



Corso di laurea in Scienze Biologiche

Corso di laurea magistrale in Scienze Biomolecolari e dell'Evoluzione

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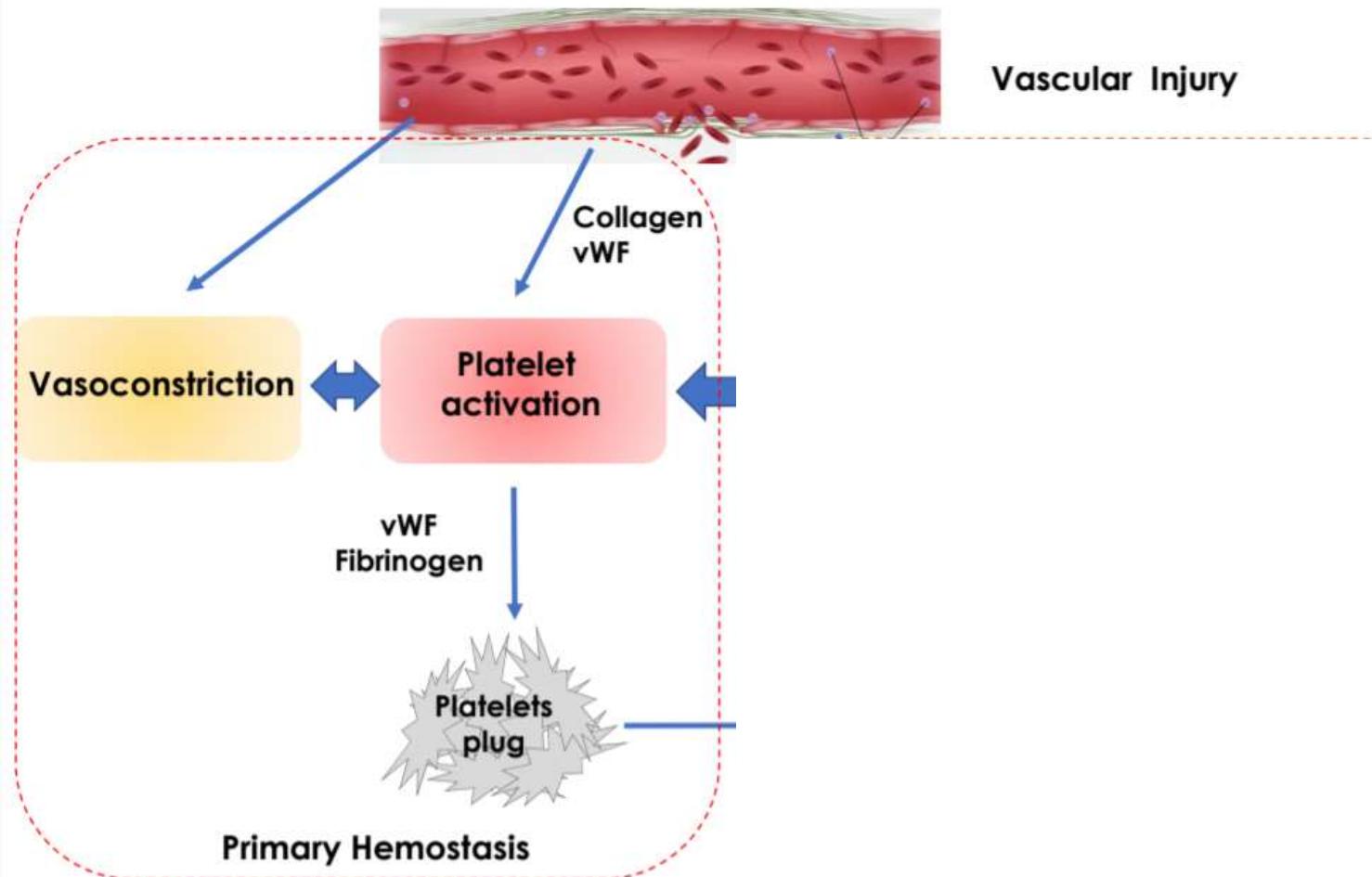
Coagulation Cascade

Activation and specificity of

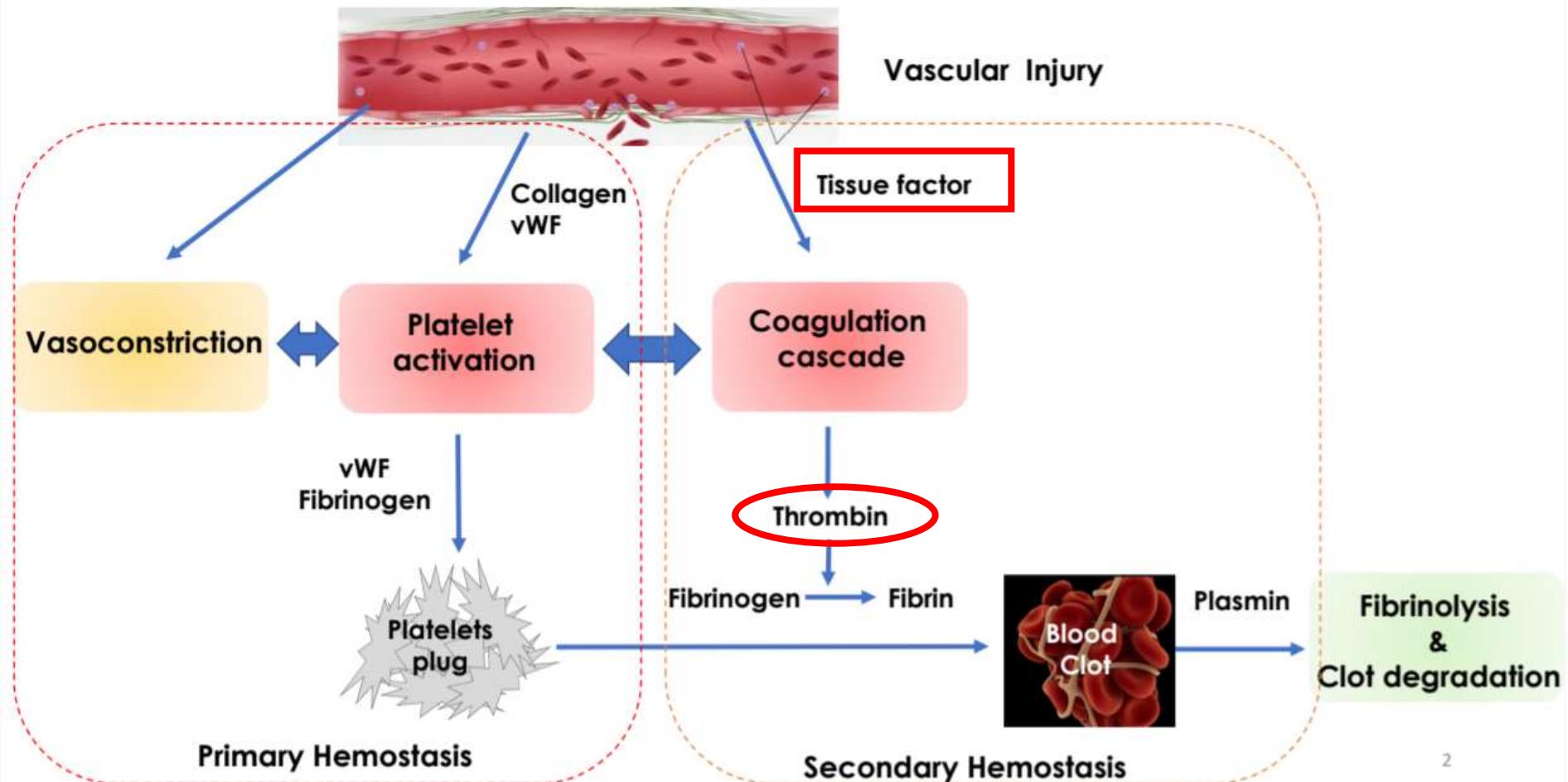
Thrombin

Da Giulia Pavani
Sara Calzavarini

Hemostatic process: an overview

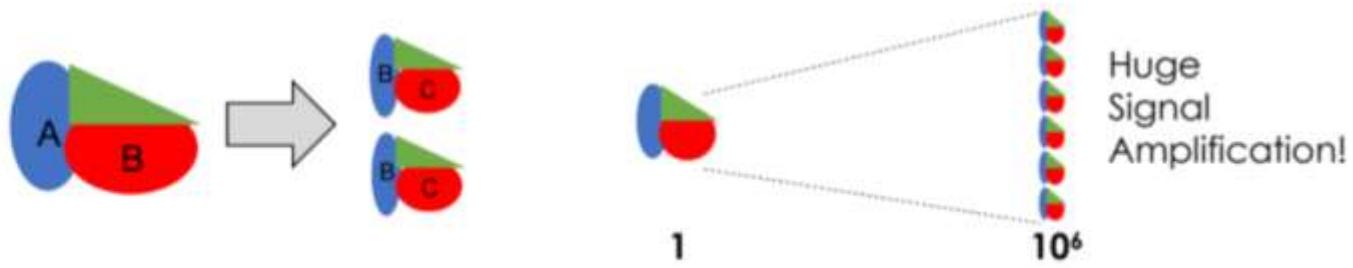


Hemostatic process: an overview



The cascade organization

Consequential enzymatic conversions of zymogens to activated enzymes

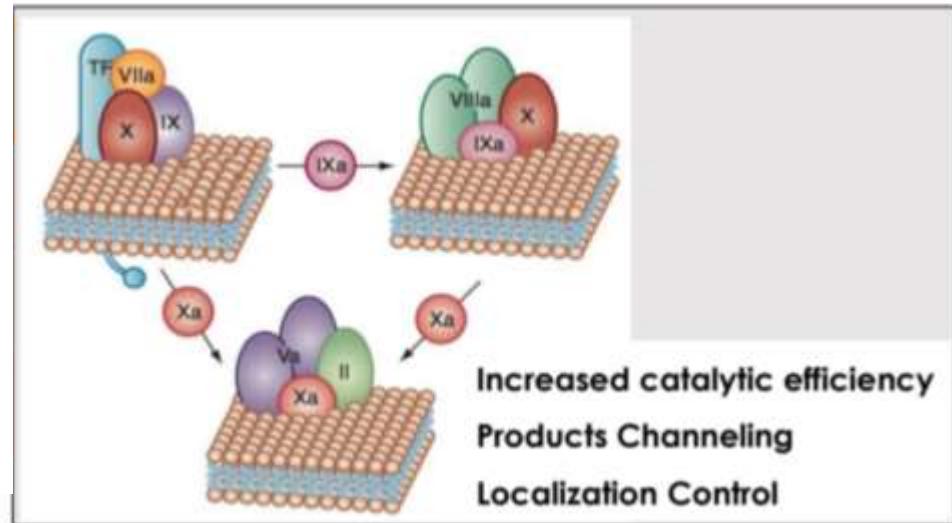
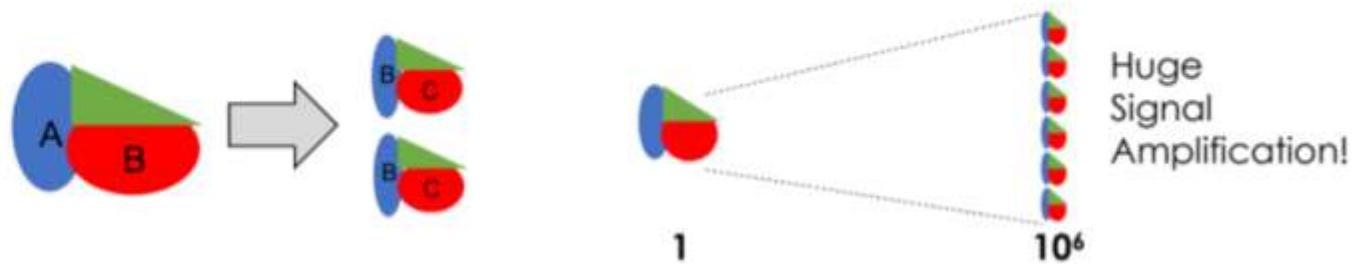


