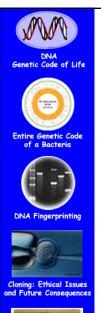


HC70A, SAS70A, & PLSS599 Winter 2022 Genetic Engineering in Medicine, Agriculture, and Law

Professors Bob Goldberg, John Harada, & Channapatna Prakash

Lecture 5 - Part Two How Are Genes Cloned & Engineered? *The Factor XIII Story*

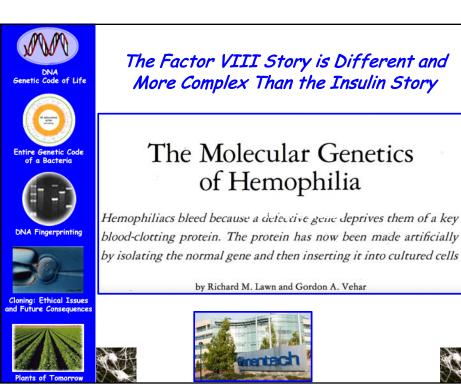
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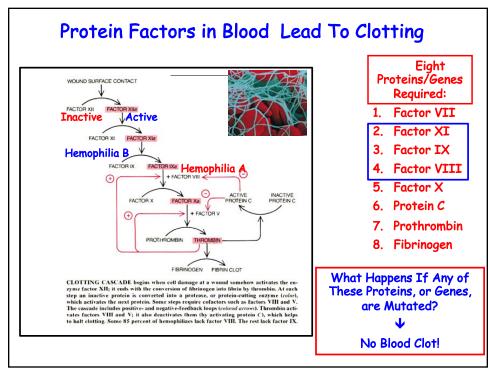
THEMES

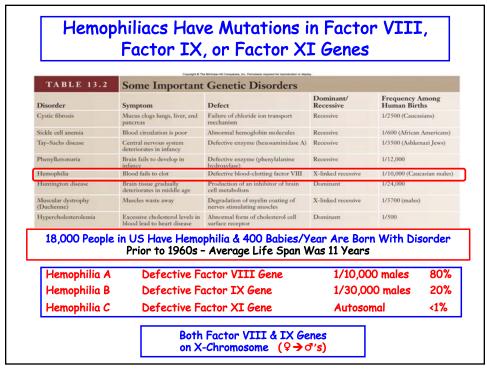
- 1. How Did the Supreme Court Indirectly Give Rise to the Biotechnology Industry?
- 2. What Strategies Were Developed For Cloning Insulin mRNA and Expressing Insulin in Bacterial Cells? What Strategy "Won" Out?
- 3. What is Hemophilia and How is it Inherited?
- 4. How Can a Disease Gene Be Found When It is Not Known Where the Gene is Expressed?
- 5. What Vectors Can Be Used For Cloning DNA?
- 6. What is the Advantage of Using a Virus Vector For Constructing Genome Libraries?
- 7. How To Make a Library of the Human Genome?
- 8. How Find a Gene With Only a Knowledge of the Protein Sequence?
- 9. How Use DNA Testing to Detect Factor VIII Disease Alleles?
- 10. How Isolate a Factor VIII cDNA Clone?
- 11. Genomic vs. cDNA Libraries
- 12. How Produce Factor VIII Protein For Use as a Drug

