

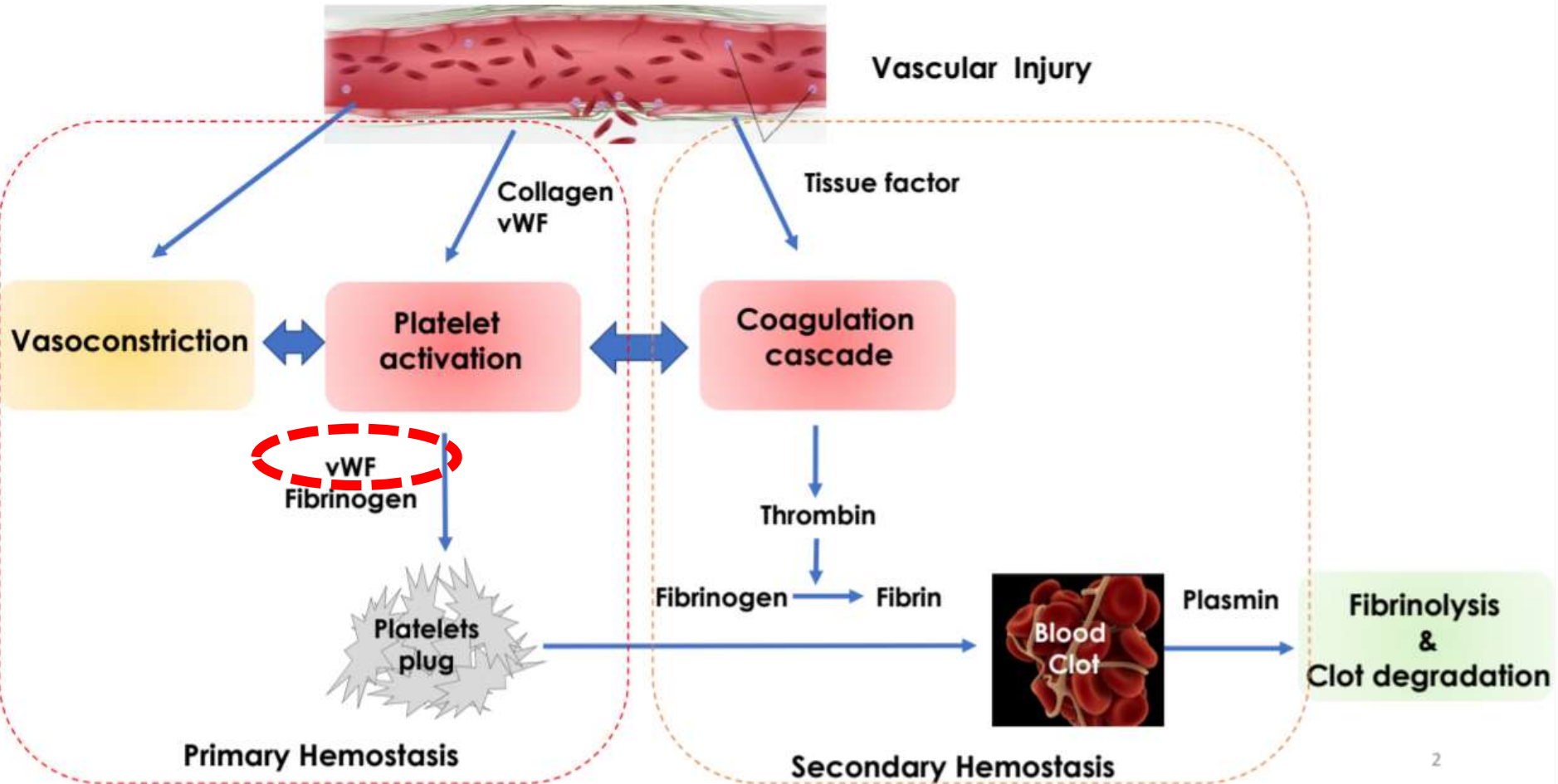
Corso di laurea in Scienze Biologiche
Corso di laurea magistrale in Scienze Biomolecolari e dell'Evoluzione

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Hemostatic process: an overview

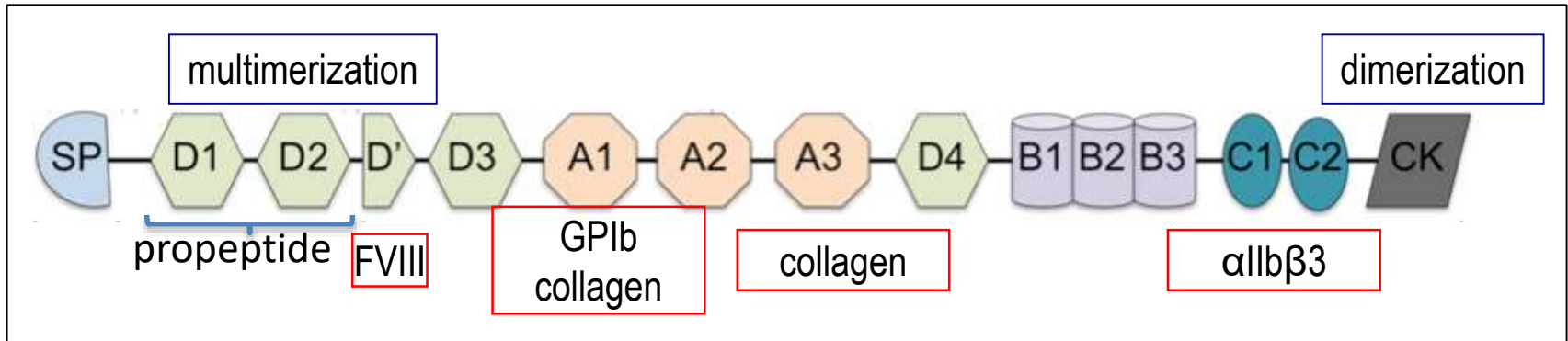


Von Willebrand factor (VWF)

- **VWF is a Large Plasmatic & Multimeric protein**
- **Produced by Endothelial cells and MEG**
- **Highly regulated biosynthesis**
- **Constitutive secretion & storage**

Von Willebrand factor (VWF)

Monomer: domain arrangement



Role of VWF in haemostasis:

- **Primary haemostasis**
- **Chaperone for coagulation FVIII**

Coagulation Cascade

It takes place on **macromolecular complex**:

Complex name	Enzyme (active)	Cofactor	Substrate (zymogen)	Catalytic Efficiency
Extrinsic Tenase	FVIIa	TF	FX	$>15 \times 10^6$
Intrinsic Tenase	FIXa	FVIIIa	FX	$>10^6$
Prothrombinase	FXa	FVa	Prothrombin	$>3 \times 10^5$

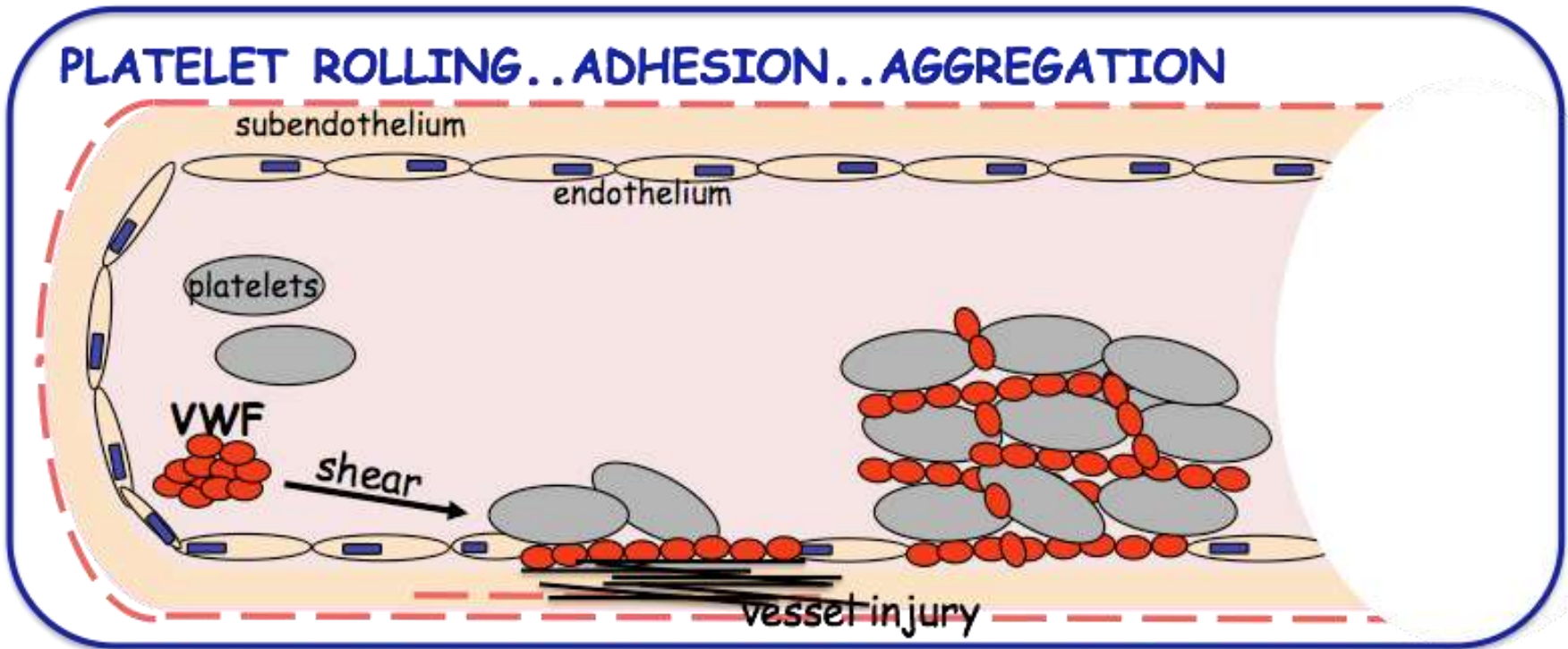
Chaperone for coagulation FVIII

VWF

- Improves stability of FVIII protein structure
- Protects FVIII from proteolysis by phospholipid-dependent proteases (activated Protein C)
- Prevents premature clearance by scavenger-receptors (such as LRP1)

Patients with no detectable VWF
> have a secondary deficiency of FVIII

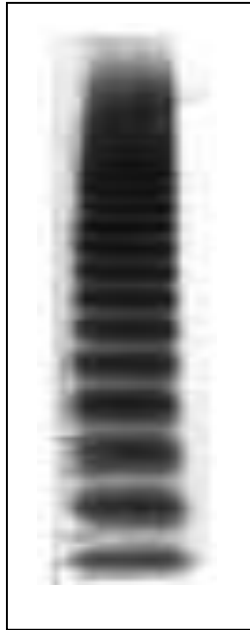
Primary haemostasis



At sites of vascular injury VWF binds to exposed collagen and allows platelets to roll and adhere to the damaged sub-endothelium.