Clotting factor number	Clotting factor name	Plasma concentration (mg/L)
I	Fibrinogen	3000
II	Prothrombin	100
III	TF Tissue Factor	-
IV	Calcium	-
V	Proaccelerin, labile factor	10
VII	Stable factor, proconvertin	0.5
VIII	Antihemophilic factor A	0.1
IX	Antihemophilic factor B or Christmas factor	5

Clotting factor number	Clotting factor name	Plasma half-life (h)	Plasma concentration (mg/L)
I	Fibrinogen	90	3000
II	Prothrombin	65	100
III	TF Tissue Factor	-	-
		-	-
V	Proaccelerin, labile factor	15	10
VII	Stable factor, proconvertin	5	0.5
VIII	Antihaemophilic factor A	10	0.1
IX	Antihaemophilic factor B or Christmas factor	25	5

The cascade organization

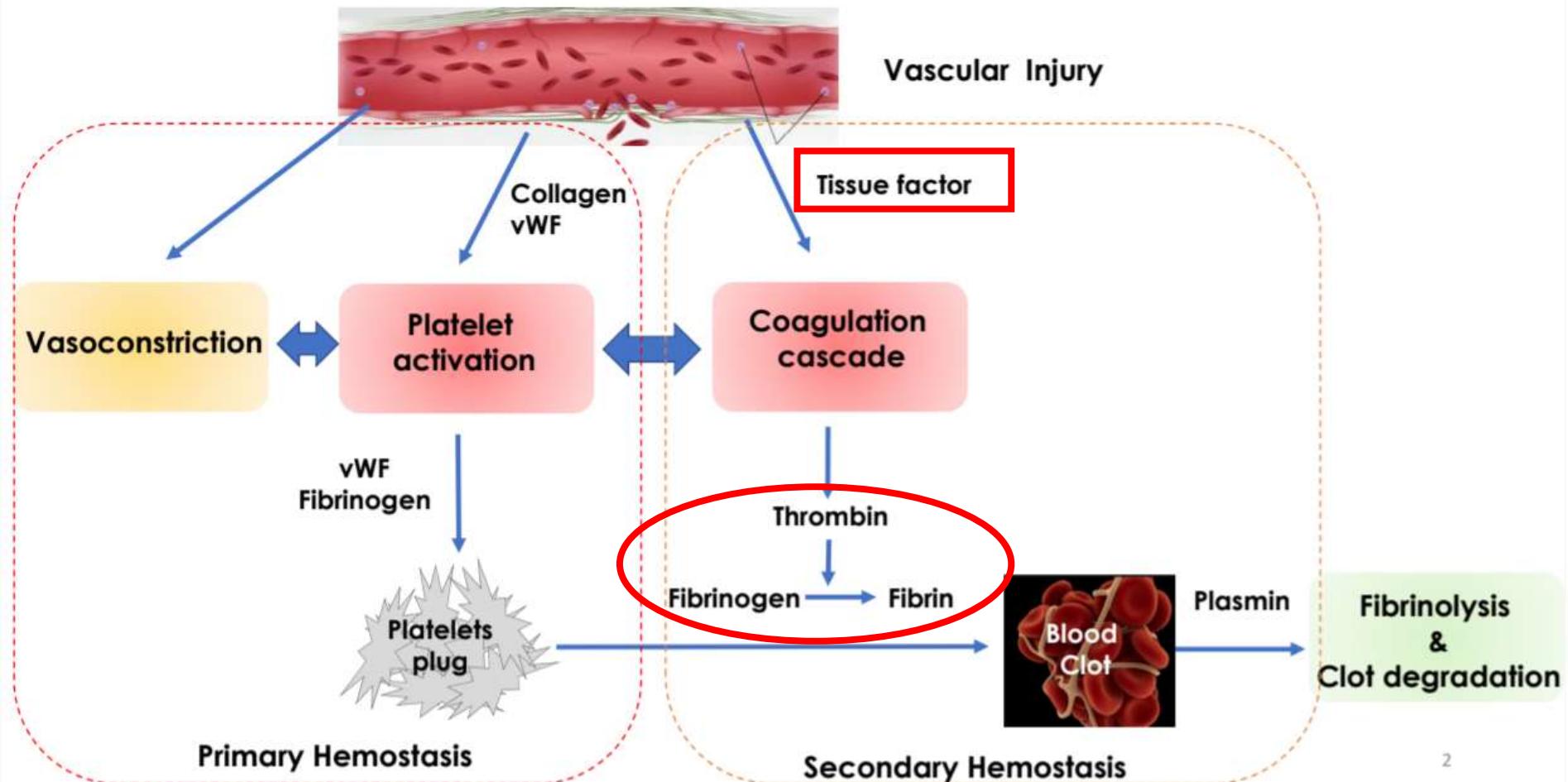
Consequential enzymatic conversions of zymogens to activated enzymes



