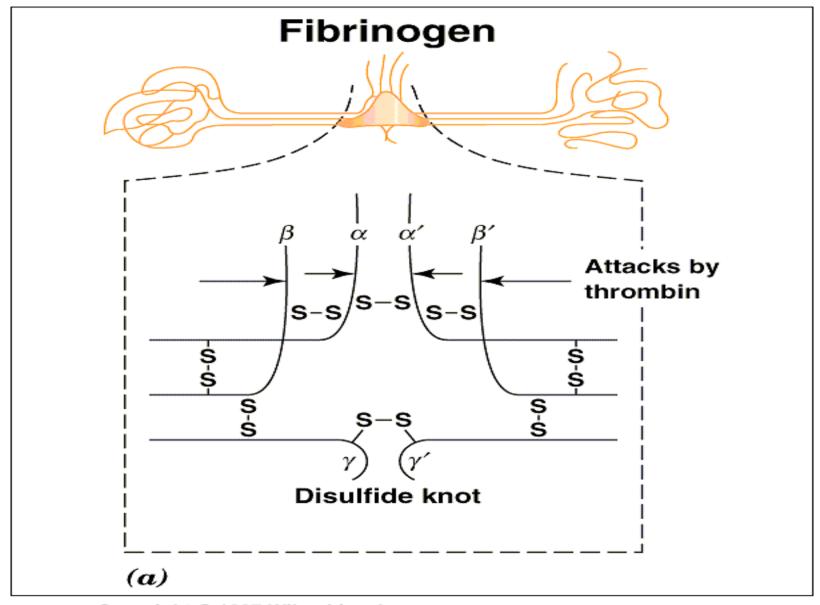
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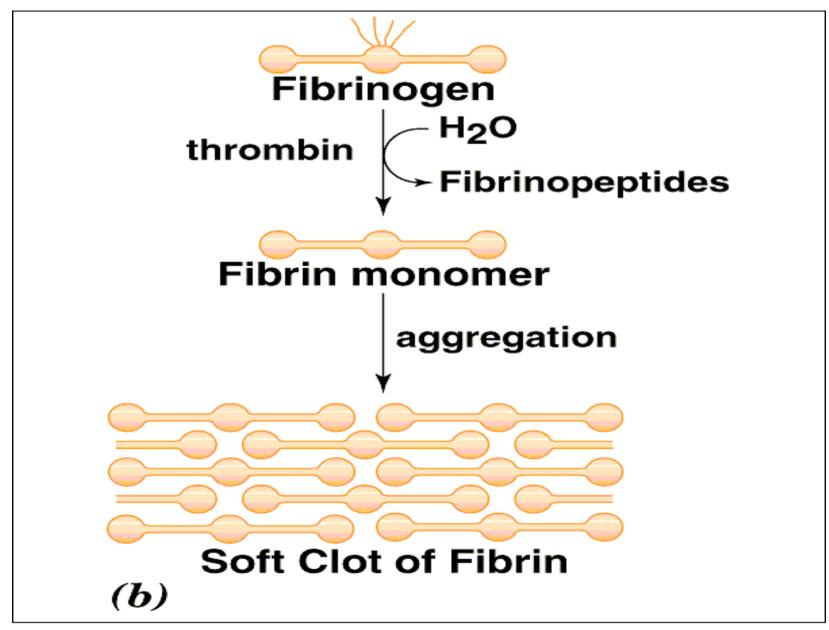
ATTIVAZIONE PROTEOLITICA DI TROMBINA