Once platelets become activated VWF allows platelet-platelet interaction during thrombus formation.

Von Willebrand factor (VWF)



Normal plasma
sample

High-molecular-weight multimers
(HMWM)

Low-molecular-weight multimers
(LMWM)

20.000kDa



500KDa(dimers)

- HMWMs are the most important for VWF function in haemostasis

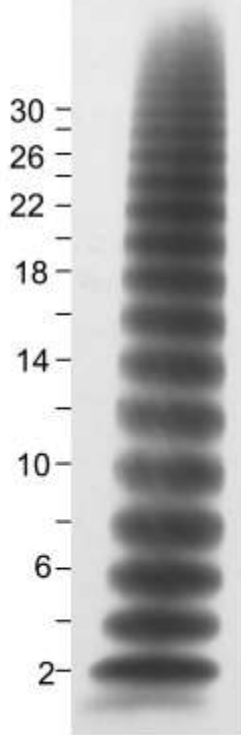
- VWF size in plasma is regulated by the enzymatic activity of a metalloprotease (ADAMTS13)

Dimerization and multimerization of VWF



Monomer
 0.25×10^6 Da

> 30 monomers !!



Dimer
 0.5×10^6 Da



Tetramer
 1.0×10^6 Da



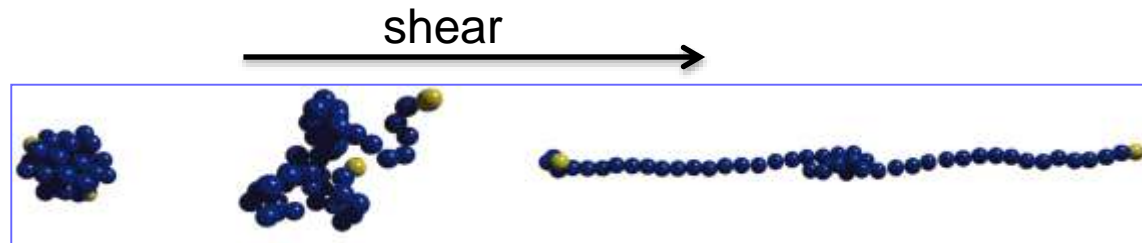
Hexamer
 1.5×10^6 Da



Multimer
 $\geq 7.5 \times 10^6$ Da

Primary haemostasis

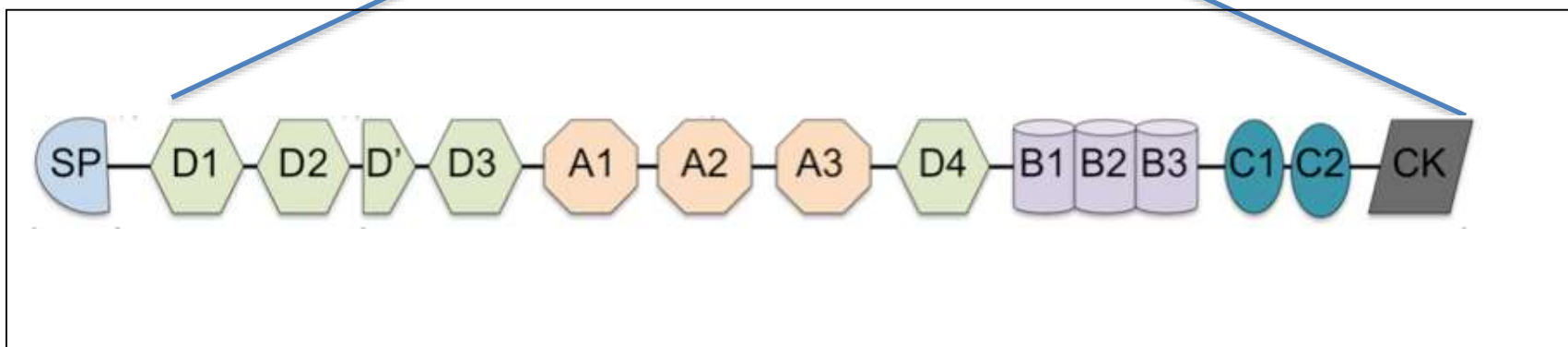
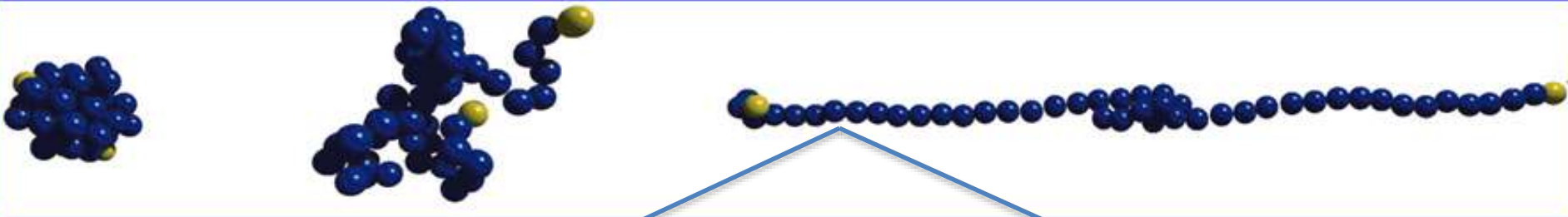
Structure-function relationship



VWF conformation depends on the applied shear force (blood flow)

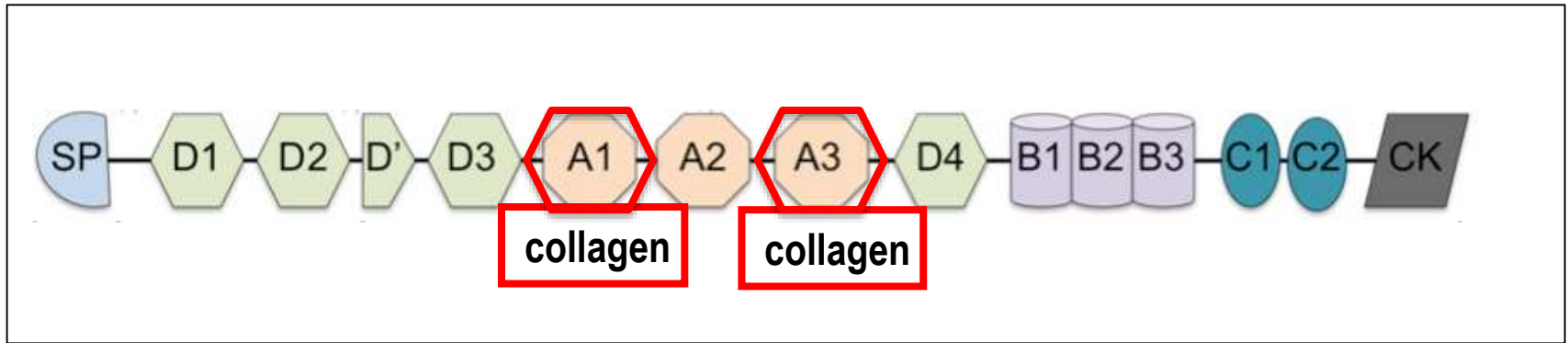
- In normal conditions VWF circulates in a globular state
- In case of high-shear rate VWF elongates into its active, platelet-binding conformation

a multimeric

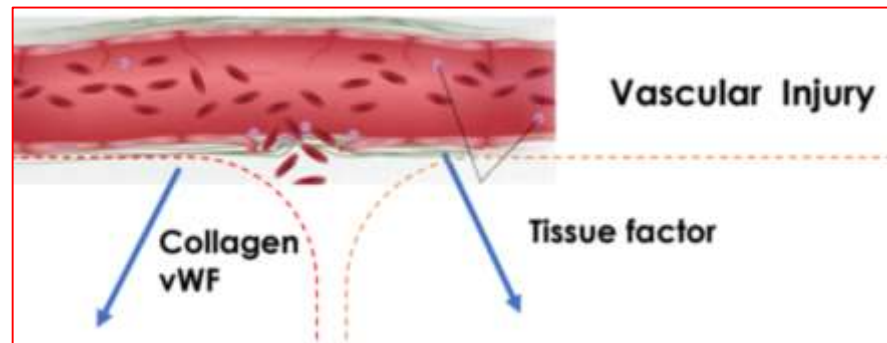


..... and multidomain protein

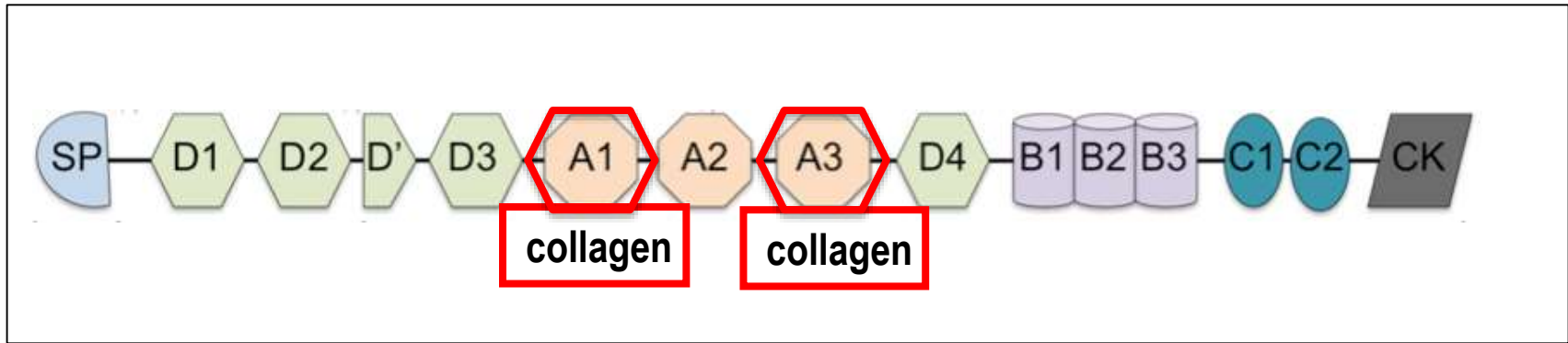
VWF-collagen binding



- Two sites of VWF-collagen binding in the A1 and A3 domains
- At sites of vascular injury sub-endothelial collagen becomes exposed to flowing blood and VWF can bind to it.

