

Structure Determination of a Bioengineered
Human/Porcine Factor VIII
for Hemophilia A Treatment,

and

Improvements to the Human Factor VIII Model



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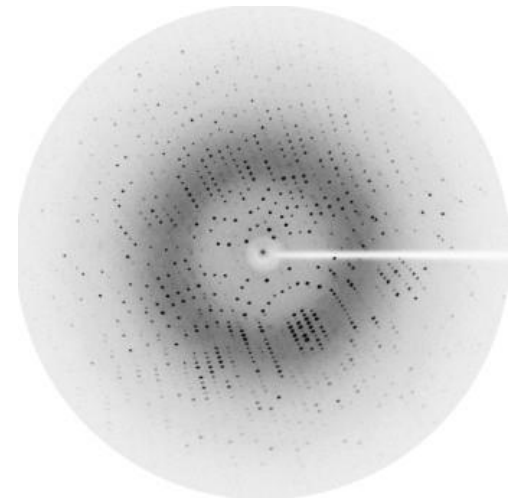
2018 GRADUATE STUDENT SYMPOSIUM



This talk will:



- Introduce:
 - The bleeding disorder Hemophilia A
 - The Blood Coagulation Cascade
 - Blood Coagulation Factor VIII (FVIII) protein
- 3-D molecular structure of novel human/porcine chimeric FVIII for Hemophilia A therapy
- Implications for future research