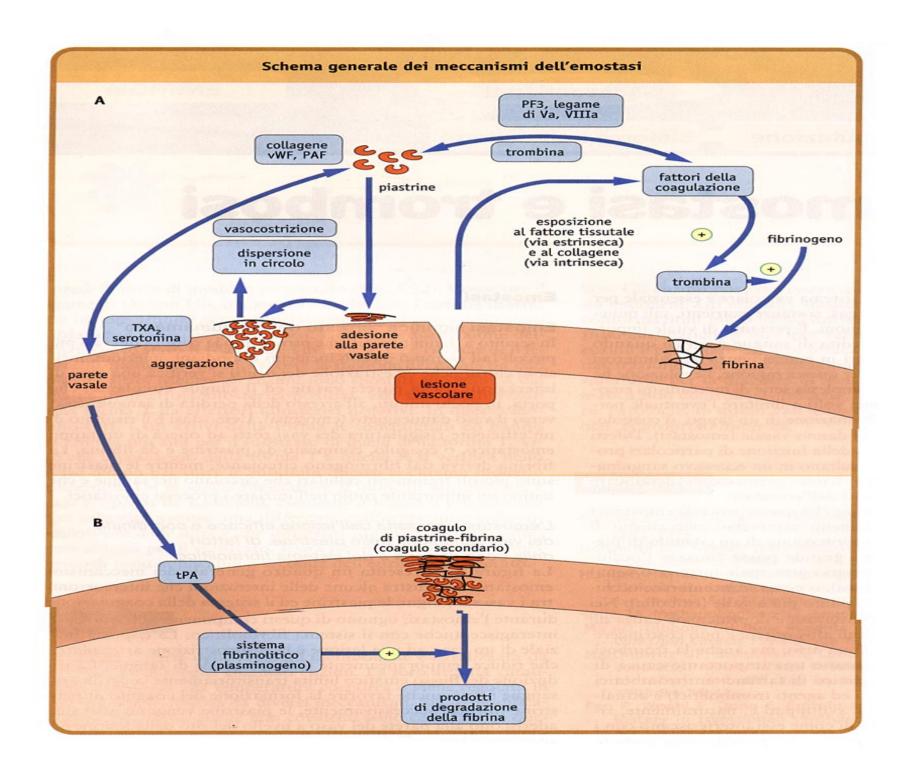
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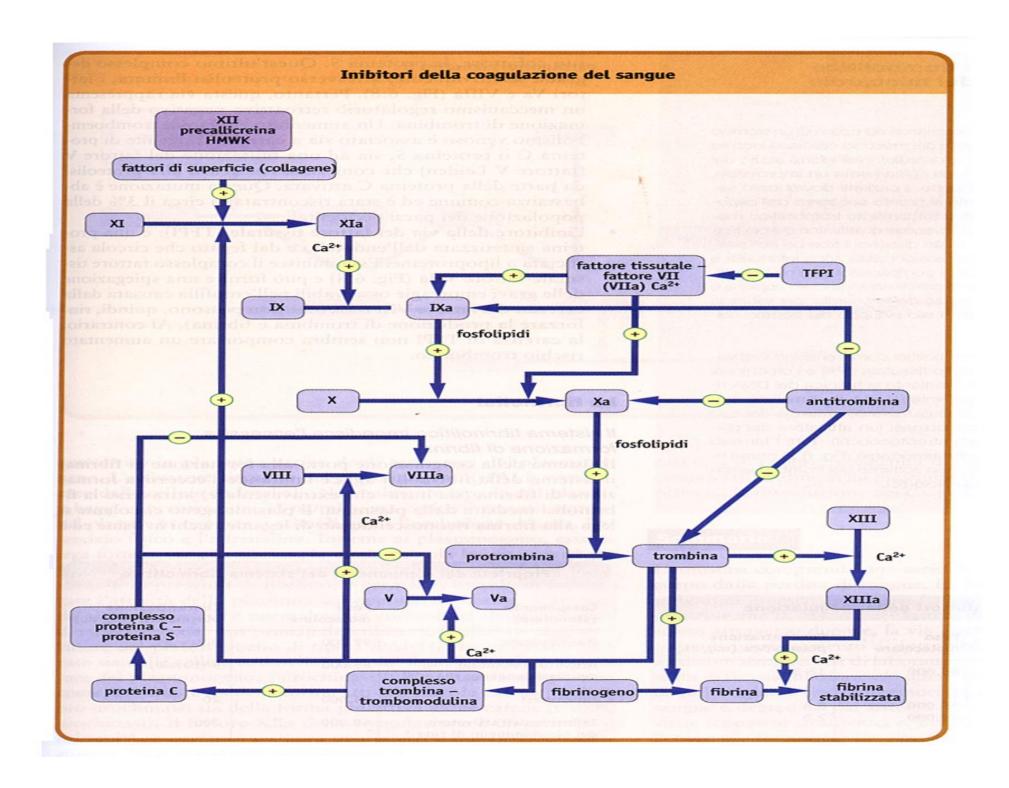


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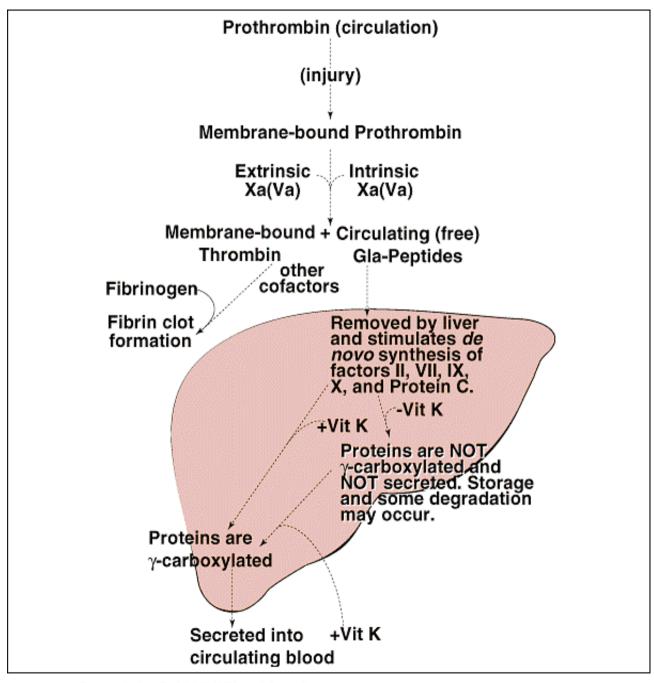