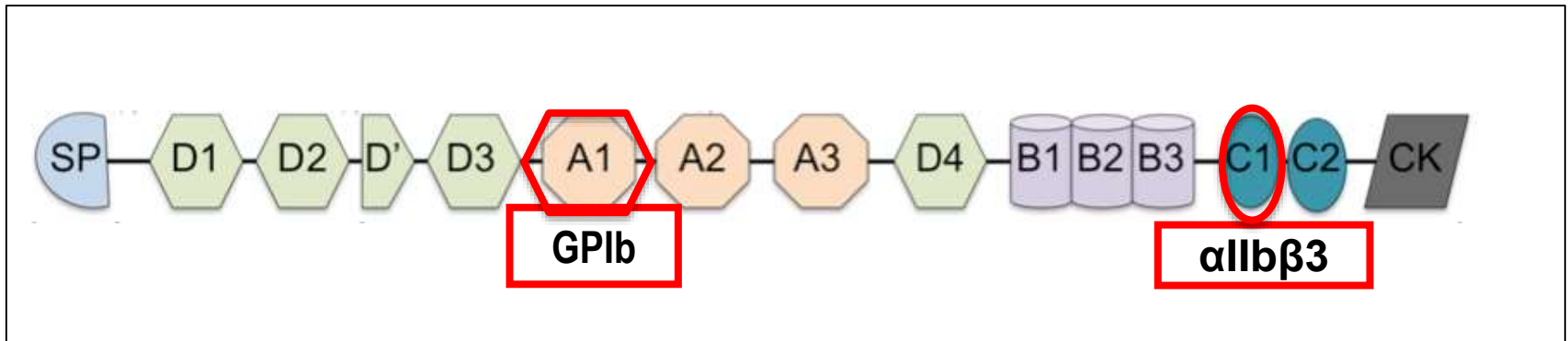


VWF-collagen binding 2

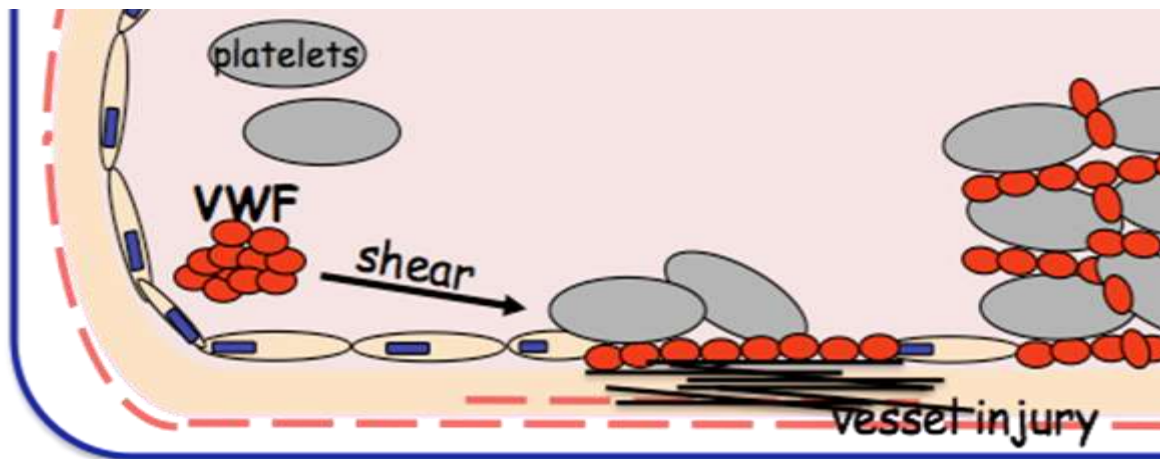


- Mutations in the A3 domain are associated with defective VWF binding to collagen type I and III and to bleeding phenotypes
- The larger VWF multimers have a much higher affinity for collagen. Thus, collagen binding is an indirect measure of multimer size and VWF activity

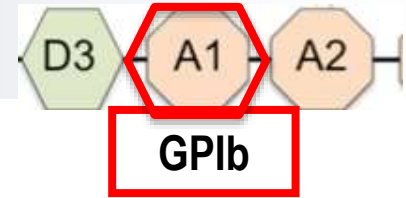
VWF-platelet binding



- Two sites of interaction between VWF and platelets:
 - VWF A1 & platelet GPIb complex
 - VWF RGDS sequence in the C1 domain & platelet αIIbβ3 integrin



VWF-platelet binding A1&GPIb

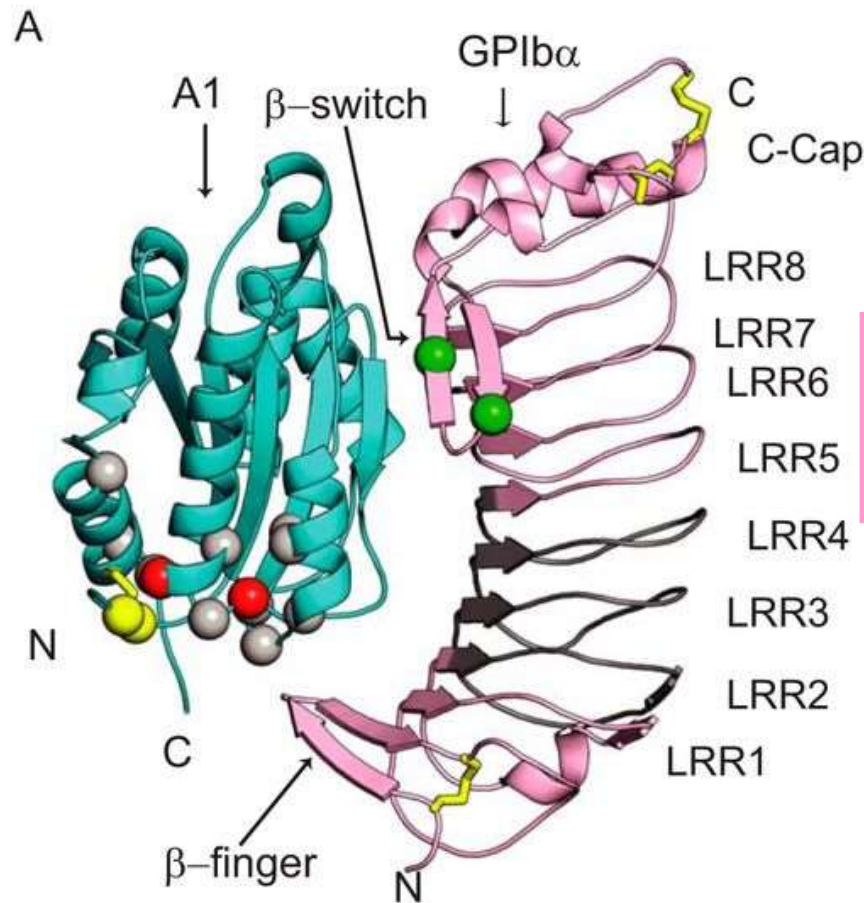


- GPIb is a complex that binds with the A1 domain of VWF
- The physiologic stimulus that induces VWF binding to platelet GPIb α is shear > meaning that VWF needs to be in its elongated conformation to be able to bind GPIb α



The VWF A1-GPIb α complex.