Central Dogma of Biology

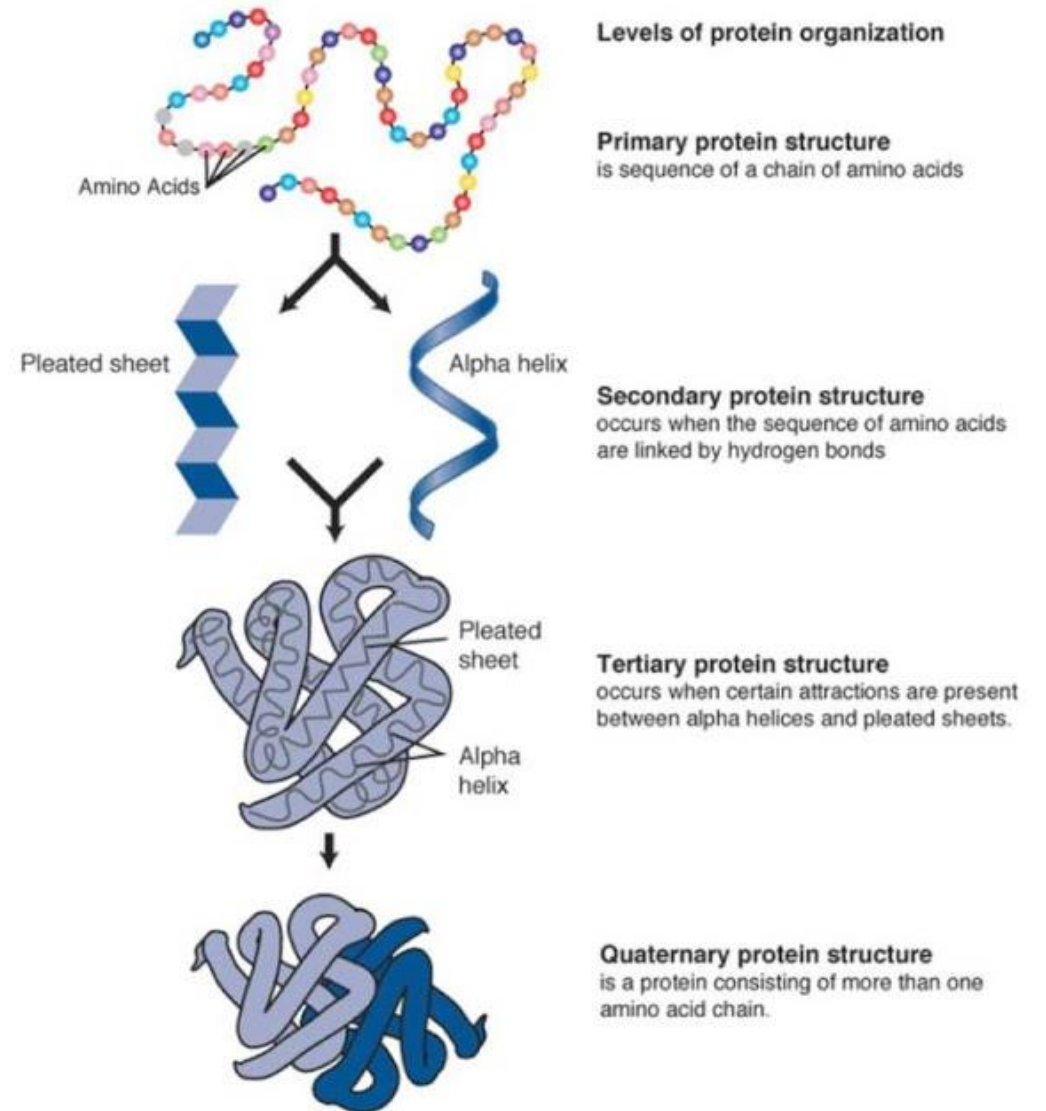
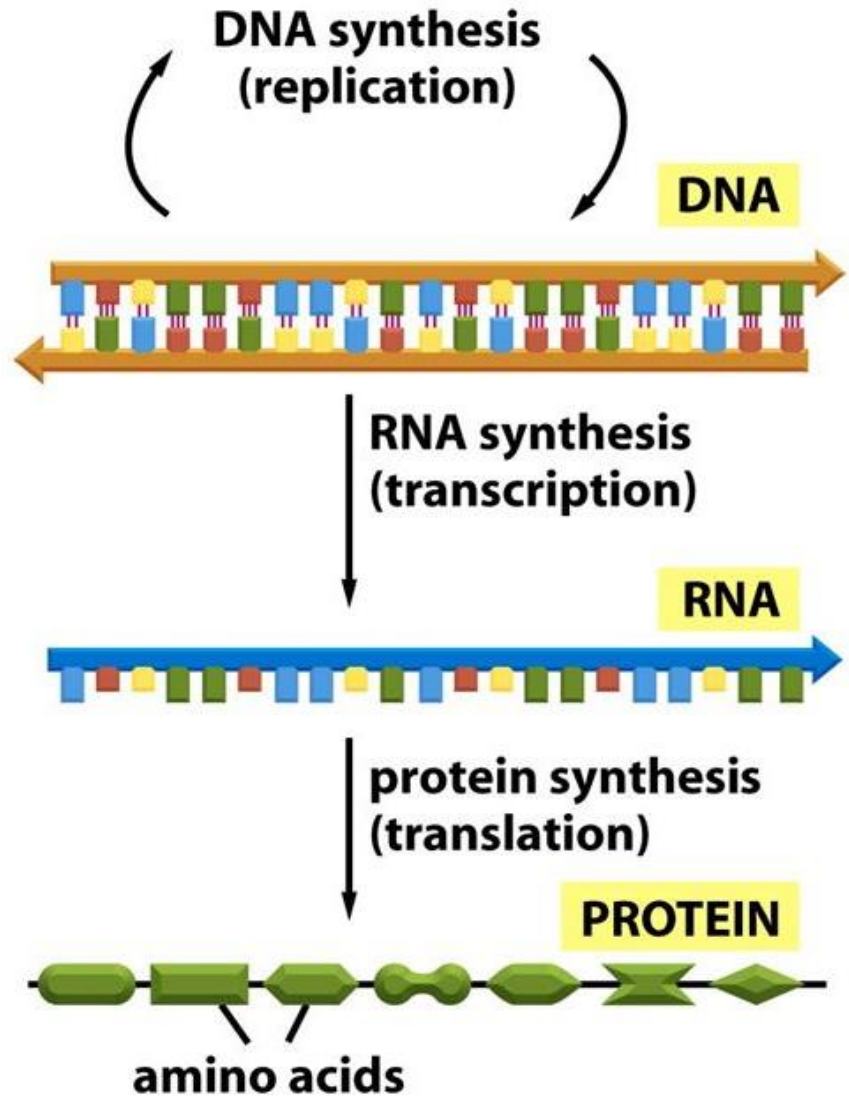
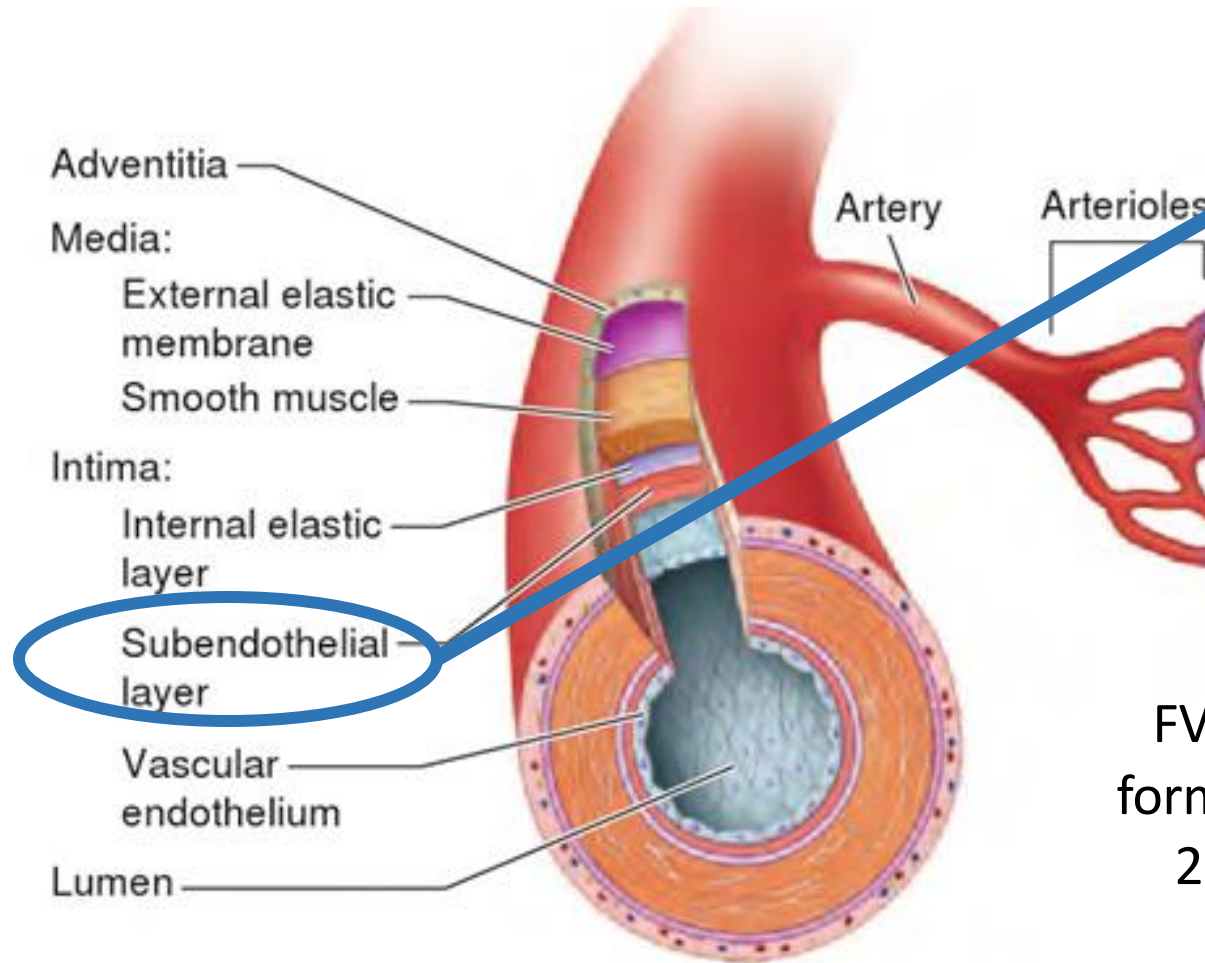
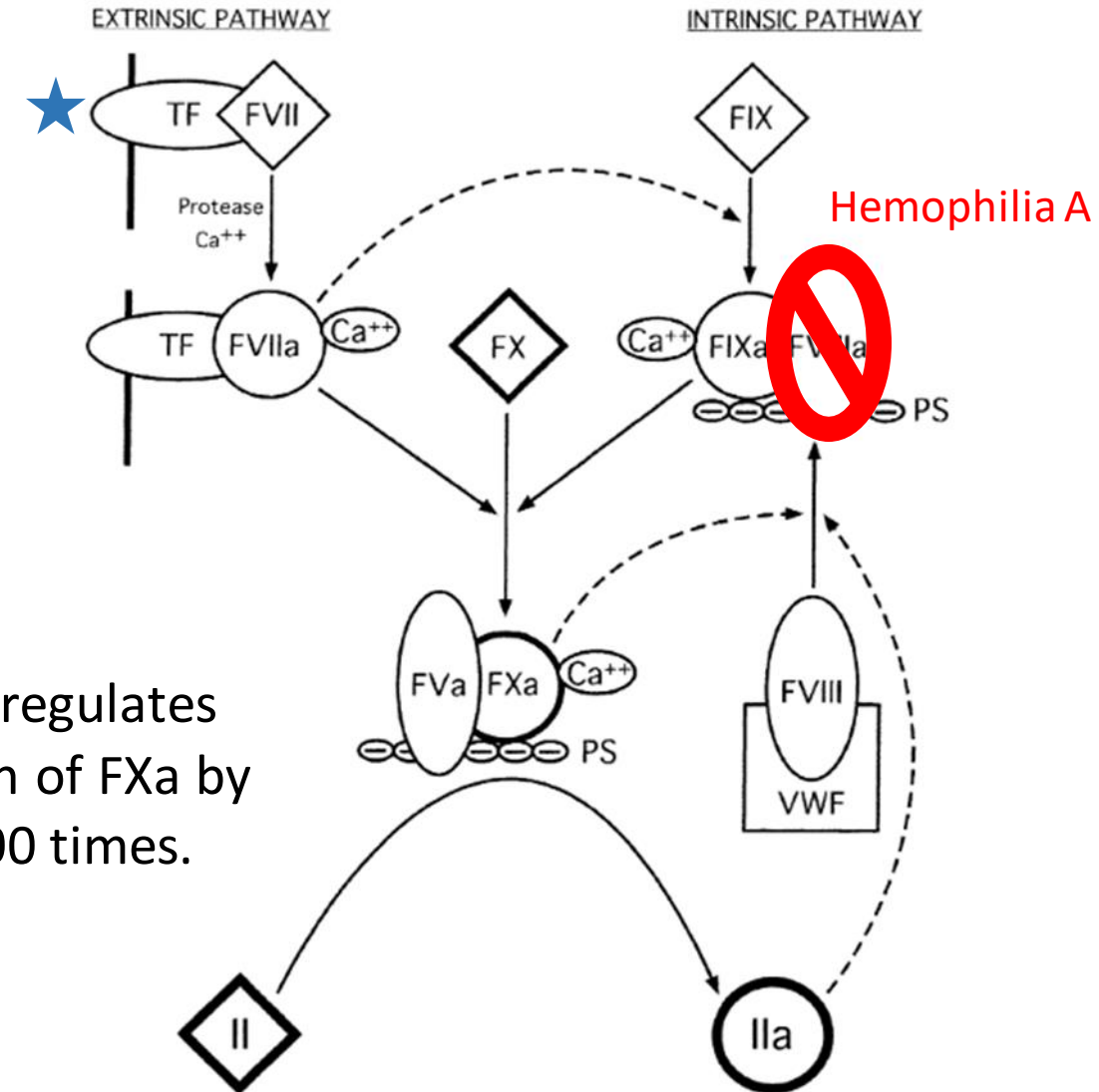