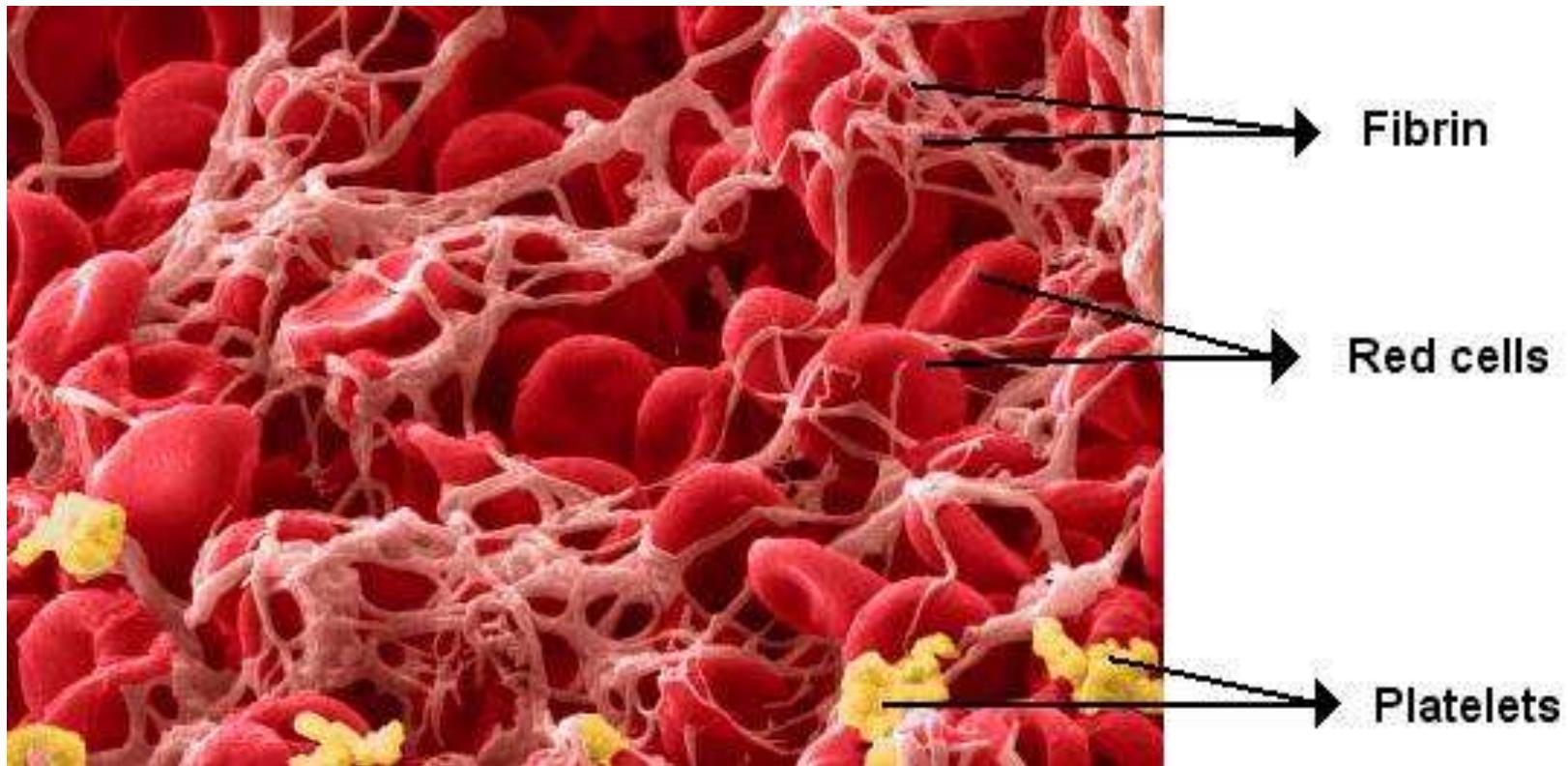
Attivazione del Fibrinogeno a Fibrina

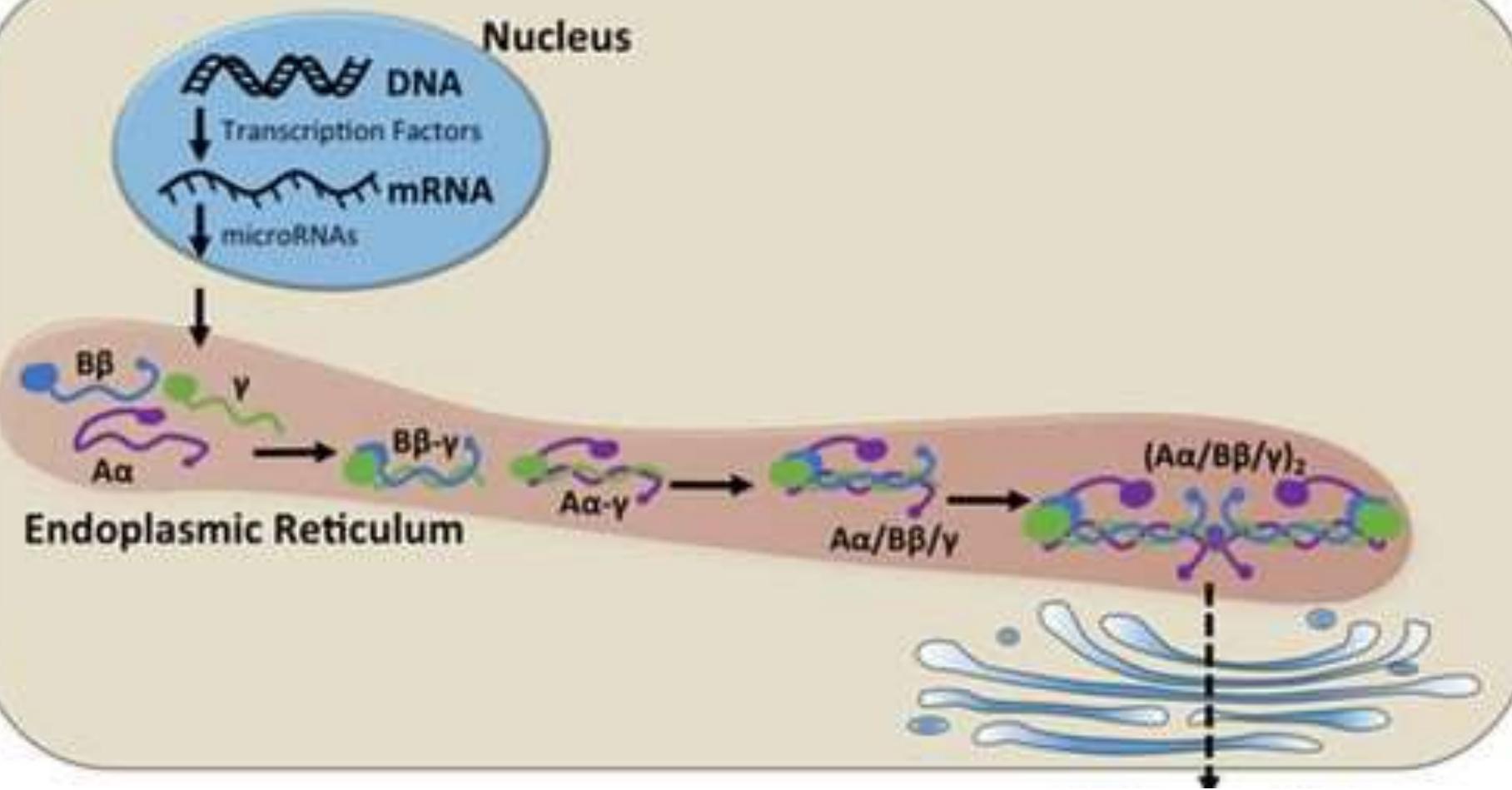
Il collante del Coagulo

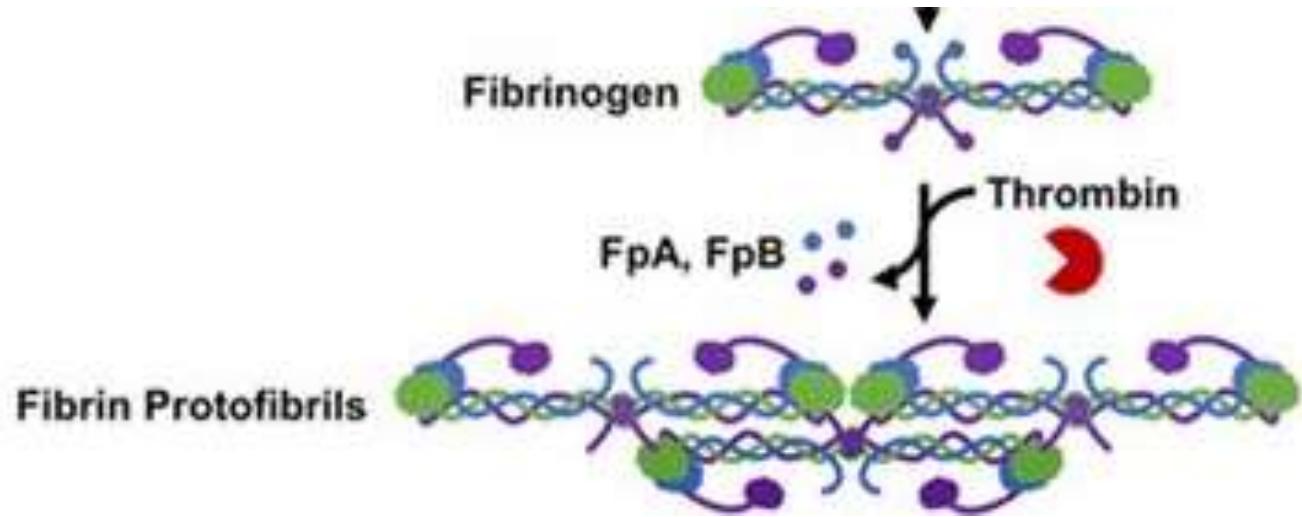
Hemostatic process: an overview



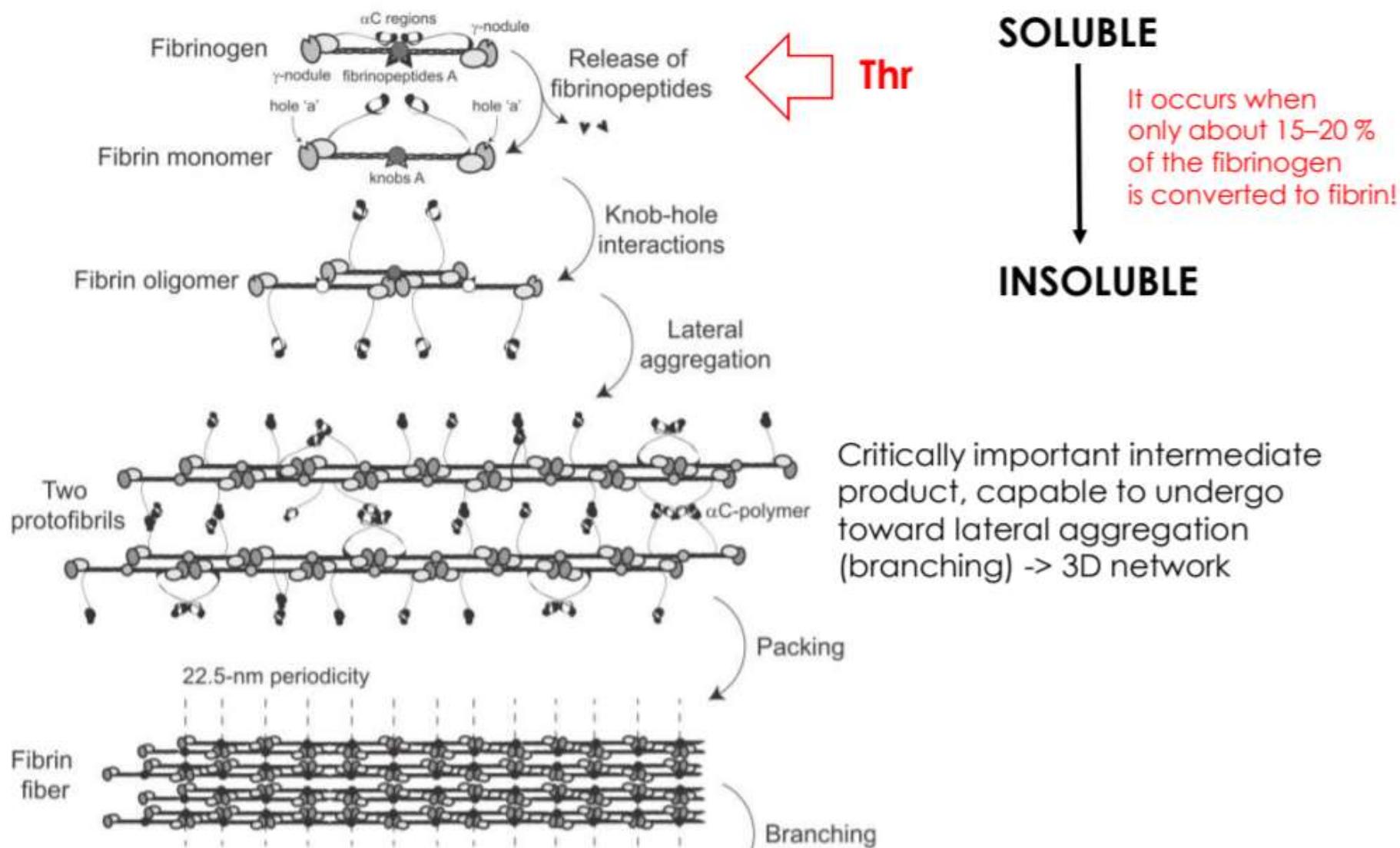
Thrombin cleavage of the plasma protein fibrinogen



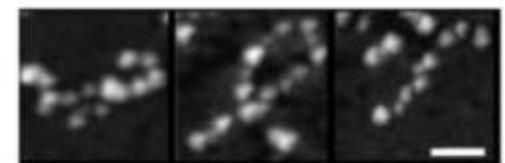
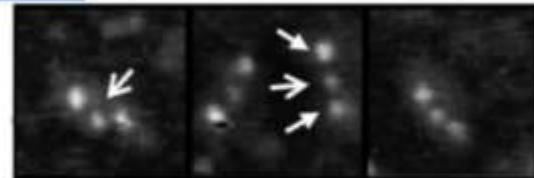
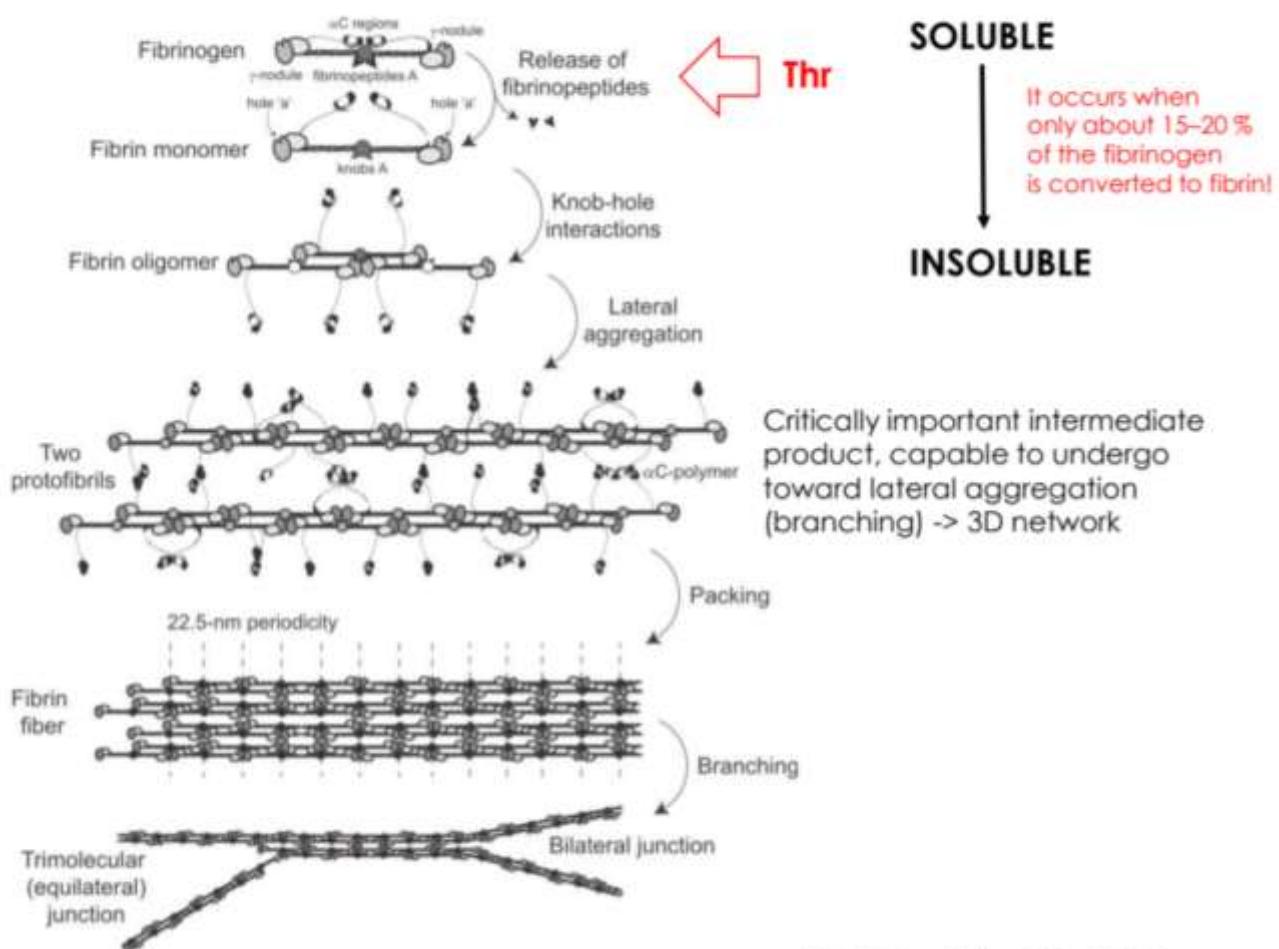