VWF A1 domain

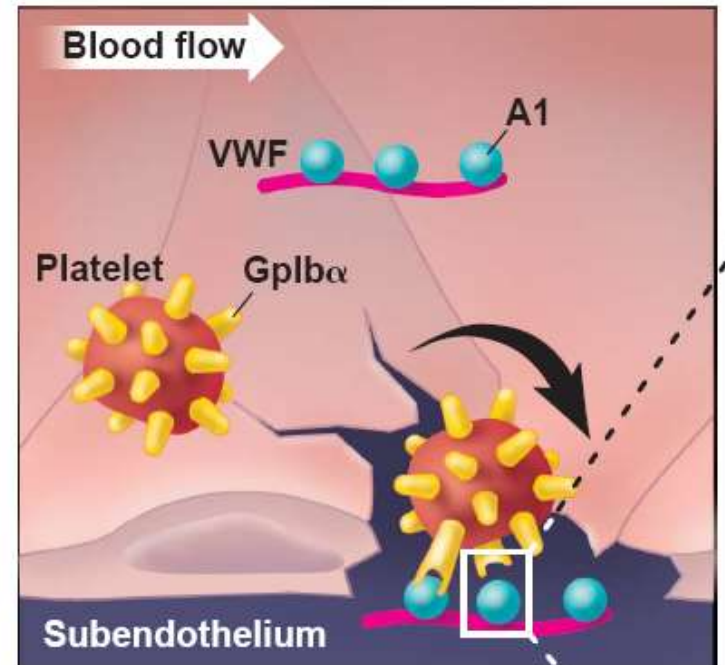


Platelet GPIb α receptor

Mark A. Blenner et al. J. Biol. Chem. 2014;289:5565-5579

VWF-platelet binding A1&GPIb

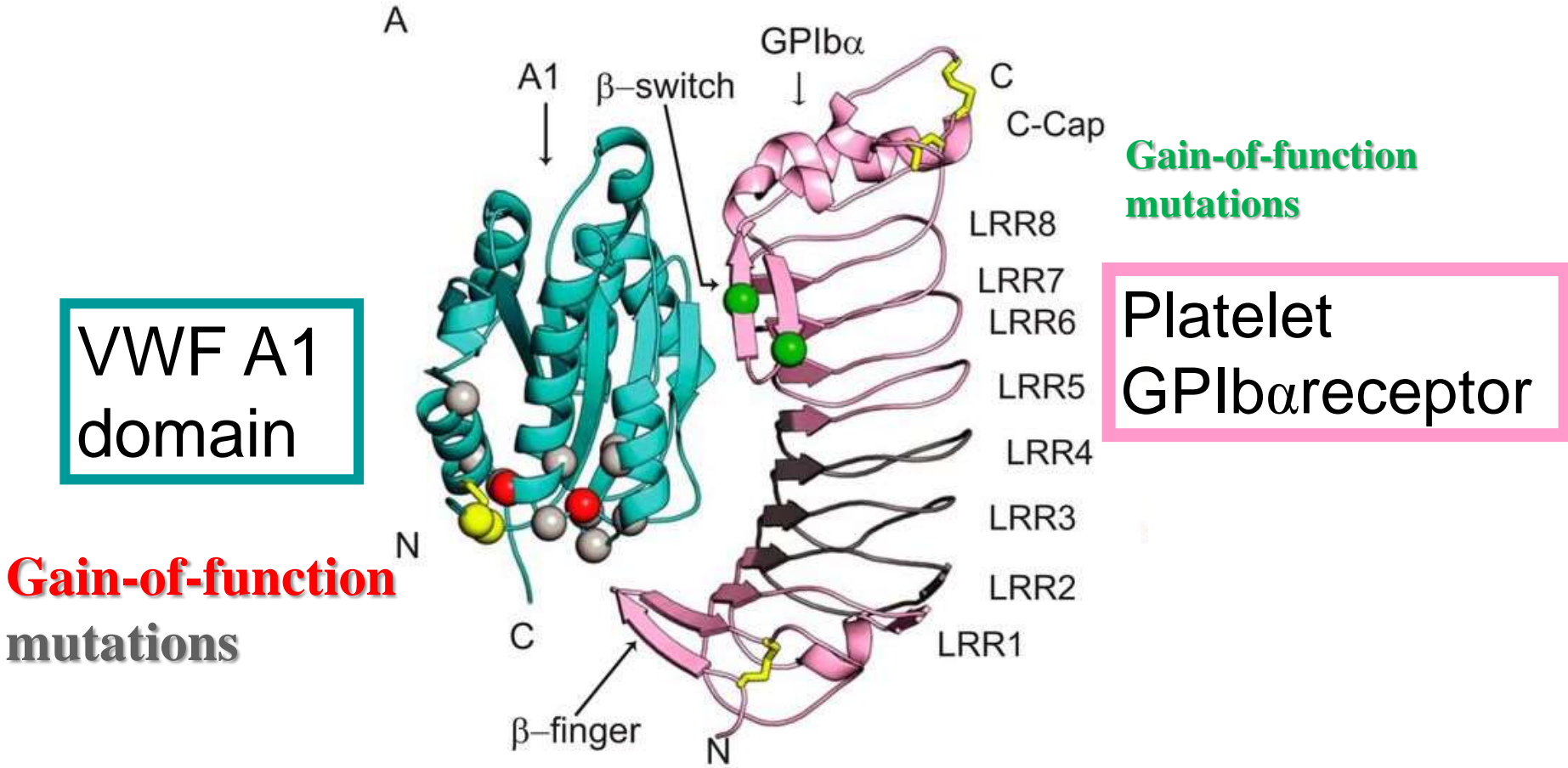
- GPIb is a complex and only GPIb α binds with the A1 domain of VWF
- The physiologic stimulus that induces VWF binding to platelet GPIb α is shear > meaning that VWF needs to be in its elongated conformation to be able to bind GPIb α 
- VWF-GPIb α binding is reversible: the high number of GPIb α molecules on platelet surface and the high concentration of A1 domains within VWF multimers allow **platelet rolling**



VWF-platelet binding A1&GPIb and Von Willebrand Disease

- There are different types of mutations that affects VWF-GPI α interaction and cause bleeding
- Some of the these mutations turn VWF/GPIb α **in the active conformation** > resulting in constitutive (instead of shear-dependent) VWF-platelet interaction
 - *Gain-of-function* mutations in the A1 domain of VWF
 - *Gain-of-function* mutations in GPIb α

The VWF A1-GPIb α complex.



Mark A. Blenner et al. J. Biol. Chem. 2014;289:5565-5579

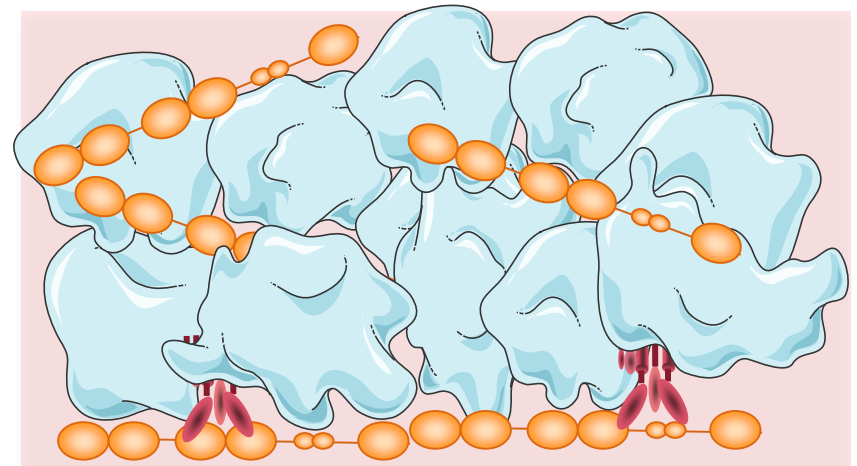
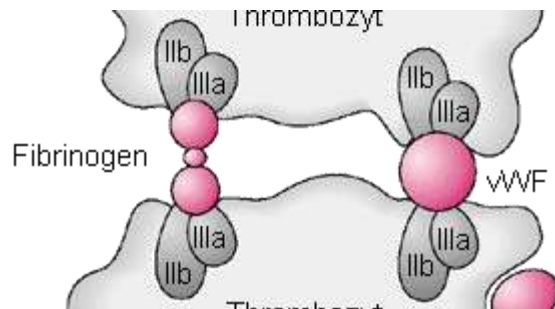


VWF-platelet binding RGDS(C1)& α IIb β 3

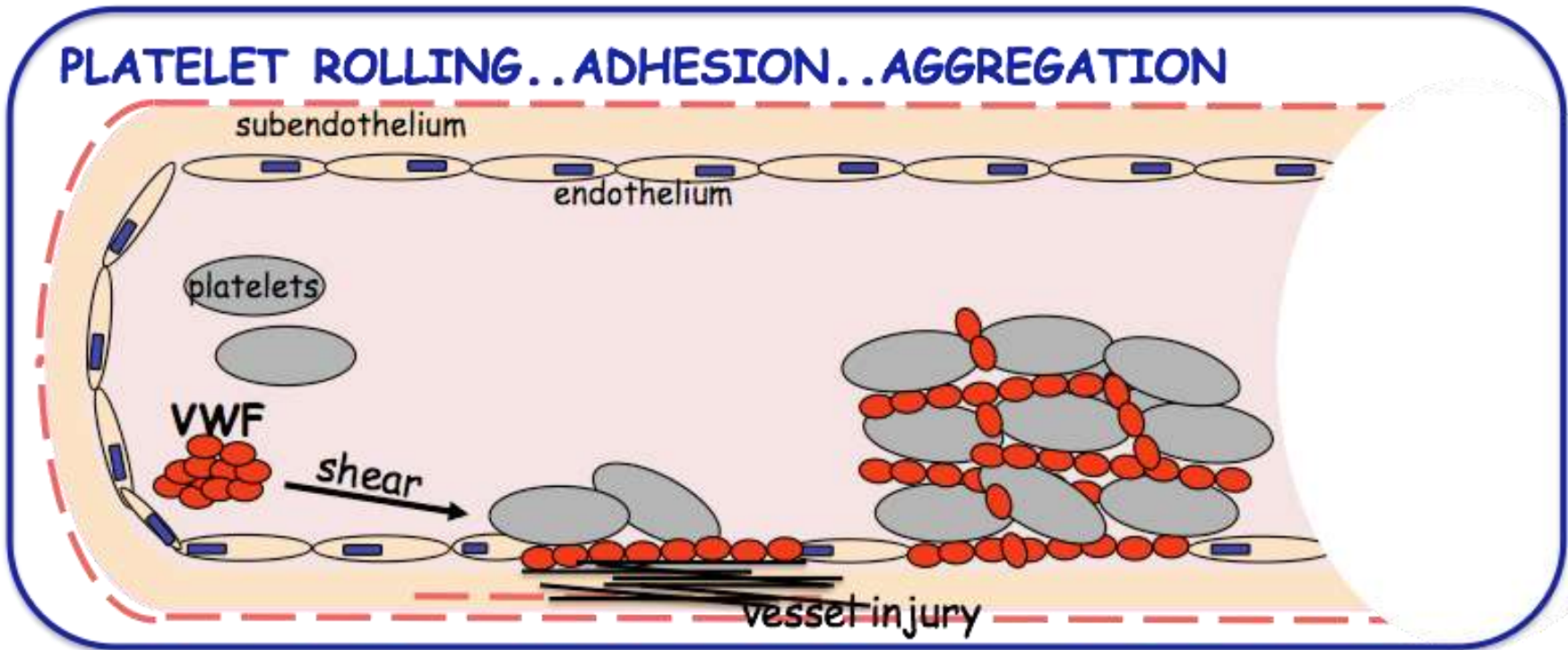


α IIb β 3

- Integrin α IIb β 3 becomes available for VWF binding only after platelet activation and a consequent conformational change
- VWF needs to be in its active conformation to bind α IIb β 3
- α IIb β 3-VWF interaction is **irreversible** and allows stable platelet-platelet interaction during thrombus formation