Figure 1-4 *Molecular Biology of the Cell*, Fifth Edition (© Garland Science 2008)

Hemostasis



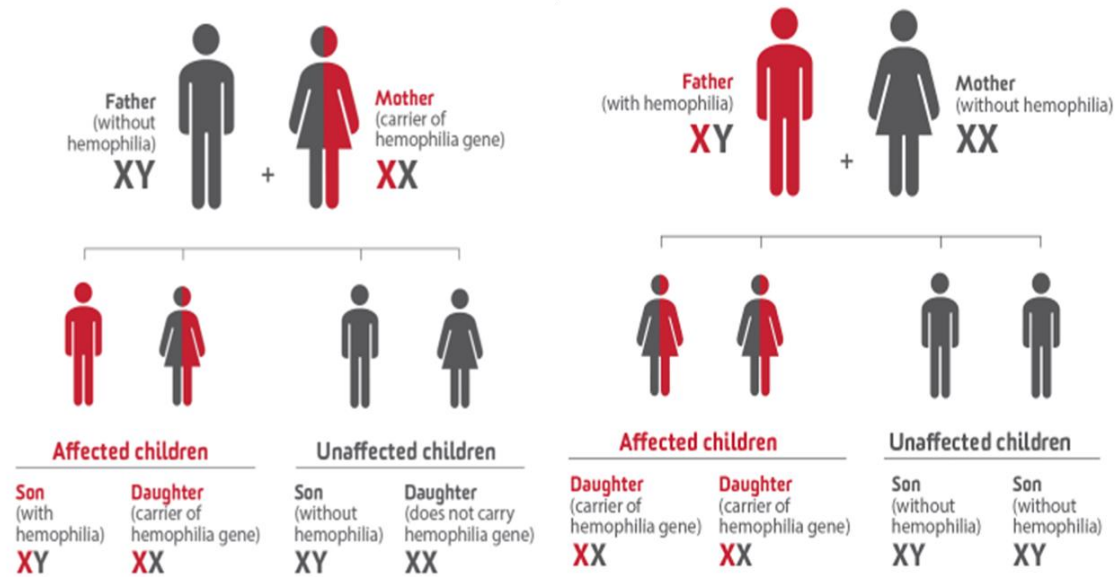
Blood Coagulation Cascade



FVIII upregulates formation of FXa by 200,000 times.