Fibrinogen to fibrin

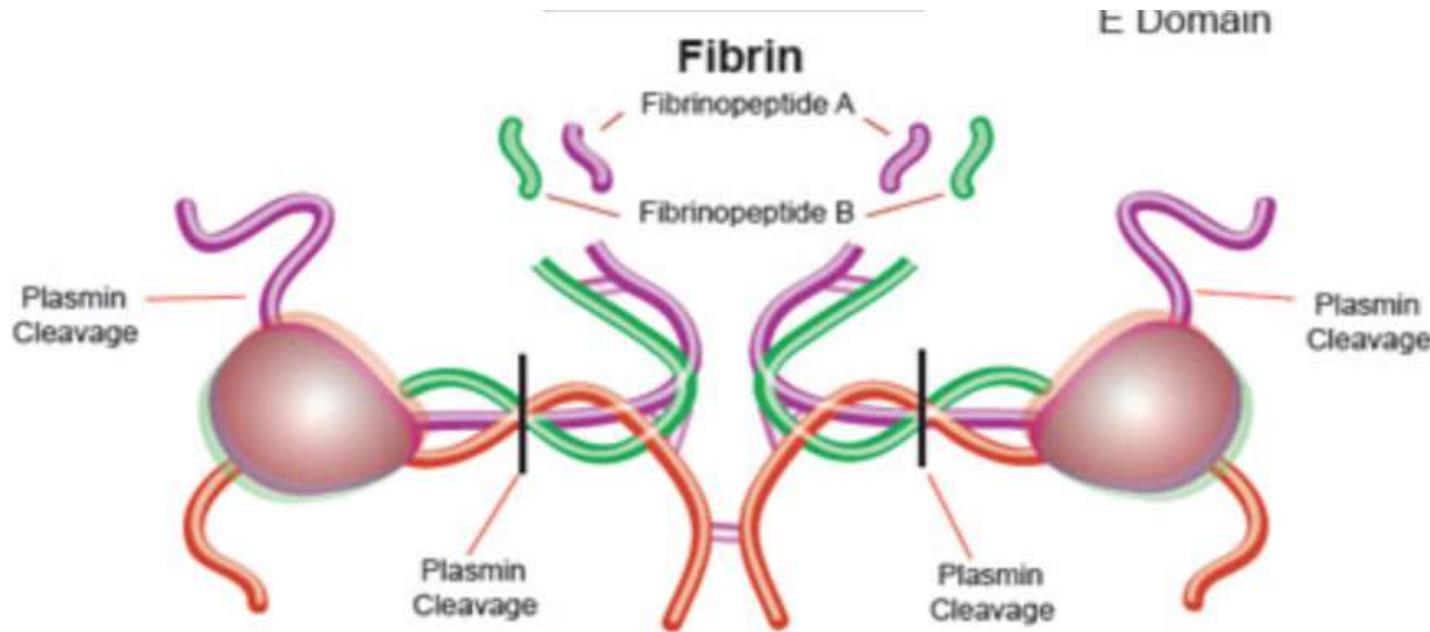


Fibrinogen to fibrin



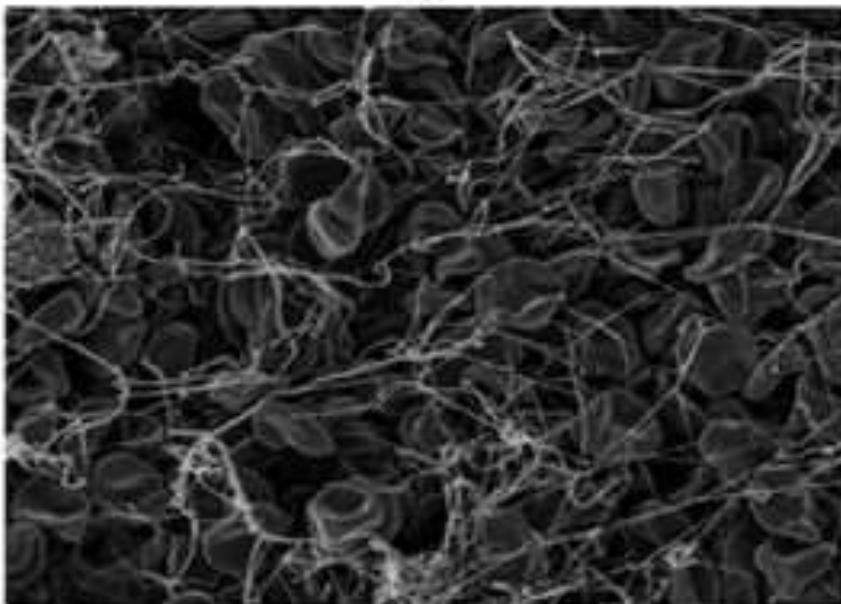
Weisel JW and Litvinov RI Blood 20013

Thrombin /Plasmin cleavage of the plasma protein fibrinogen/fibrin



Modifiers of Fibrin Clot Formation, Structure, and Stability

Concentrations of:	Metal ions	Blood cells	Polyphosphates	Post-translational
Procoagulants	pH	Vascular cells	DNA & histones	modification
Anticoagulants	Temperature	Cellular vesicles	Heparin	Blood flow
Fibrinogen variants			Protamine	Others?



Diseases Associated with Abnormal Fibrin(ogen) Structure and Stability

Coronary Artery Disease

Myocardial Infarction

Ischemic Stroke

Venous Thromboembolism

Abdominal Aortic Aneurysm

Smoking

Chronic Kidney Disease

In-stent Thrombosis

Cirrhosis

Hemophilia

Others?



The Blood Coagulation Response:

Various Cells (endothelium, platelets, etc.)

Hidden Cofactors (tissue factor)

Zymogens (FXI, FVII, FIX, FX, II, etc.)

Procofactors (FVIII and FV)

Quiescent State

Other Functions

Injury

Activated Cells

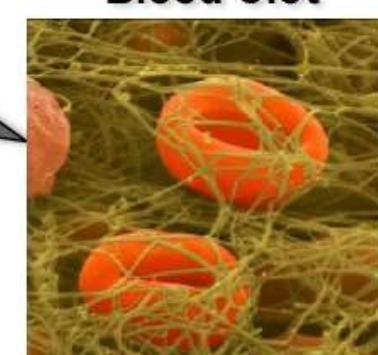
Exposed TF

Proteases

Cofactors

Initiation of Coagulation

Thrombin



astrographics.com; Dennis Kunkel

Bleeding
(Hemophilia)