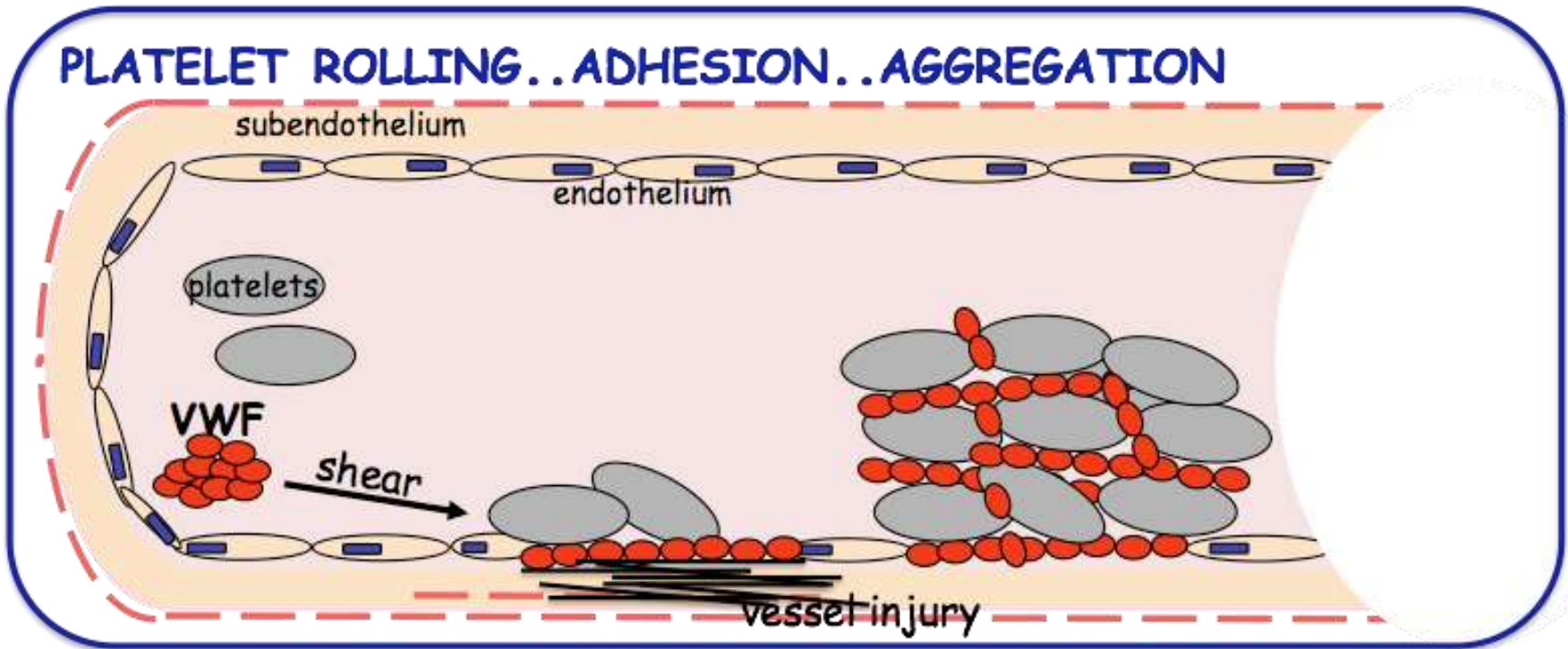
Role of VWF in primary haemostasis



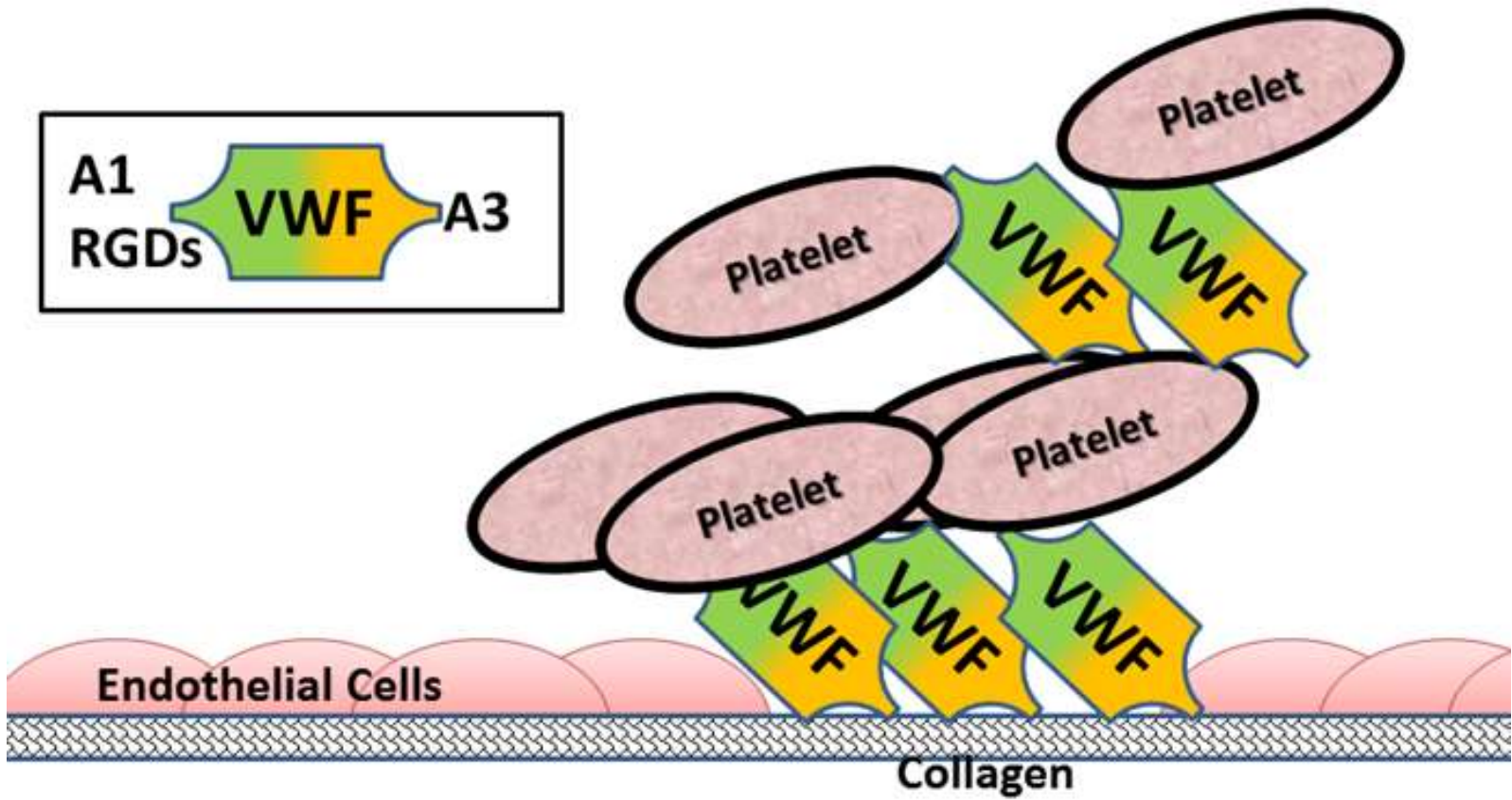
1 Vascular injury:

- exposed sub-endothelial matrix component (such as collagen)
 - High shear stress elongates VWF in its active conformation
- VWF binds to collagen & to platelet GPIIb/IIIa (> platelet rolling)

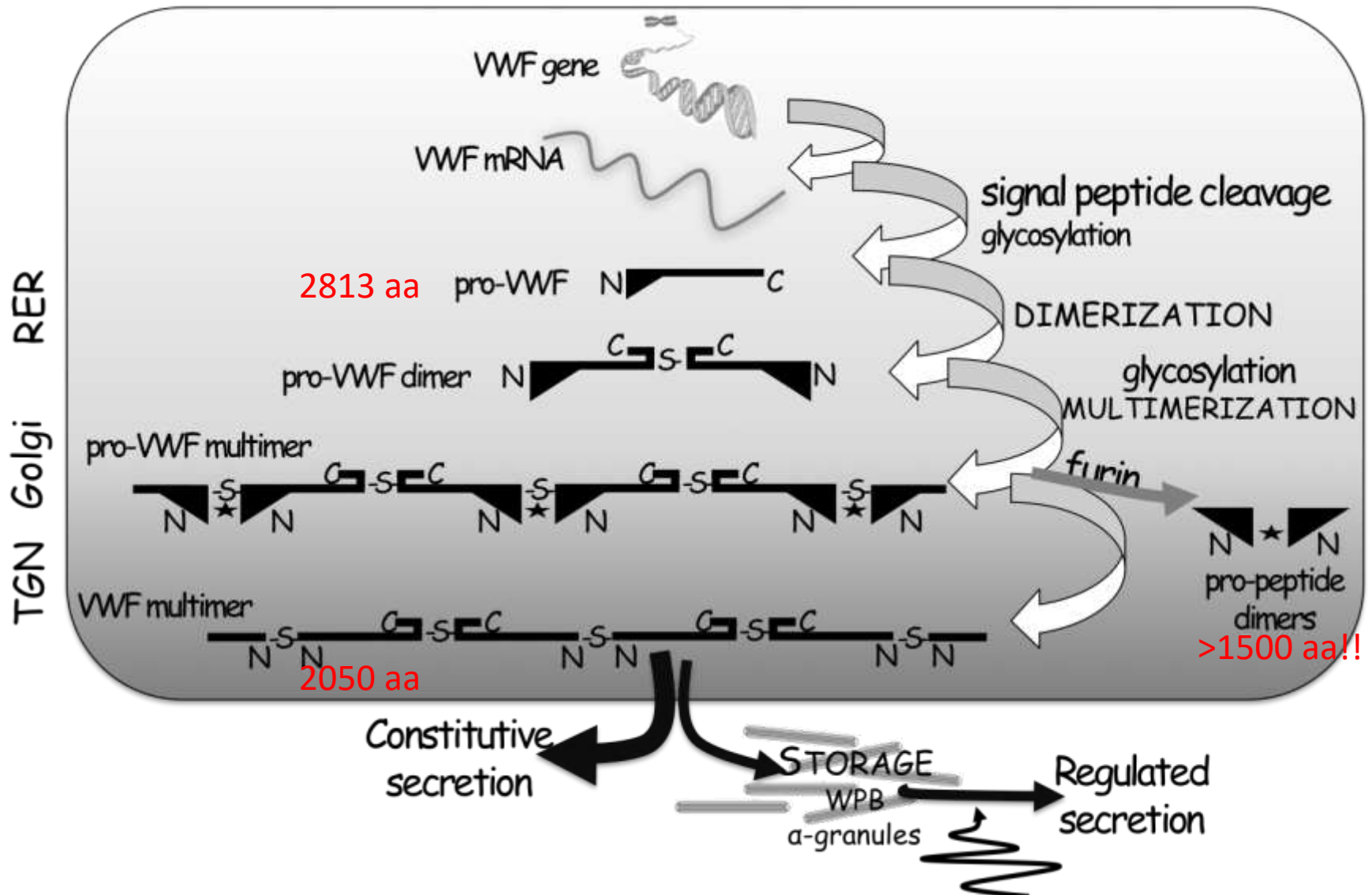
Role of VWF in primary haemostasis



2 platelet interaction with VWF and subendothelium allows their adhesion and activation > conformational change in $\alpha\text{IIb}\beta\text{3}$ integrin > interaction with VWF RGDS & fibrinogen > platelet-platelet interaction > thrombus