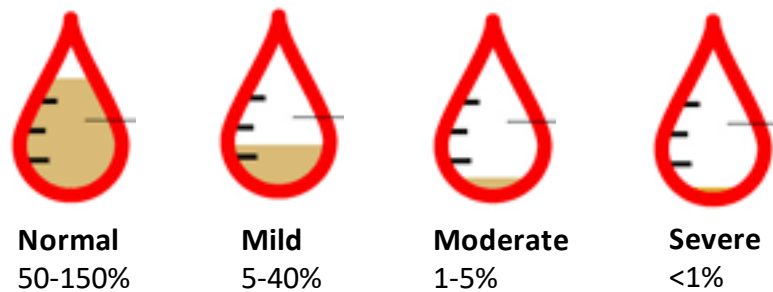
Hemophilia A and Factor VIII (FVIII)

- X-linked disease that affects 1 in 5000 males worldwide



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Hemophilia Degrees of Severity (% FVIII or FIX)



Resolution: 3.7Å
SpaceGroup: 4₁2₁2



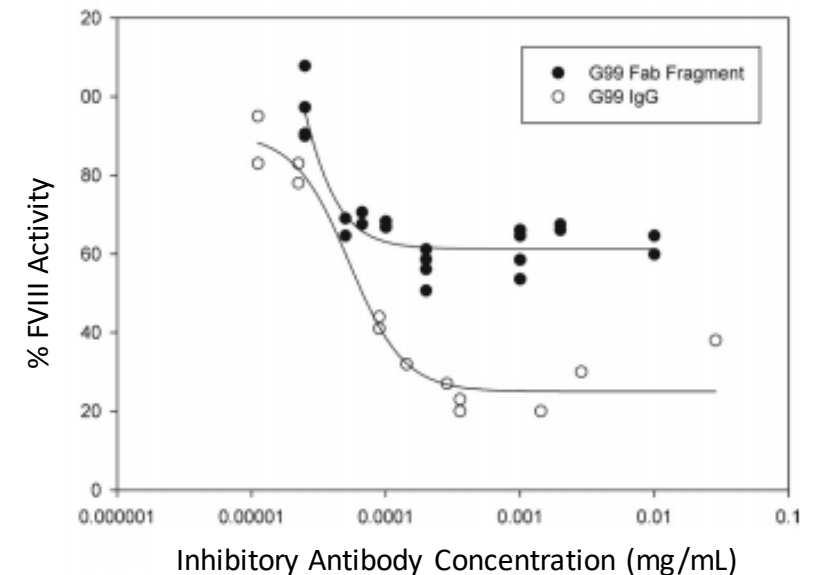
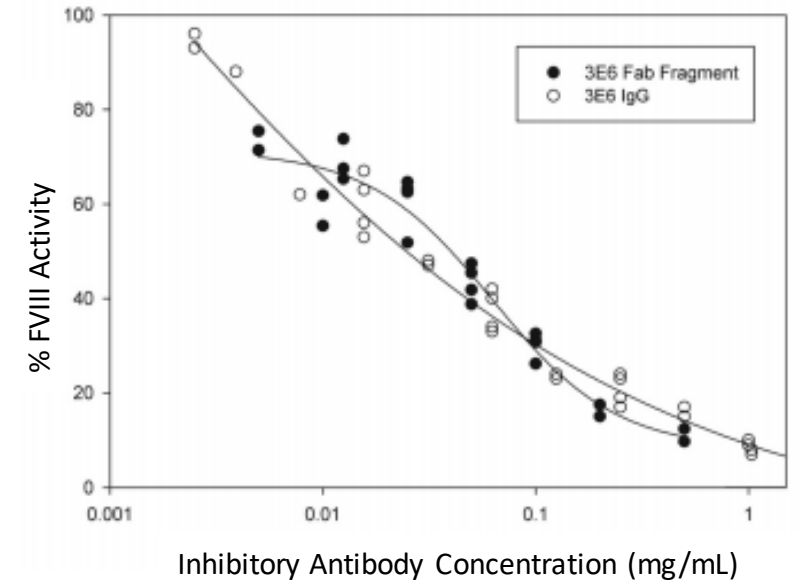
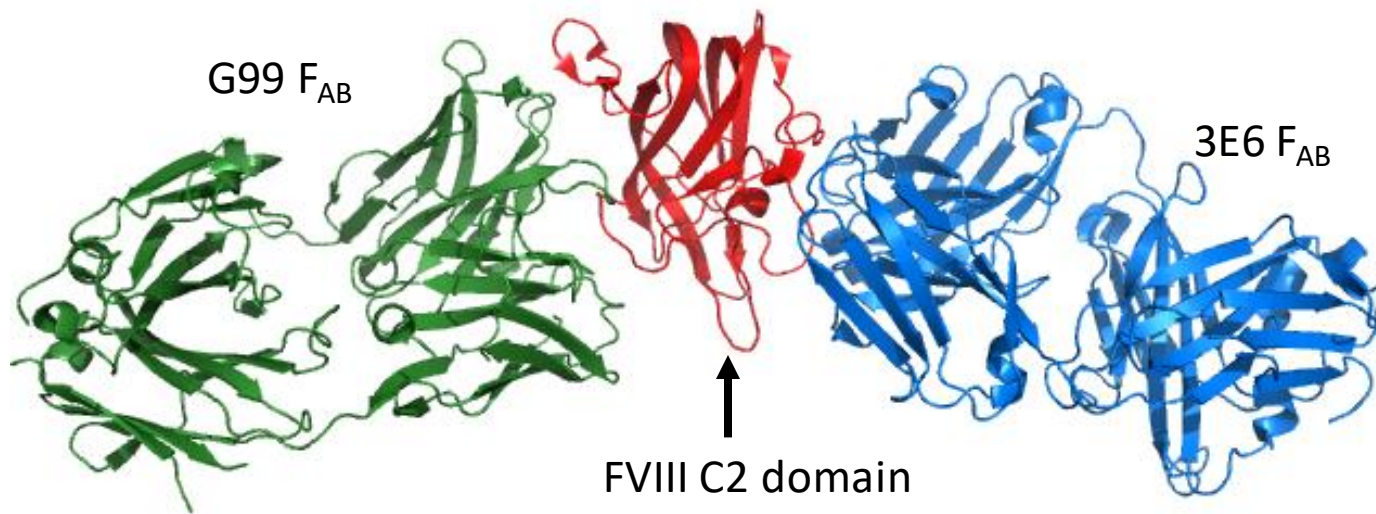
- Hemophilia A replacement therapy with recombinant FVIII

Shen et. al, *Blood*, 2008, 111, 1240-1257

Treatment Complication

- ~30% patients receiving FVIII concentrate replacement therapy acquire an inhibitory alloantibody.