Pathology

Too Little

Too Much

Thrombosis

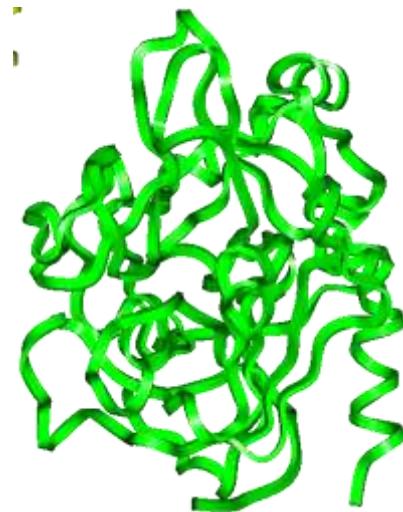
SPECIFICITA E MOLTEPLICITA SUBSTRATI

Trypsin and Thrombin have similar structures

Trypsin



Thrombin

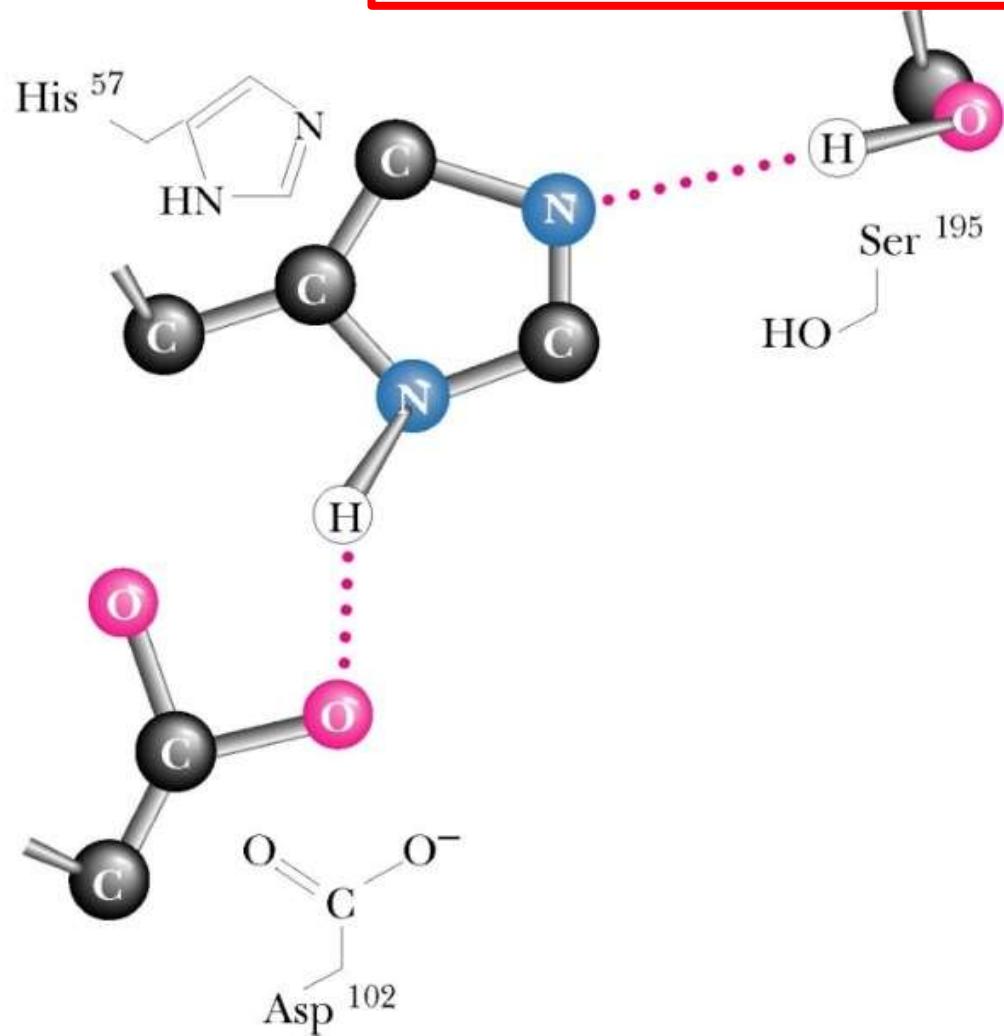
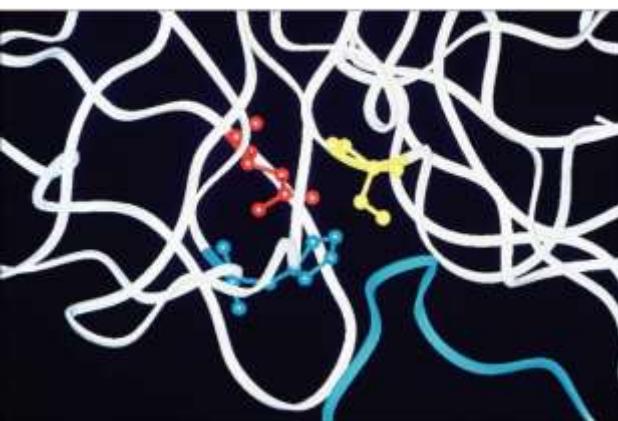


Serina proteasi

classe di enzimi proteolitici il cui meccanismo catalitico è basato sulla presenza di un residuo di serina nel sito attivo.

- tripsina
 - chimotripsina
 - elastasi
 - trombina
 - plasmina
 - attivatore tissutale del plasminogeno
- } enzimi digestivi sintetizzati nel pancreas e secreti nell'apparato digerente come proenzimi inattivi o zimogeni
- acetil colinesterasi= non è una proteasi ma una serina esterasi il cui meccanismo d'azione è correlabile a quello delle serina proteasi. Essa idrolizza il neurotrasmettitore acetilcolina nello spazio sinaptico interneuronico.

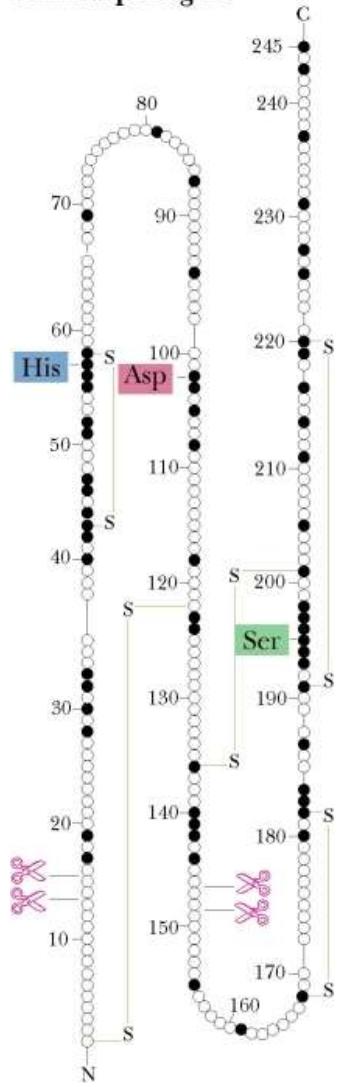
Serin-Proteasi



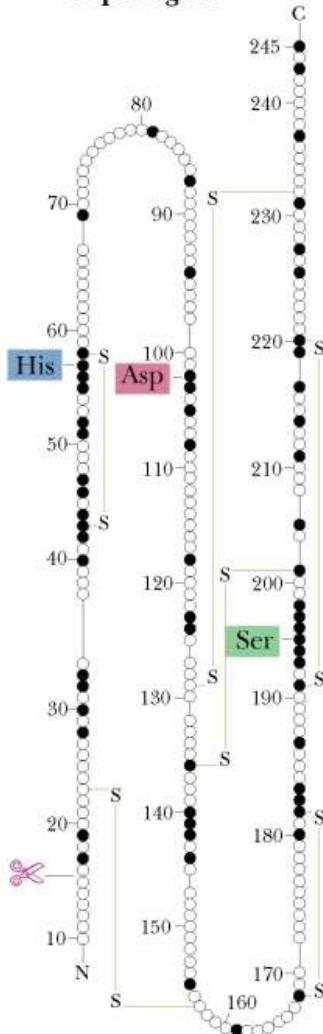
Tre residui polari (His⁵⁷, Asp¹⁰², Ser¹⁹⁵) formano la cosiddetta *triade catalitica* in corrispondenza del sito attivo il quale è costituito da una depressione sulla superficie dell'enzima.

Tripsina, chimotripsina ed elastasi catalizzano tutti la stessa reazione:
la scissione di una catena polipeptidica.

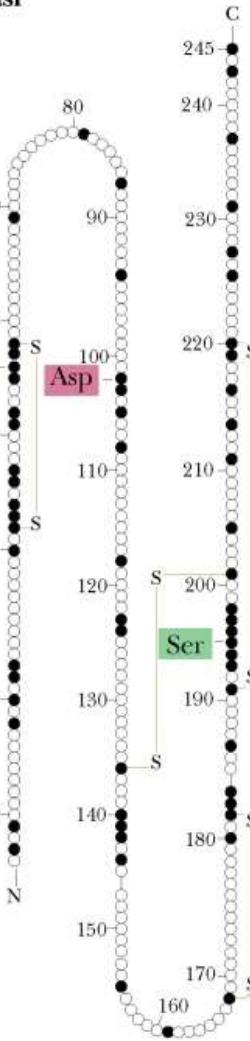
Chimotripsinogeno



Tripsinogeno



Elastasi



Tripsina: agisce su aa basici
(arginina, lisina)

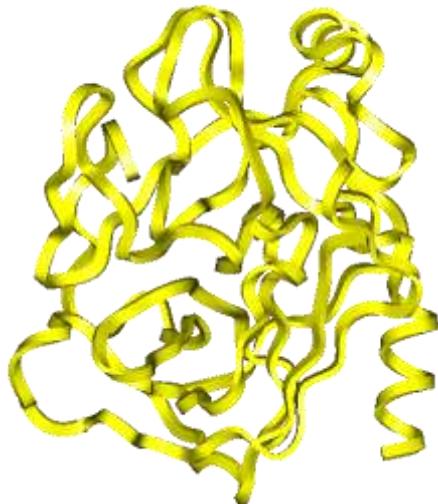
Chimotripsina: agisce su aa
aromatici

Elastasi: agisce su piccoli
residui di aa neutri

Trypsin and Thrombin have similar structures

Trypsin

- Cleaves peptides on the C-term of **Lys** and **Arg** amino acid residues



Thrombin

- Cleaves peptides at **Arg** (Pro, **Arg**, Ser/Ala/Gly/Thr, not acidic)



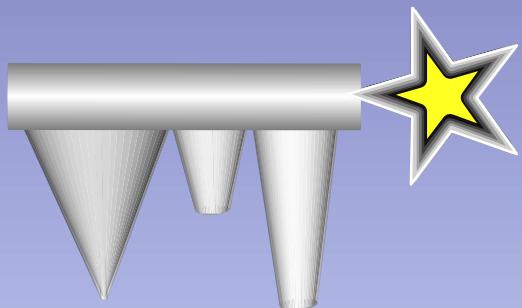
Thrombin cleaves different substrates

- Thrombin cleaves after Arg residues

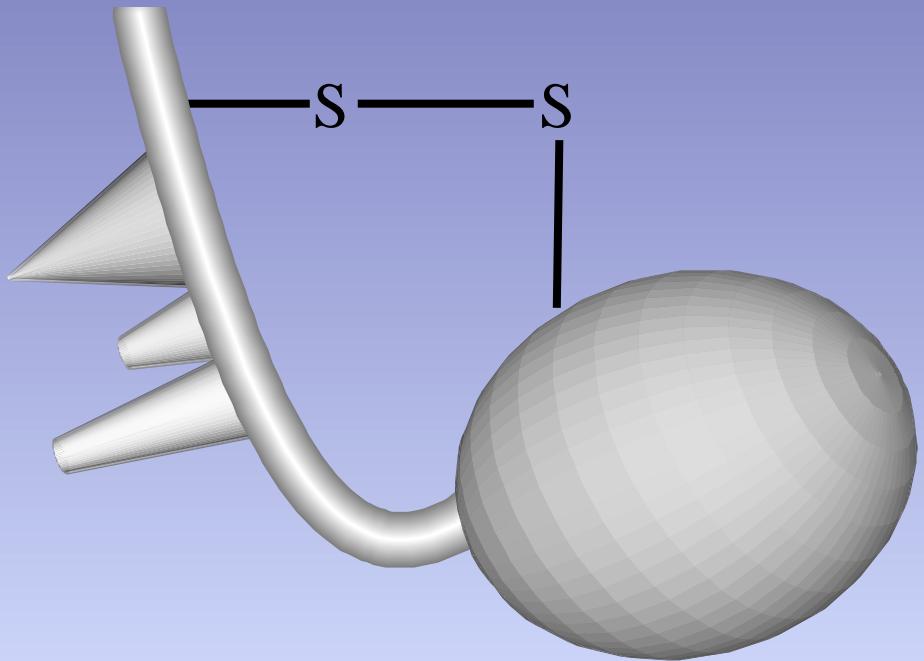
Cleavage Sites for Natural Thrombin Substrates