Von Willebrand factor (VWF) –biosynthesis-

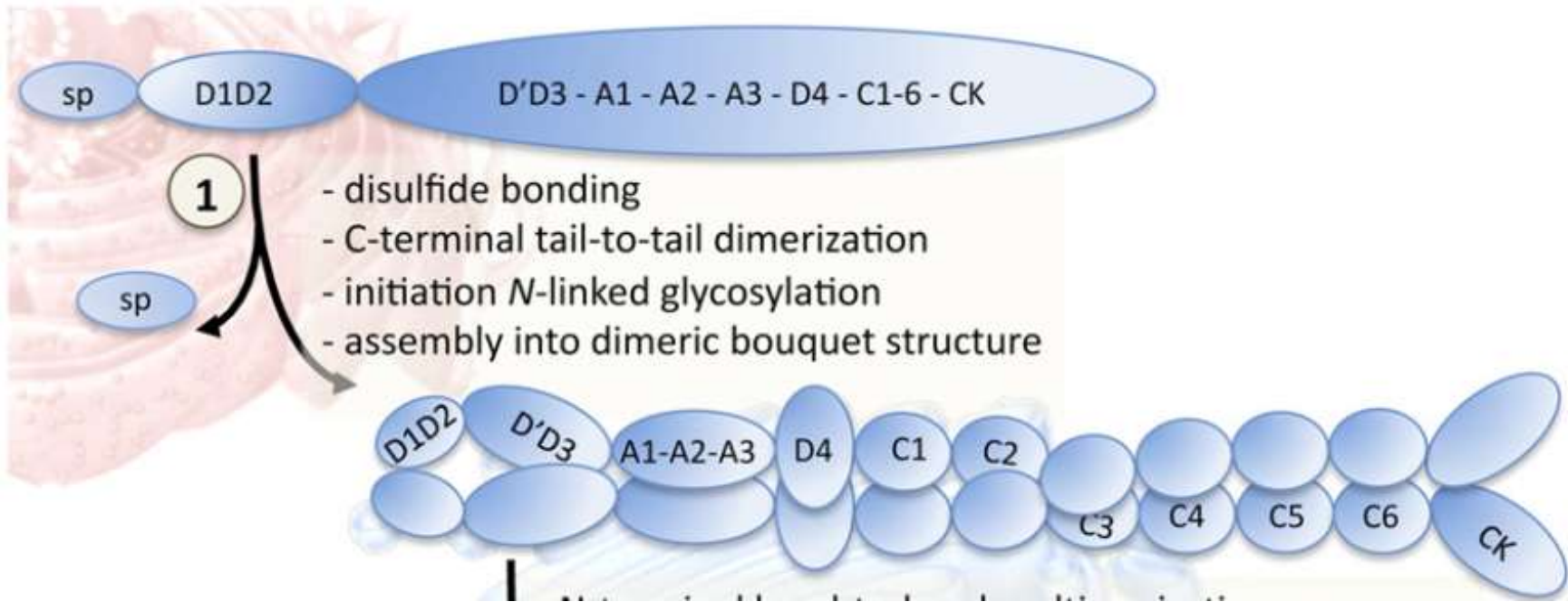


Von Willebrand factor (VWF) –storage-

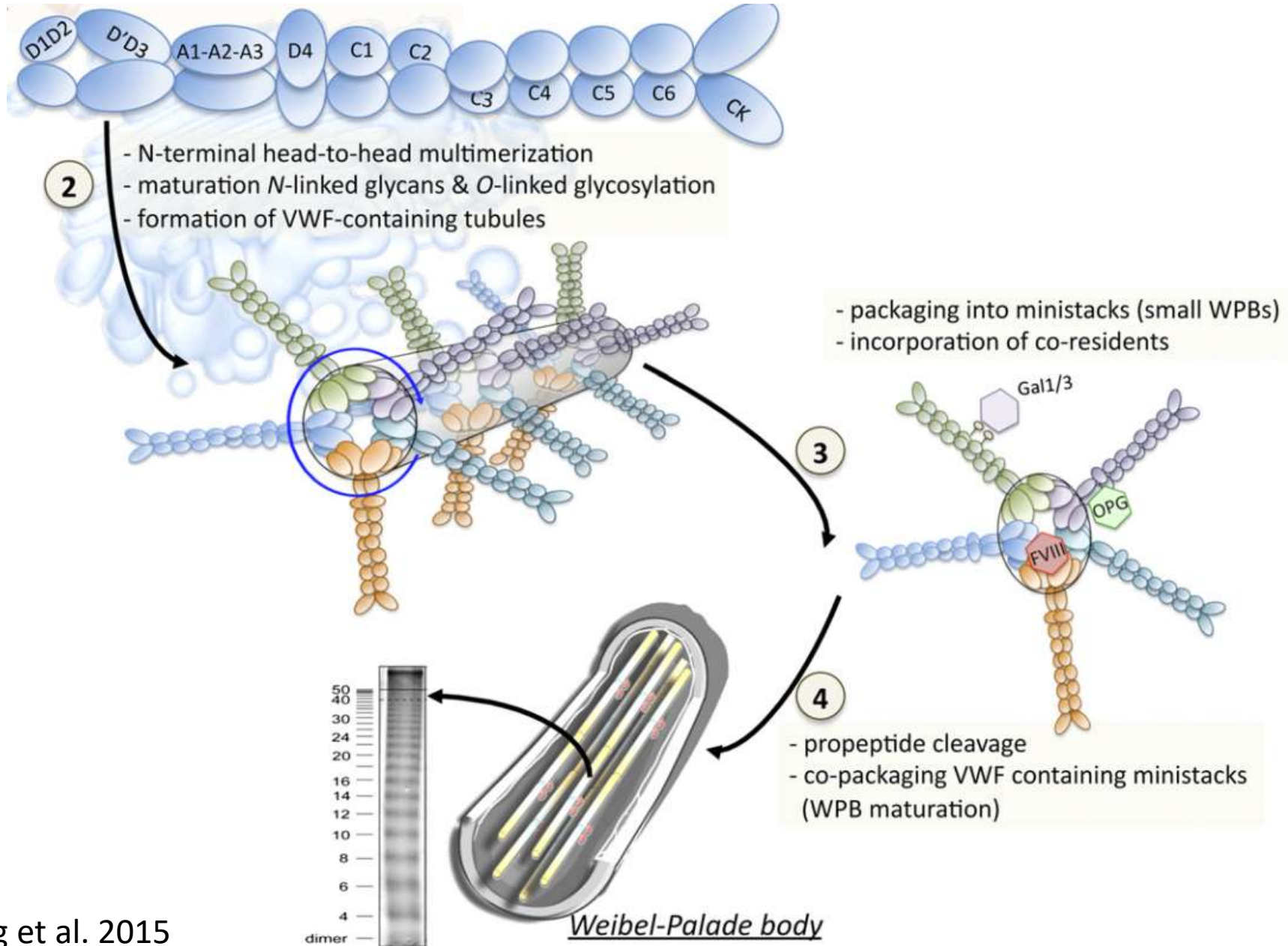
Weibel-Palade bodies

- Exclusively present in endothelial cells
- Highly structured
- Storage of large VWF multimers
- Regulated secretion following cellular stimulation