Bray et al, *Blood*, 1994, 83 (9), 2428-2435



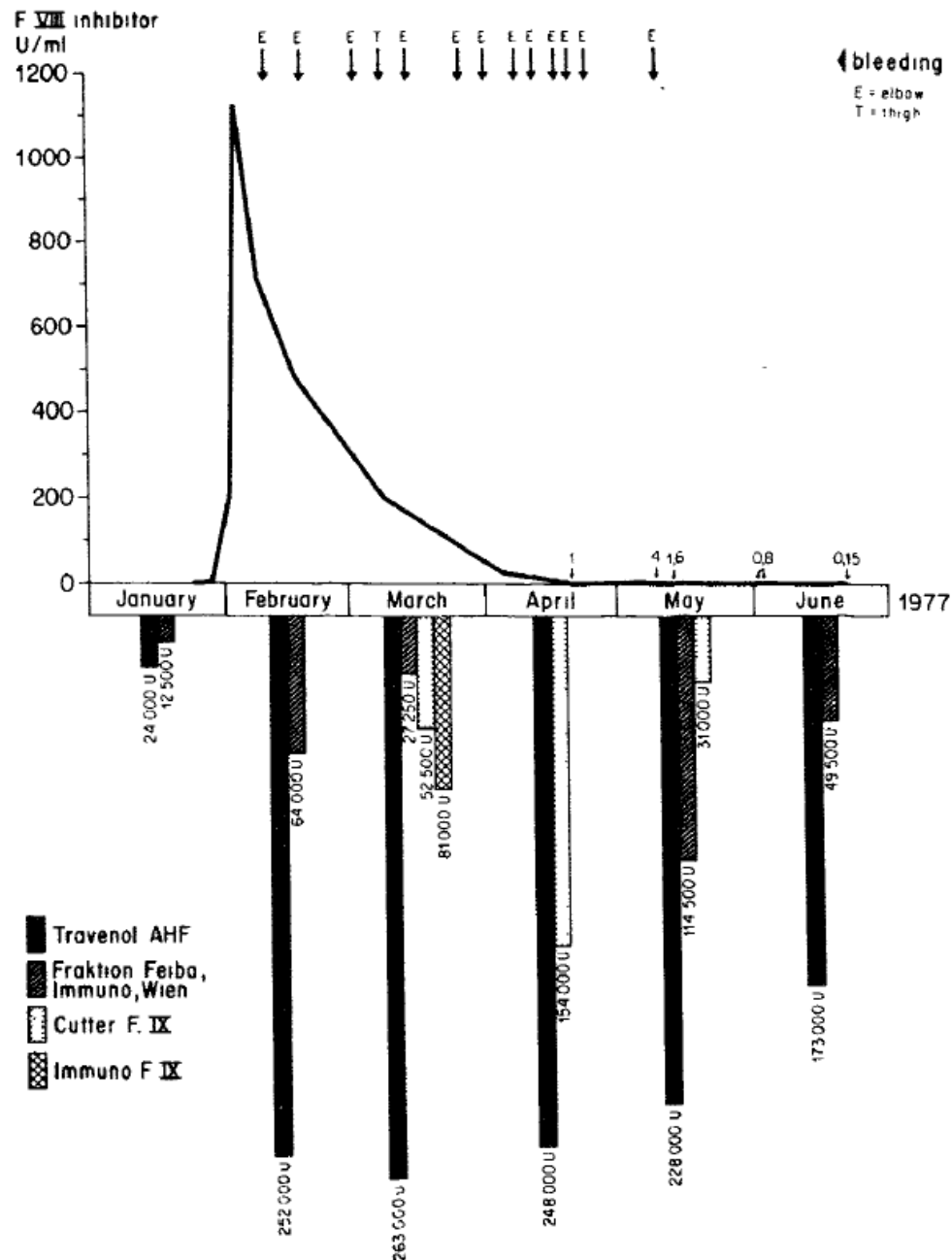


Fig.—Factor VIII inhibitor during treatment.

Brackmann, H.h., and J. Gormsen. *The Lancet*, 1977, 310, p.933.

Immune Tolerance Induction (ITI)

Table 1

Predictors of success following immune tolerance induction (ITI): data from the International Immune Tolerance Registry (IITR) and North American Immune Tolerance Registry (NAITR) [5,11].