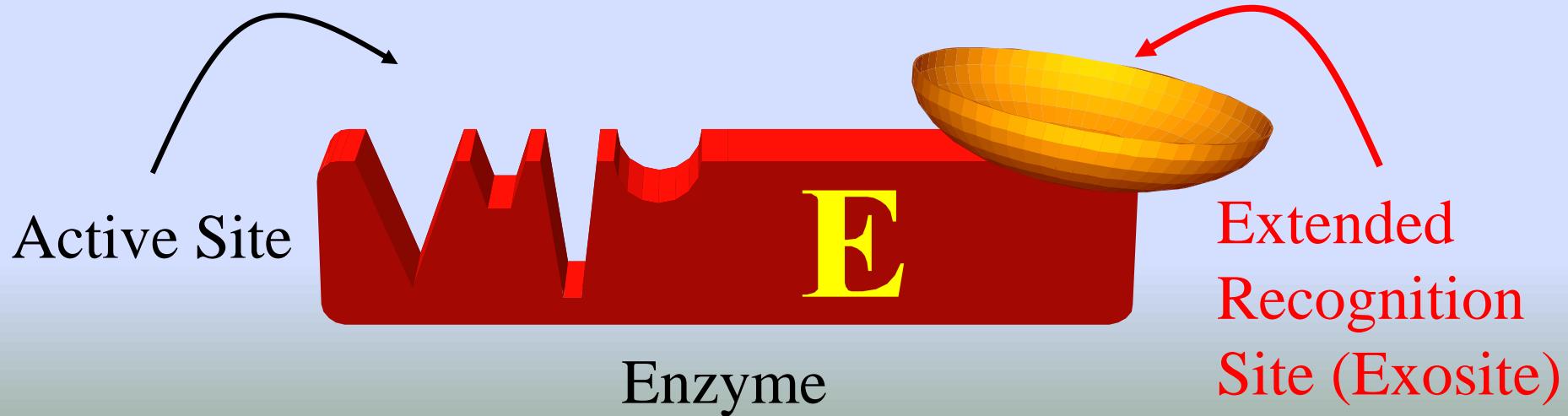
	P4	P3	P2	P1	P1'	P2'	P3'
Fibrinogen (A)	Gly	Gly	Val	Arg	Gly	Pro	Arg
Fibrinogen (B)	Phe	Ser	Ala	Arg	Gly	His	Arg
FV (709)	Leu	Gly	Ile	Arg	Ser	Phe	Arg
FV (1018)	Leu	Ser	Pro	Arg	Thr	Phe	His
FV (1545)	Trp	Tyr	Leu	Arg	Ser	Asn	Asn
FVIII (372)	Ile	Gln	Ile	Arg	Ser	Val	Ala
FVIII (740)	Ile	Glu	Pro	Arg	Ser	Phe	Ser
FVIII (1689)	Gln	Ser	Pro	Arg	Ser	Phe	Gln
FXIII	Gly	Val	Pro	Arg	Gly	Val	Asn
PAR1	Leu	Asp	Pro	Arg	Ser	Phe	Leu
PAR4	Pro	Ala	Pro	Arg	Gly	Tyr	Pro
FXI	Ile	Lys	Pro	Arg	Ile	Val	Gly
PC	Val	Asp	Pro	Arg	Leu	Ile	Asp
TAFI	Val	Ser	Pro	Arg	Ala	Ser	Ala
AT	Ile	Ala	Gly	Arg	Ser	Leu	Asn



Oligopeptidyl
Substrate



Protein Substrate

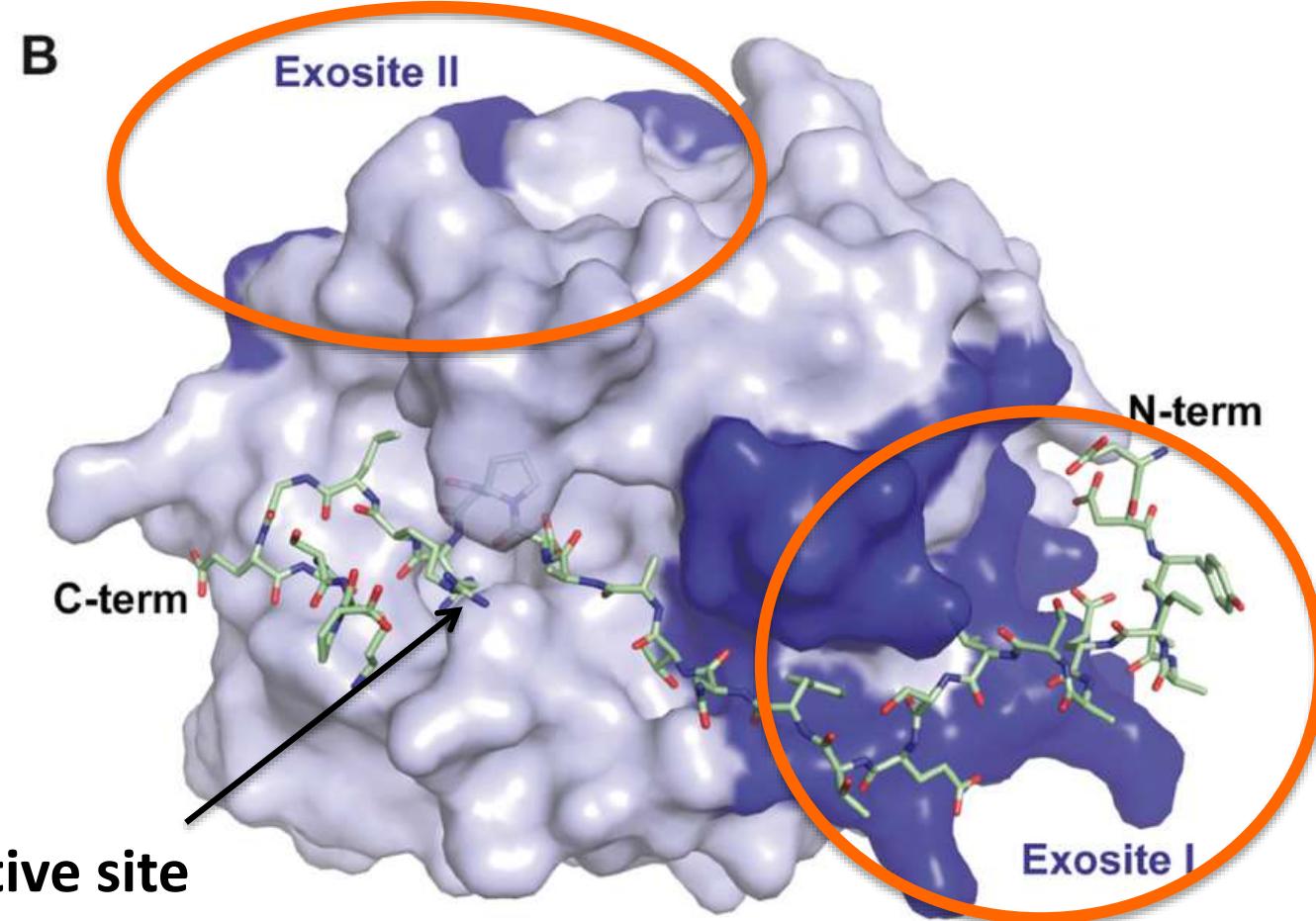


Active Site

Enzyme

Extended
Recognition
Site (Exosite)

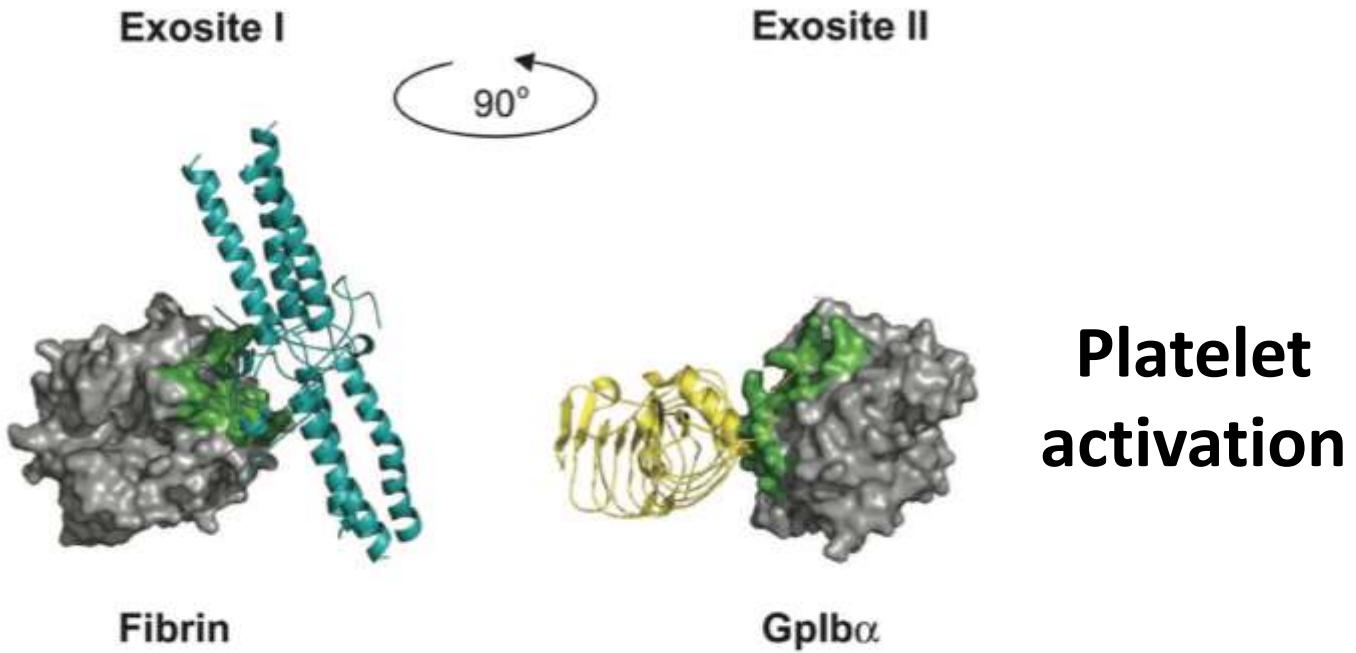
Thrombin X-ray structure



Exosite binding determines substrate specificity

- Thrombin targets are restricted due to specific interactions between the protein substrate and residues outside the catalytic cleft termed **Exosite**
- Extended interactions at exosites drive substrate affinity and contribute to substrate specificity.