Von Willebrand factor (VWF) –storage-



Von Willebrand factor (VWF) –storage-



VWF-related pathologies

+

**Thrombotic
Thrombocytopenic
Purpura (TTP)**

Thrombosis

VWF



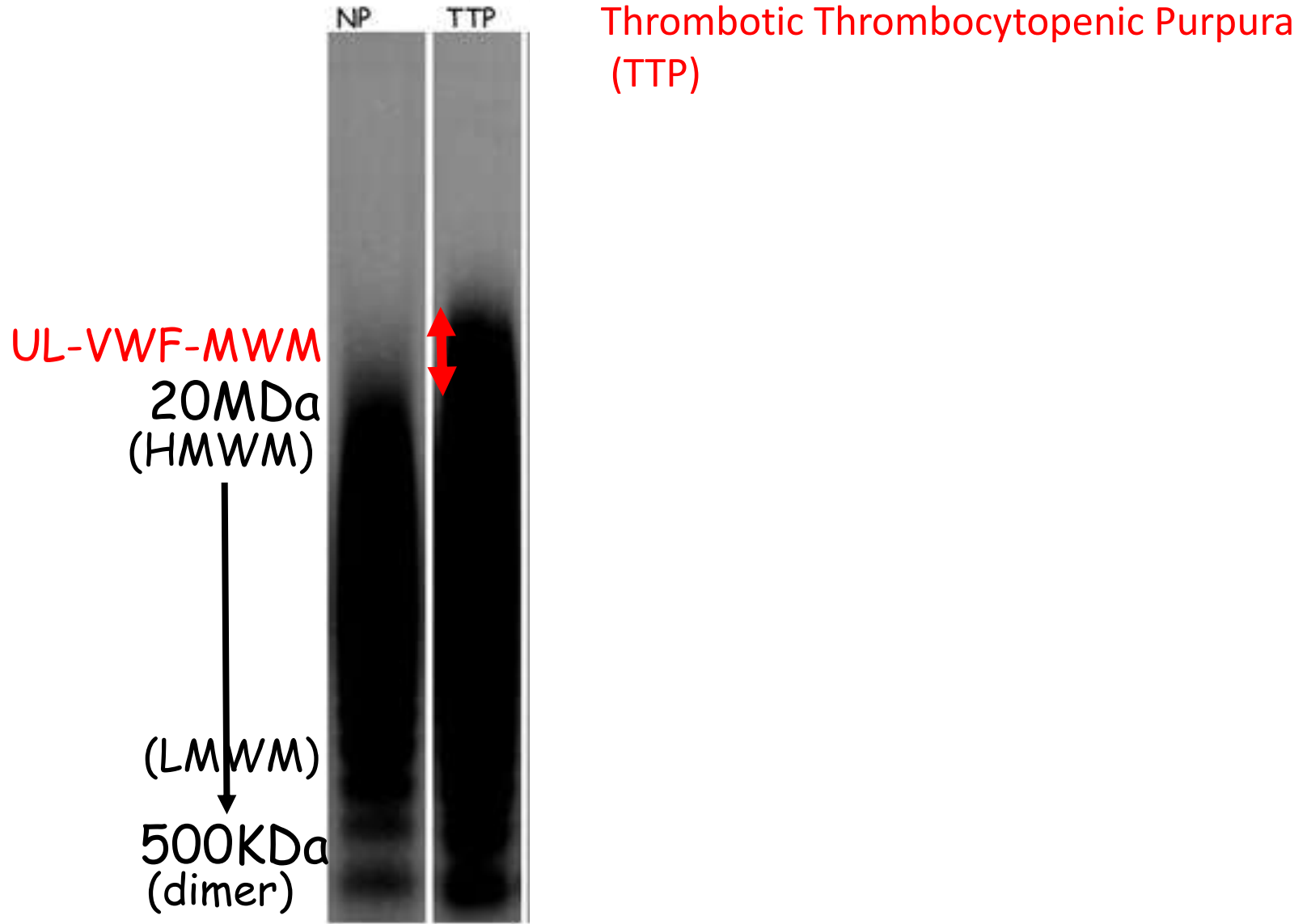
Haemostasis

-

**von Willebrand
Disease (VWD)**

Bleeding

- Presence of Ultra-large (UL)-VWF-MWMs in plasma



VWF-related pathologies

Thrombotic Thrombocytopenic Purpura (TTP)

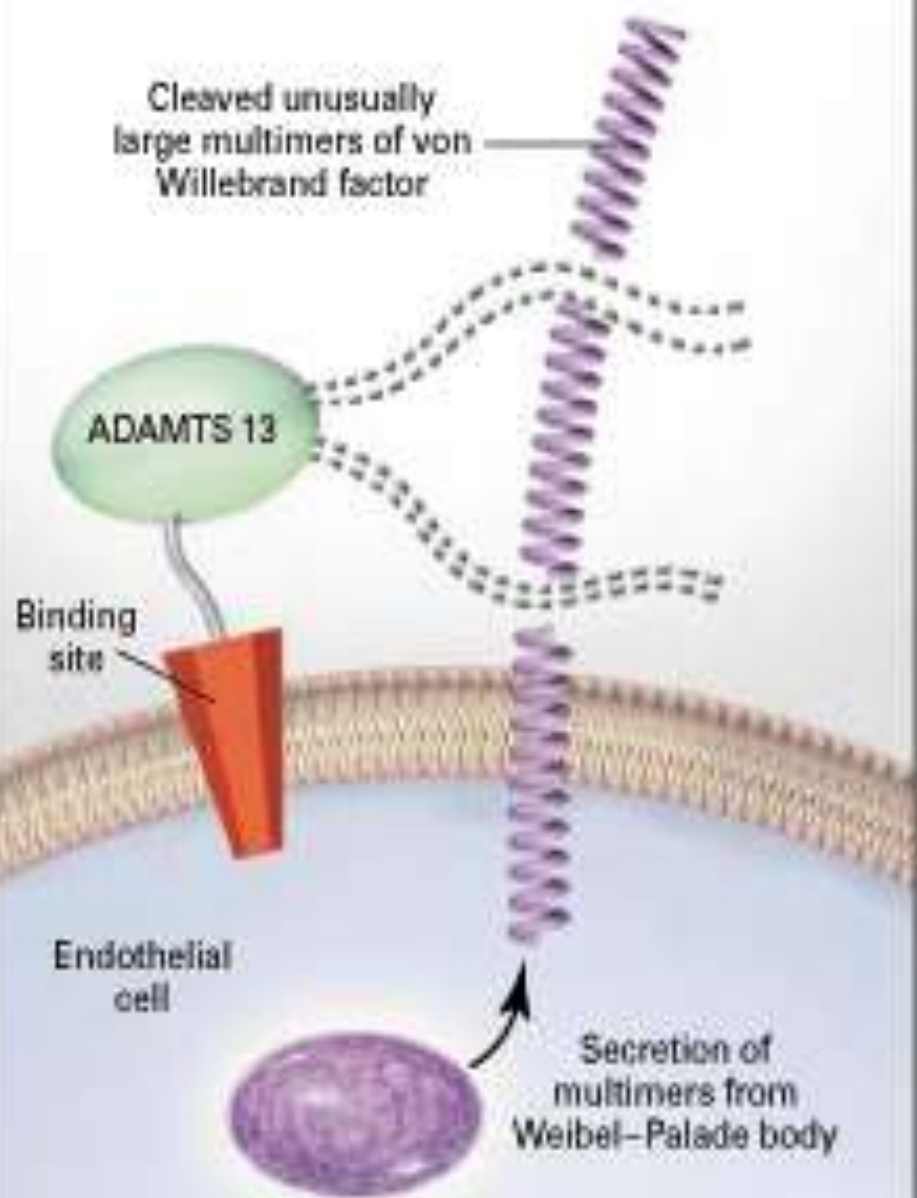
- Systemic disorder characterized by inappropriate deposition of VWF and platelet rich thrombi throughout the microvasculature, thrombocytopenia, organ failure and death
- Presence of Ultra-large (UL)-VWF-MWMs in plasma

VWF-related pathologies

Thrombotic Thrombocytopenic Purpura (TTP)

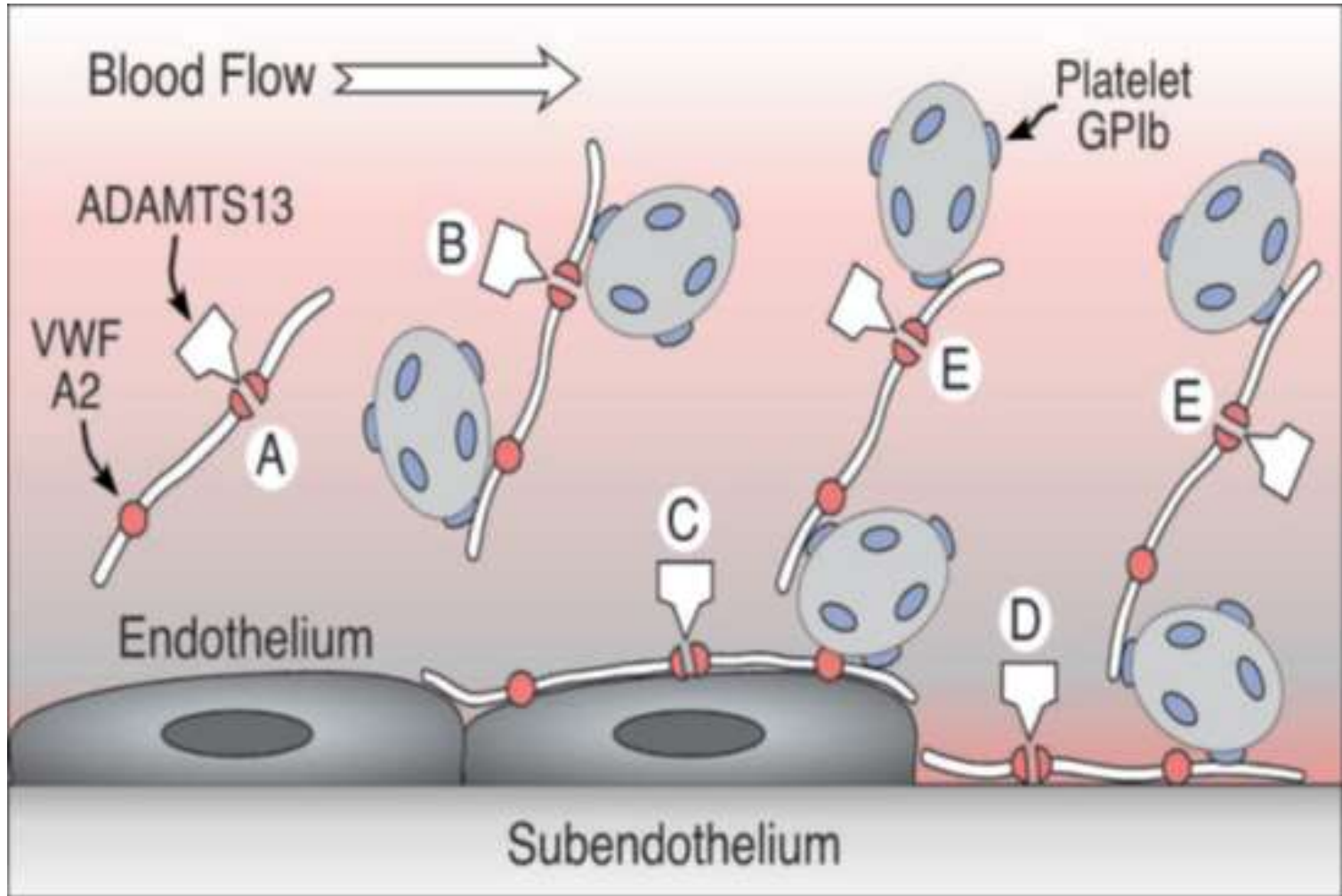
- TTP results from the deficiency of the metalloprotease ADAMTS13 that cleaves circulating VWF
- **Rare inherited TTP** = Over 50 mutations in the ADAMTS13 gene have been identified in patients with familial TTP
- **More frequent acquired TTP** > due to inhibitory anti-ADAMTS13 autoantibodies (more frequent in woman)