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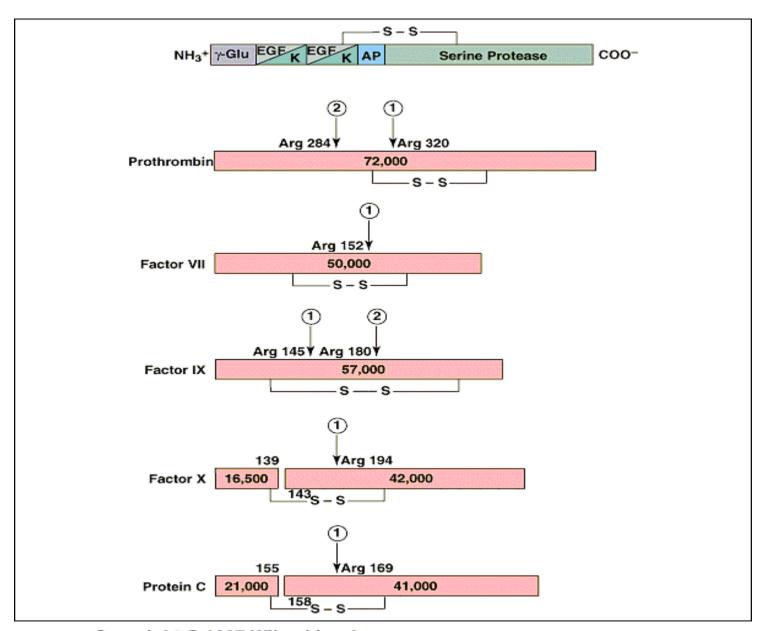




Ruolo dei residui di ac. γ carbossiglutammico nella regolazione della sintesi de novo dei fattori della coagulazione

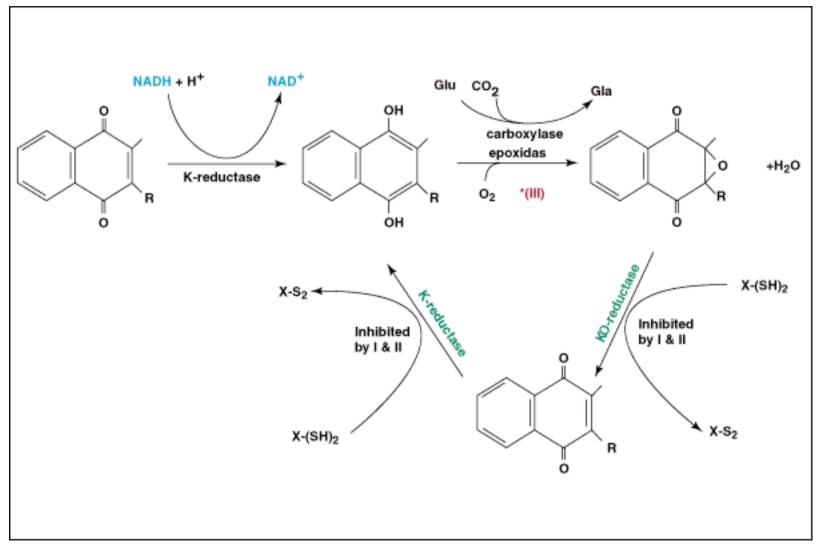


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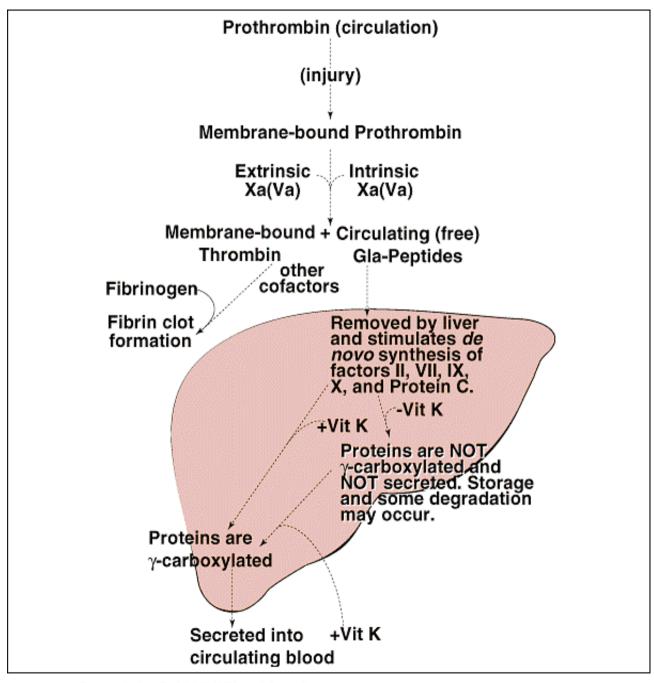
CICLO DELLA VITAMINA K



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Ruolo dei residui di ac. γ carbossiglutammico nella regolazione della sintesi de novo dei fattori della coagulazione



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