**Clot
stabilization**



**Activation
of the
anticoagulant
pathway**

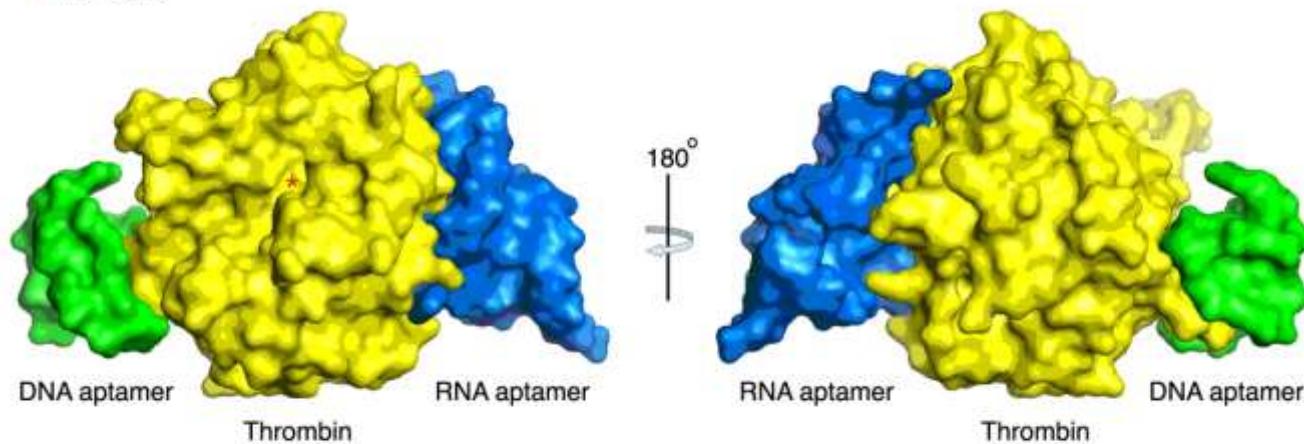


**Platelet
activation**

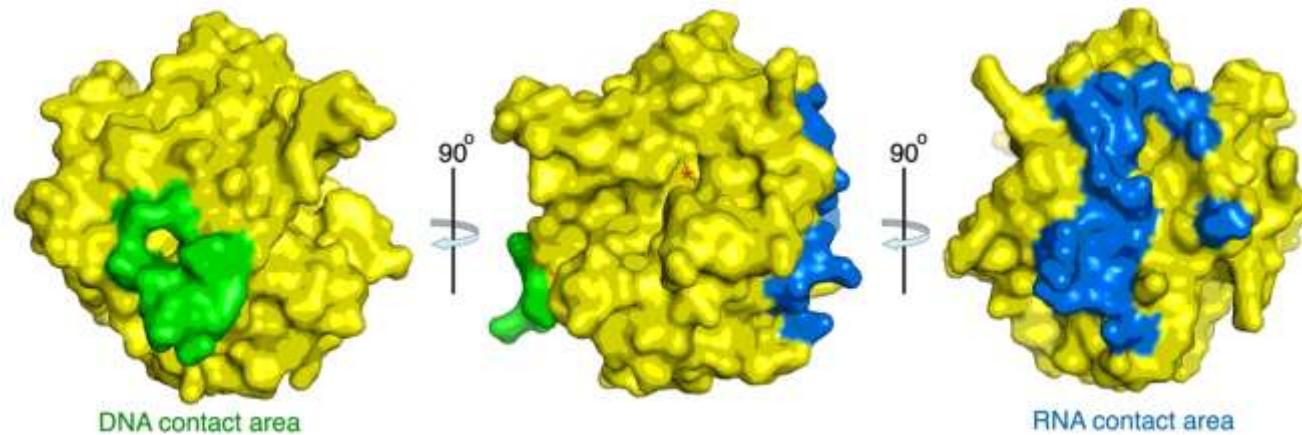
Inhibition

Exosites are good targets to inhibit specific functions of thrombin

A *Active site



B

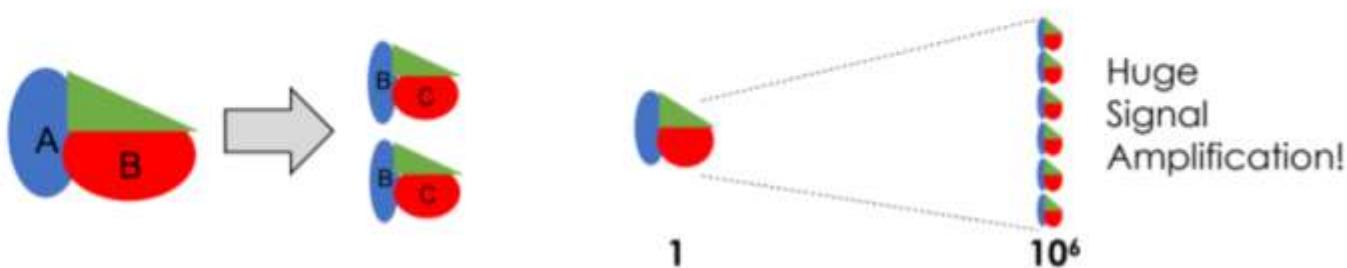


RNA

ATTIVAZIONE della Trombina

The cascade organization

Consequential enzymatic conversions of zymogens to activated enzymes



It takes place on **macromolecular complex**:

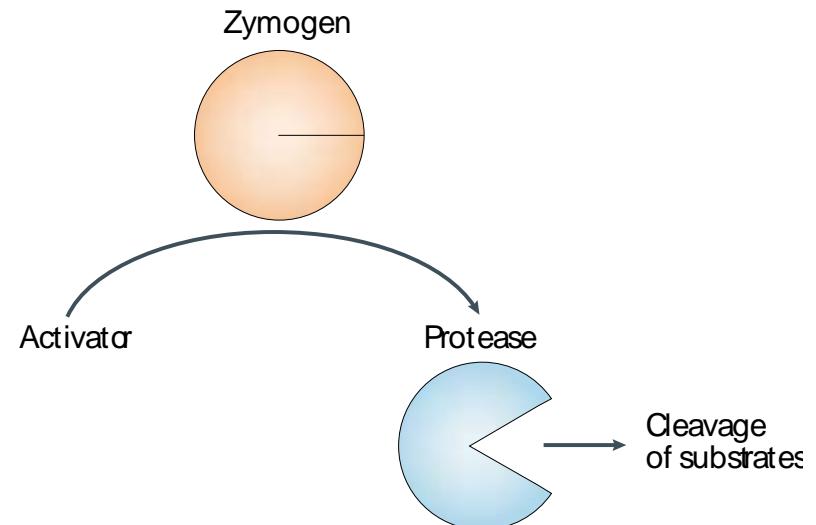
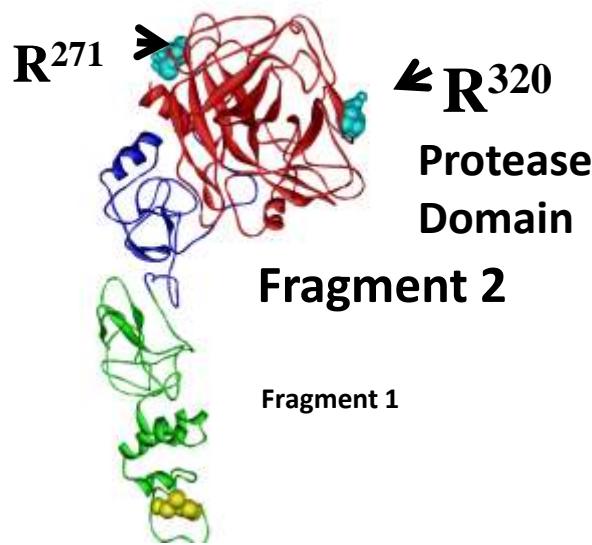
Complex name	Enzyme (active)	Cofactor	Substrate (zymogen)	Catalytic Efficiency	
Prothrombinase	FXa	FVa	Prothrombin	$>3 \times 10^5$	<p>The diagram shows a macromolecular complex composed of several proteins (purple, green, yellow) embedded in a lipid bilayer (represented by orange circles). Red arrows point from the complex to the text below, which lists its properties: Increased catalytic efficiency, Products Channeling, and Localization Control.</p> <p>Increased catalytic efficiency Products Channeling Localization Control</p>

Thrombin is synthesized as a Zymogen: Prothrombin

Zymogen:

A proenzyme or inactive enzyme. It requires a biochemical change to reveal the active site for it to become an active enzyme.

Zymogens lack the structural attributes required for formation of the enzyme-substrate complex.



Prothrombin is activated to thrombin by two proteolytic cleavages

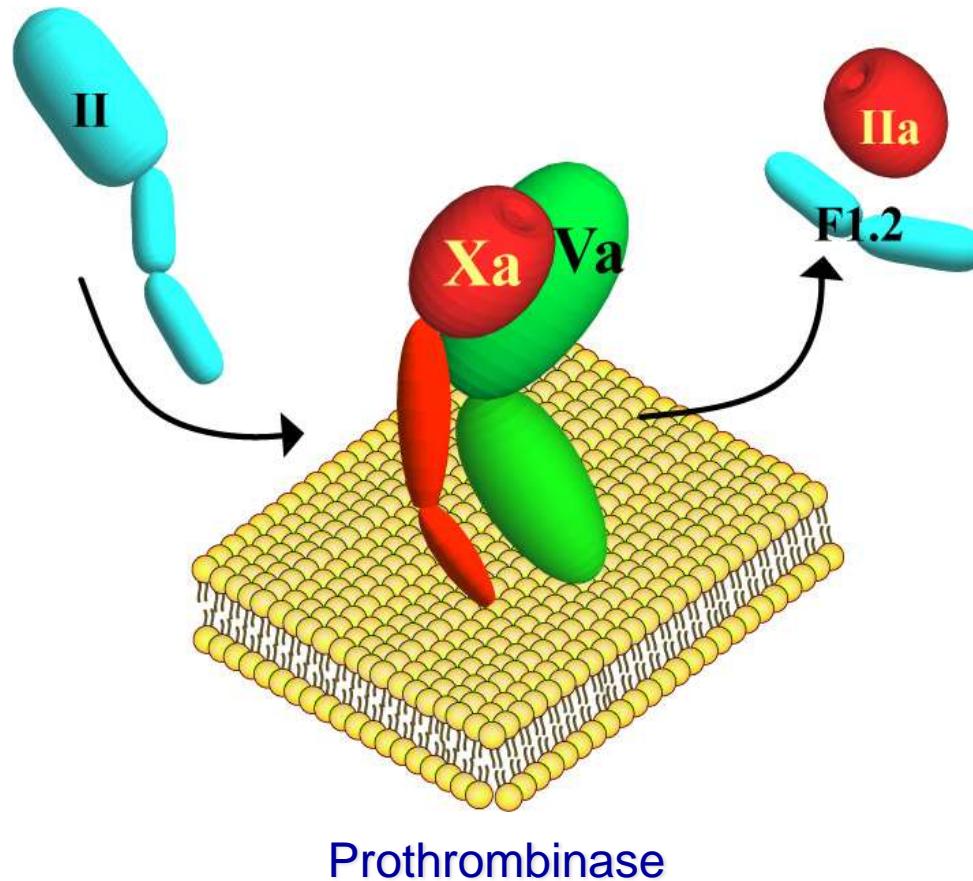


Table 1 Sites of cleavage in the human vitamin K-dependent zymogens*

Enzyme	Substrate†	P ₄	P ₃	P ₂	P ₁	↓	P _{1'}	P _{2'}	P _{3'}	P _{4'}
Xa/Va	II	I	E	G	R		T	A	T	S
	II _(15–16)	I	D	G	R		I	V	E	G
VIIa/TF, IXa/VIIIa	X _(15–16)	N	L	T	R		I	V	G	G
VIIa/TF, Xla	IX	K	L	T	R		A	E	A	V
	IX _(15–16)	D	F	T	R		V	V	G	G
VIIa/TF, Xa	VII _(15–16)	P	Q	G	R		I	V	G	G
IIa/TM	PC _(15–16)	V	D	P	R		L	I	D	G

*Sequences flanking cleavage sites relevant to the activation of the vitamin K-dependent zymogens are presented along with the relevant enzymes that catalyze these reactions. The site of bond cleavage is denoted by the arrow. †The site, in each substrate, at which cleavage is required to produce the serine proteinase is indicated as (15–16) corresponding to the homologous residue numbers in chymotrypsinogen [70].