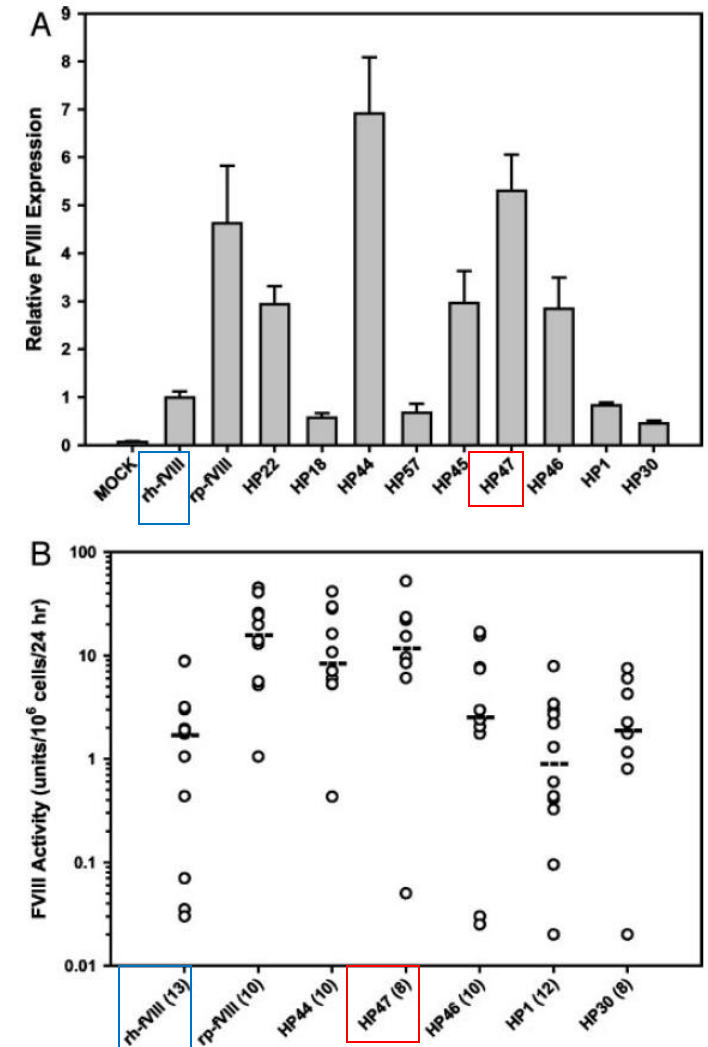
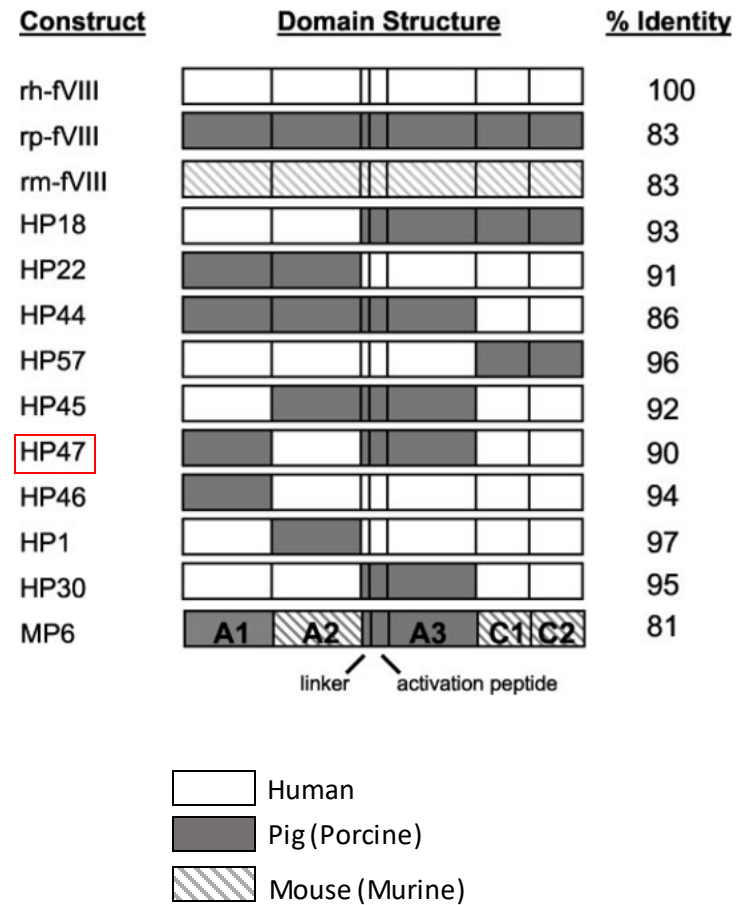
Variable	Success rate (%)		Cutoff for 'good risk'
	IITR	NAITR	
Age (years)			
< 20	70–78		< 20 years
> 20	40		
Peak titer (BU/mL)			
< 20	85–89		< 20 BU/mL
> 20	44–58		
Pre-ITI titer (BU/mL)			
< 10	78–79	83	< 10 BU/mL
> 10	41–65	40	
Dose (IU/kg/day)			
< 200	48–66	72–83	> 200 IU/kg/day
> 200	86	41	

BU, Bethesda unit.

Astermark, *Thrombosis Research*, 2011, 127, S6-S9

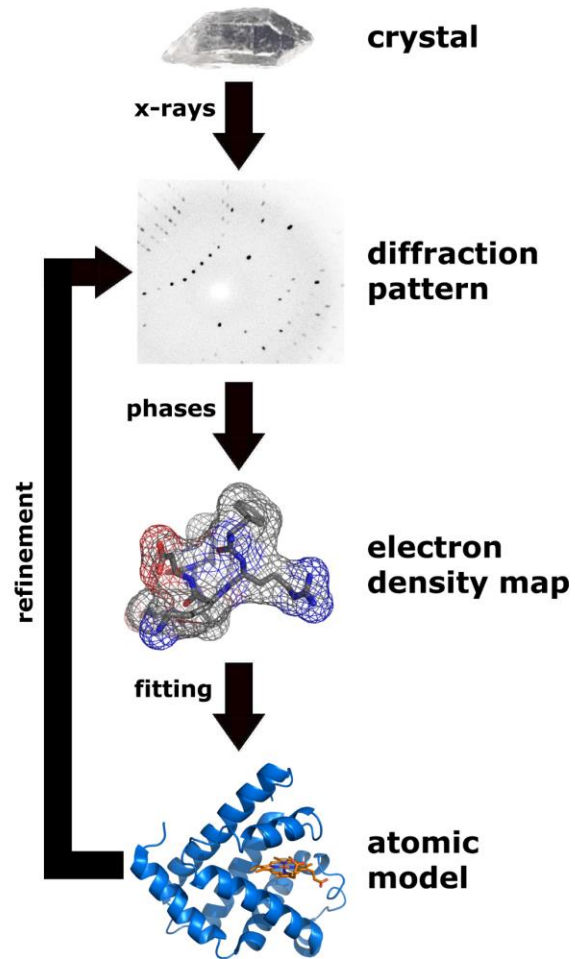
Human / Porcine Chimeric FVIII (HP47)