Normal Subject



A

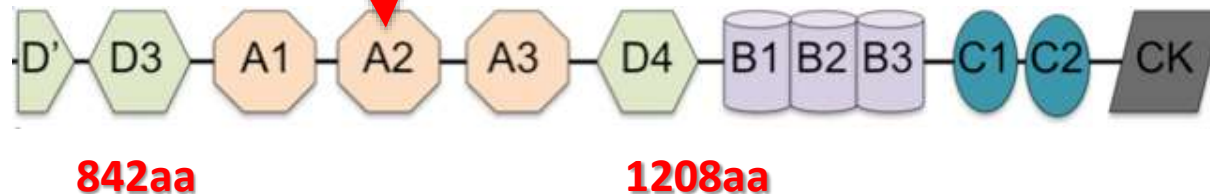
ADAMTS13



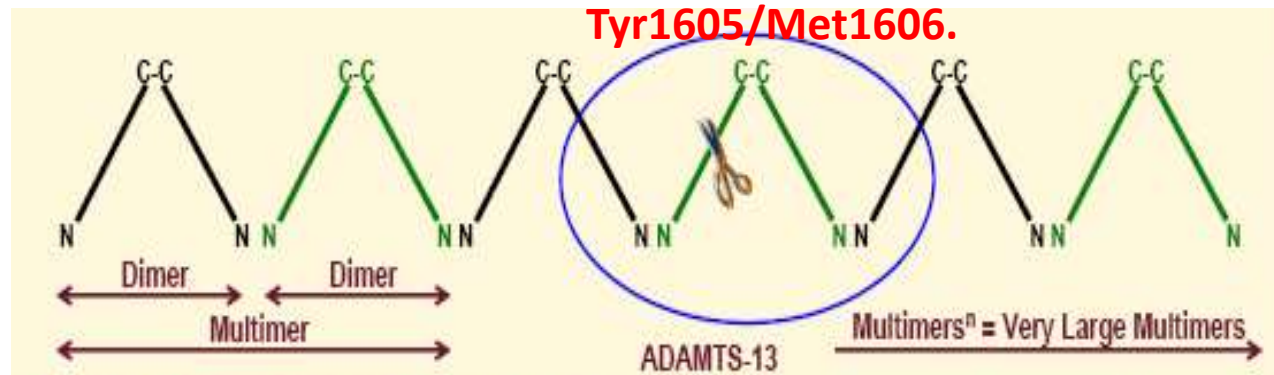
ADAMTS13

(A Disintegrin And Metalloprotease with ThromboSpondin type 1 motifs)

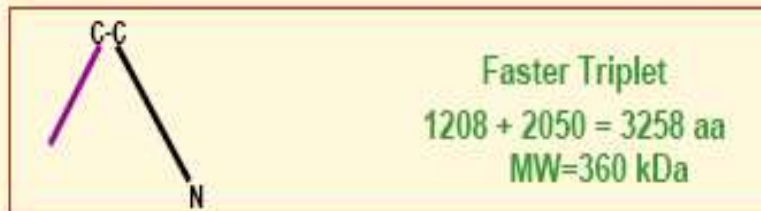
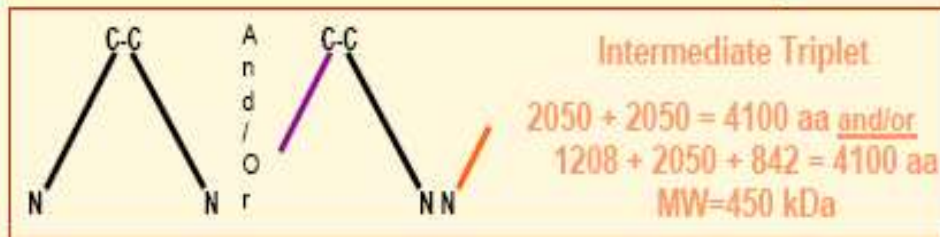
- It cleaves VWF UL-multimers **as soon as they are released**, into smaller and less thrombogenic multimers
- The only known substrate of ADAMTS13 is VWF. The **cleavage site** is in the **A2** domain between Tyr1605 & Met1606.



ADAMTS13 activity and multimer migration



VWF Proteolytic Fragments



aa = amino acids

Adapted from: Fischer BE, et al.
 Biochem J. 1998;331 (Pt 2):483-8.



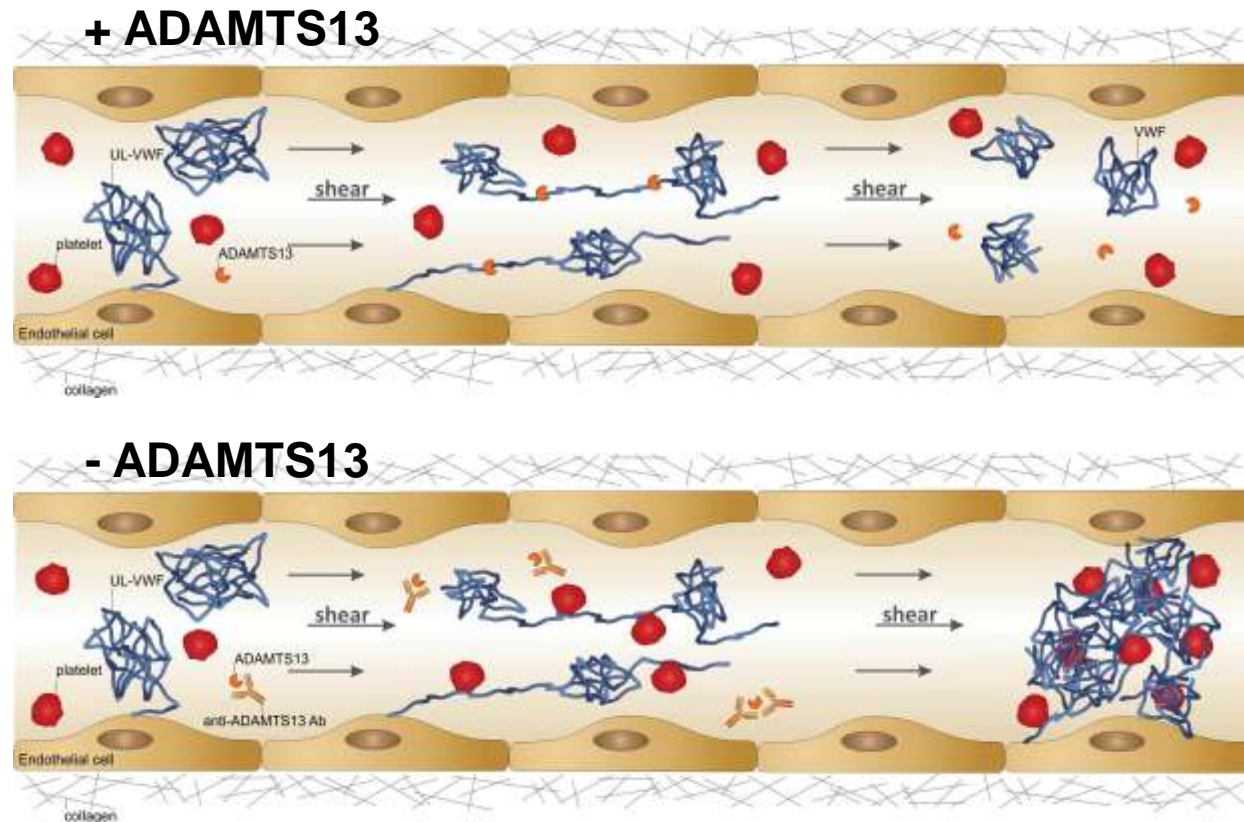
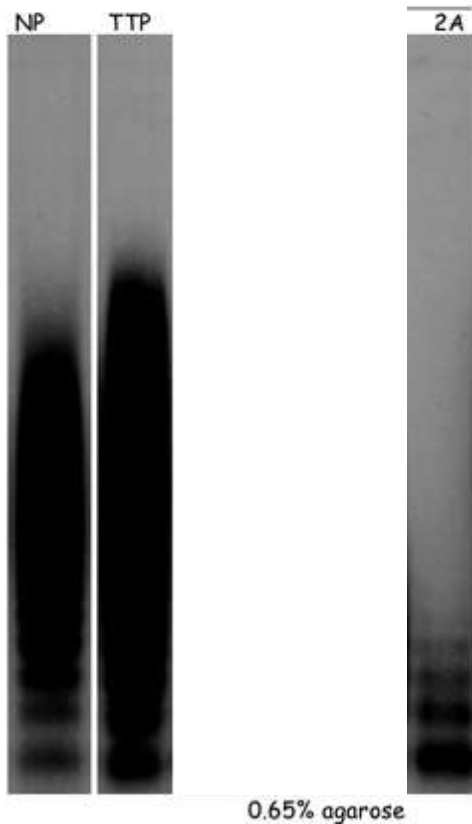
ADAMTS13

(A Disintegrin And Metalloprotease with ThromboSpondin type 1 motifs)

- Unlike most plasma proteases, ADAMTS13 is **constitutively active** in circulation
- VWF needs to be in its elongated conformation to expose the cleavage site for ADAMTS13 (shear-dependent mechanism) > **VWF auto-regulates its own cleavage**

ADAMTS13

- ADAMTS13 deficiency > TTP > UL-VWF-MWMs
- **ADAMTS13 hyper-activity (mutations in VWF that result in increased susceptibility to ADAMTS13 cleavage)** > VWD-type 2A > absence of HMWMs and increased satellite bands



Role of VWF beyond haemostasis: unexpected versatility