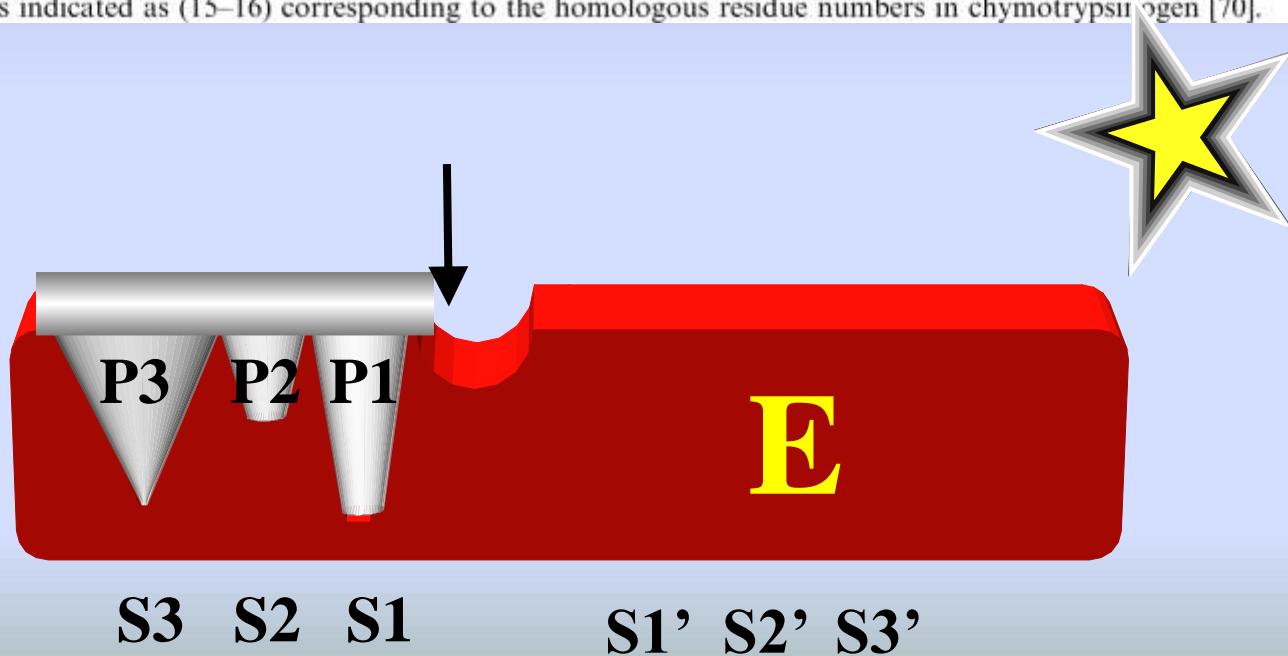
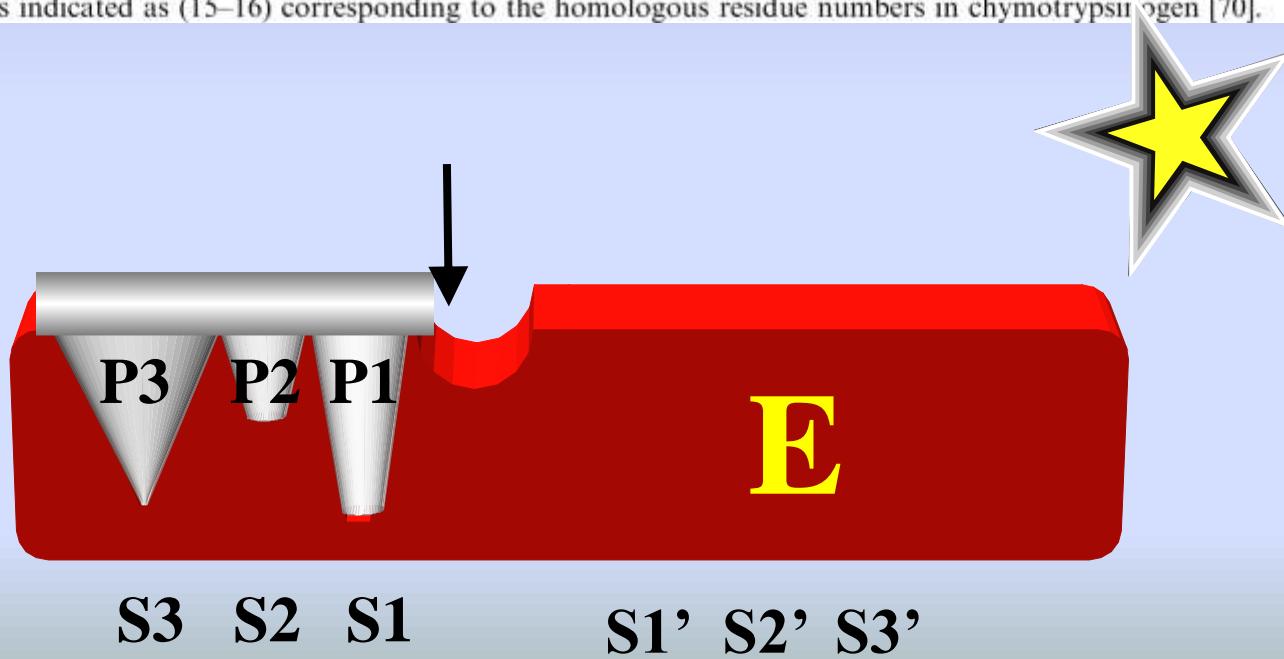


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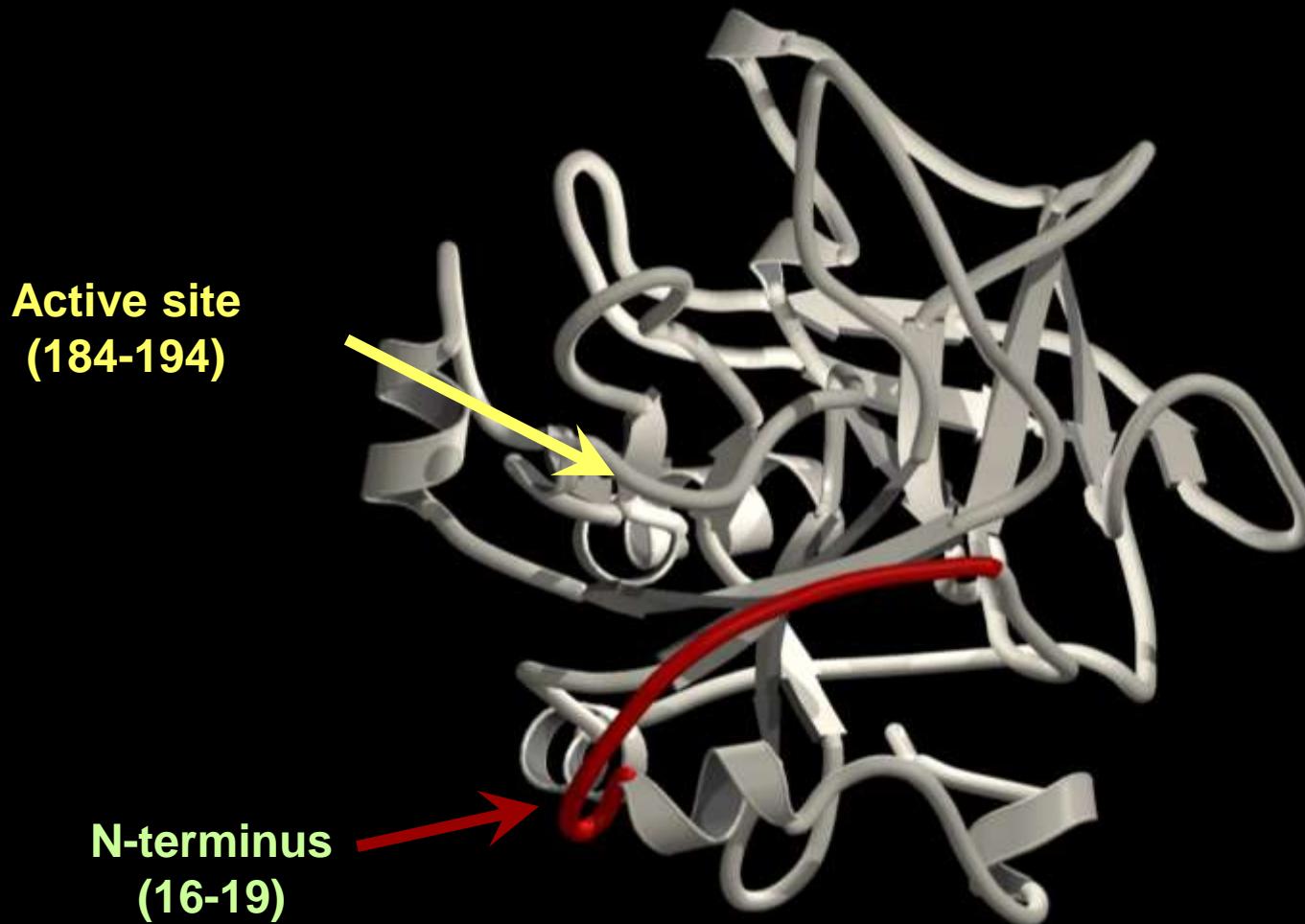
Enzyme	Substrate†	P ₄	P ₃	P ₂	P ₁	↓	P _{1'}	P _{2'}	P _{3'}	P _{4'}
Xa/Va	II	I	E	G	R		T	A	T	S
	II _(15–16)	I	D	G	R		I	V	E	G
VIIa/TF, IXa/VIIIa	X _(15–16)	N	L	T	R		I	V	G	G
VIIa/TF, Xla	IX	K	L	T	R		A	E	A	V
	IX _(15–16)	D	F	T	R		V	V	G	G
VIIa/TF, Xa	VII _(15–16)	P	Q	G	R		I	V	G	G
IIa/TM	PC _(15–16)	V	D	P	R		L	I	D	G

*Sequences flanking cleavage sites relevant to the activation of the vitamin K-dependent zymogens are presented along with the relevant enzymes that catalyze these reactions. The site of bond cleavage is denoted by the arrow. †The site, in each substrate, at which cleavage is required to produce the serine proteinase is indicated as (15–16) corresponding to the homologous residue numbers in chymotrypsinogen [70].



Serine Proteases: Conversion Pathway

- Cleavage between Arg¹⁵-Ile¹⁶ → Exposure of new N-terminus
- New N-terminus (IleVal) forms salt bridge with Asp¹⁹⁴
- N-terminal insertion leads to a conformational change in the “activation domain”



Courtesy of W. Bode,
Max Planck
Institute of Biochemistry

Prossima
MECCANISMO ATTIVAZIONE BATTERICA
Protrombina - Trombina