- hFVIII expresses poorly.
[Serum] = ~0.5nM
- Recombinant porcine FVIII displays increased cellular secretion
- Human/Porcine Chimera as high output protein therapeutic
- HP47 demonstrates higher expression and comparable activity to hFVIII

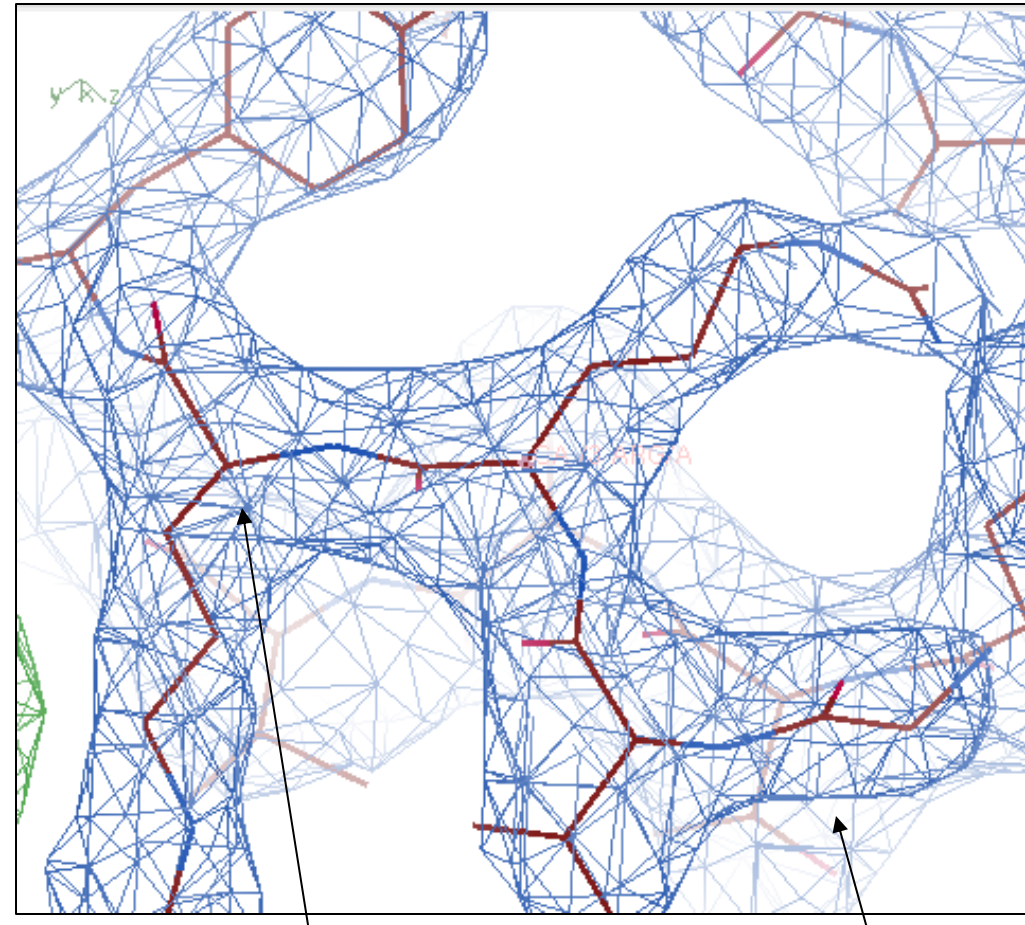


Protein Crystallography

Building the FVIII molecular model



T. Splettstoesser



FVIII amino acids
(red)

Electron density
(blue wire mesh)

- Assess model agreement to electron density with R factors
 - R_{work} and R_{free}
- Scale of 0.0 (perfect fit) - 0.6 (random/poor fit)

HP47 Crystal Structure

- At 3.2Å, highest resolution FVIII model to date
 - More confidently rebuild sections

HP47 Model Overview

- ~85% of the 1467aa HP47 sequence built into model

Quality of the current model

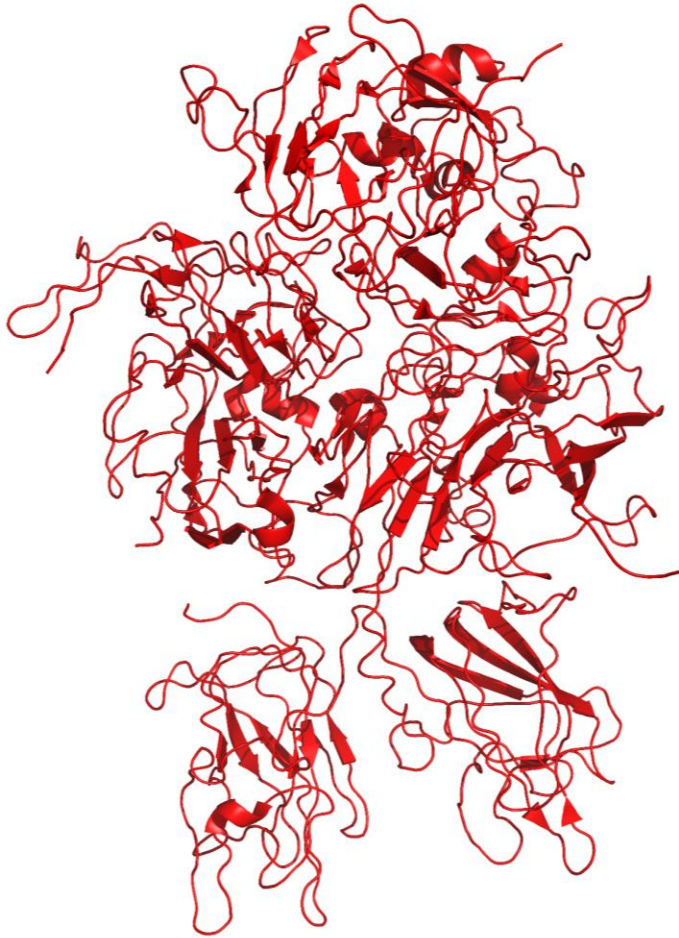
- R_{work} : 0.1972
- R_{free} : 0.2863
- Ramachandran outliers: 10.40%
- Space Group: $P2_1$



HP47



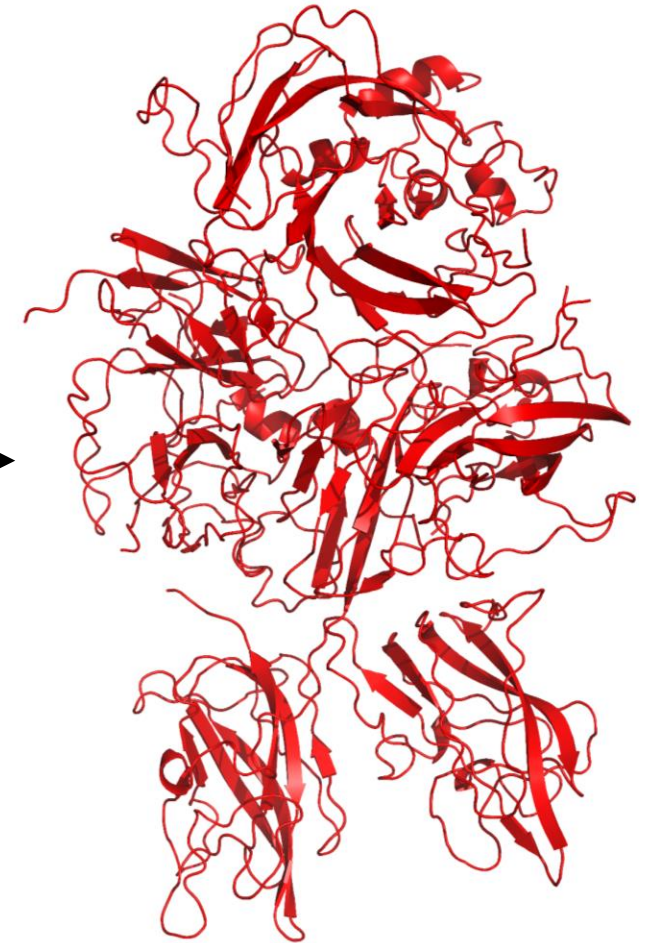
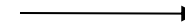
Improvements to the Human Model FVIII



2008 FVIII Structure (PDB: 2R7E)



New human/porcine FVIII structure



Newly refined FVIII Structure

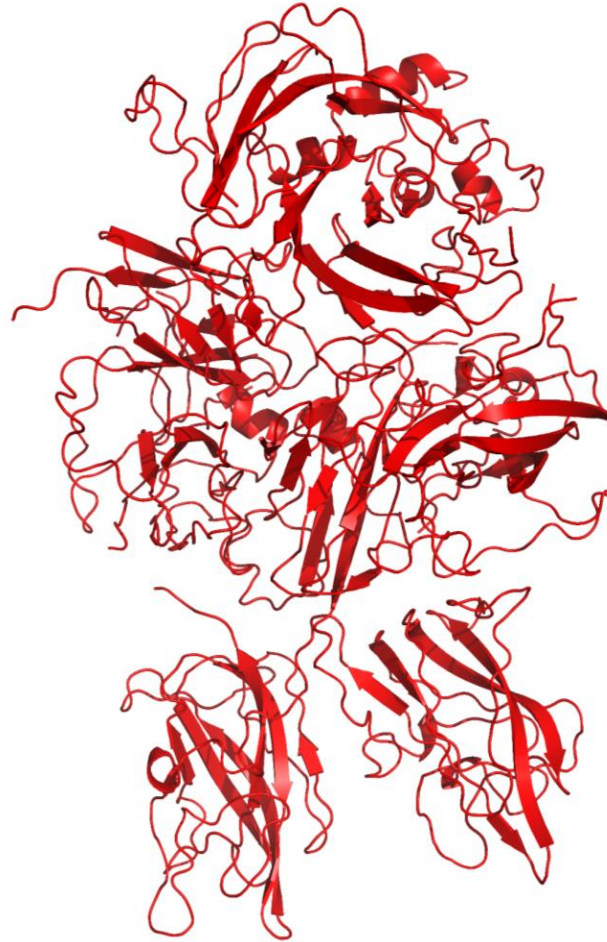
Improvements to the Human Model FVIII



HP47



New human/porcine FVIII structure



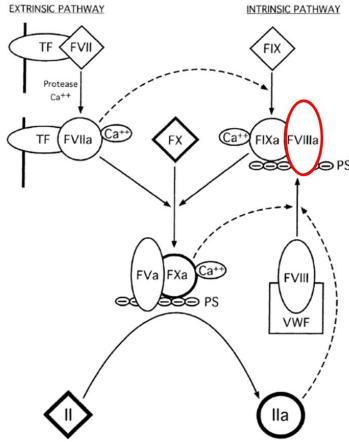
Human FVIII



Newly refined FVIII Structure

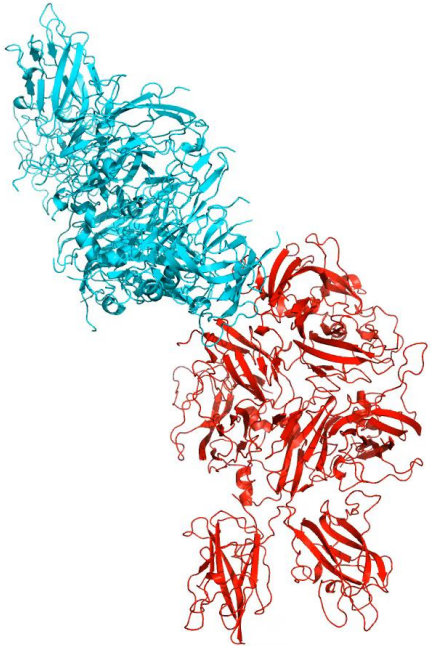
- Enhance basic understanding of FVIII's function in coagulation
- Develop more effective Hemophilia A therapeutics

Summary



- Hemophilia A is a X-linked disease caused by a lack of blood coagulation factor VIII (FVIII) protein.

- Patients who develop an immune response to FVIII rely on large quantities of improved therapeutics.



- Determined crystal structure of novel Human/Porcine Chimeric therapeutic. Improved human FVIII protein model.



Questions?