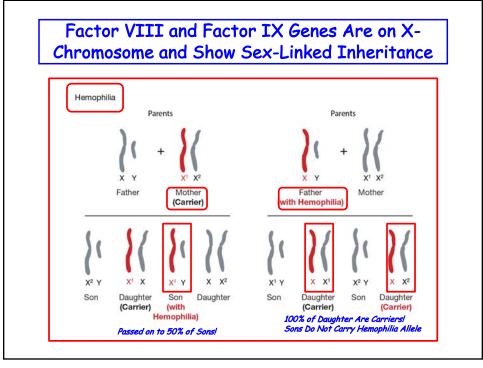
Plants of Tomorrow

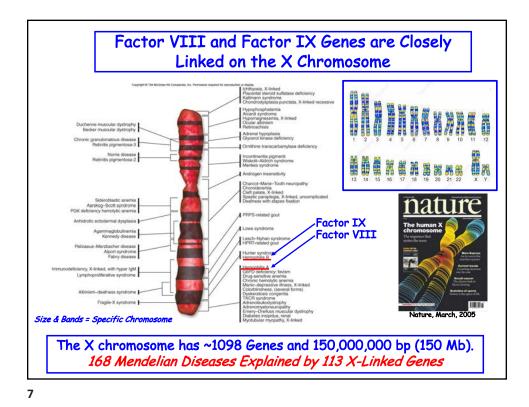


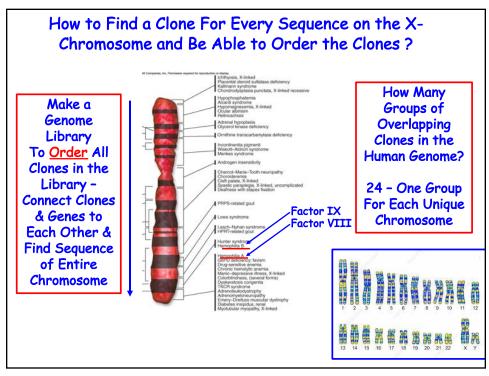


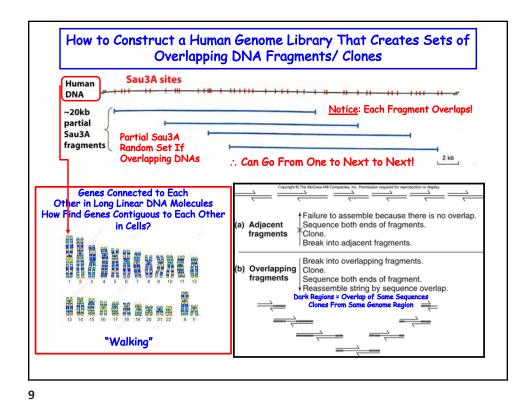


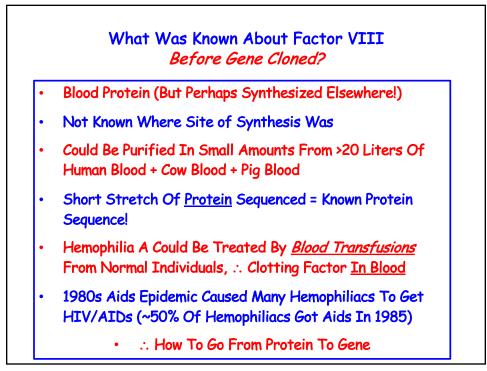


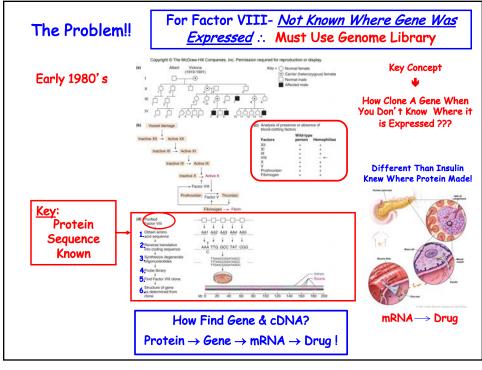


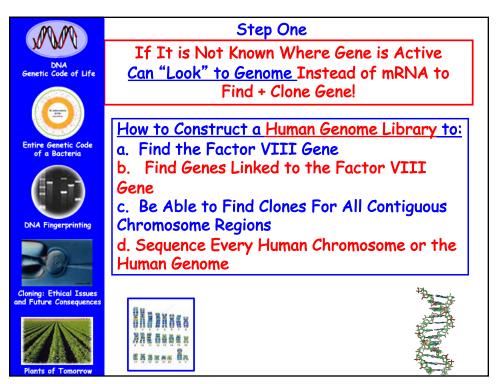


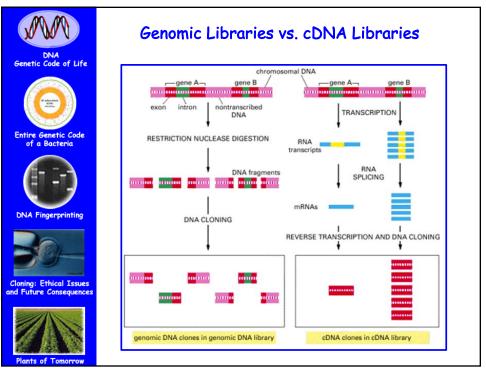


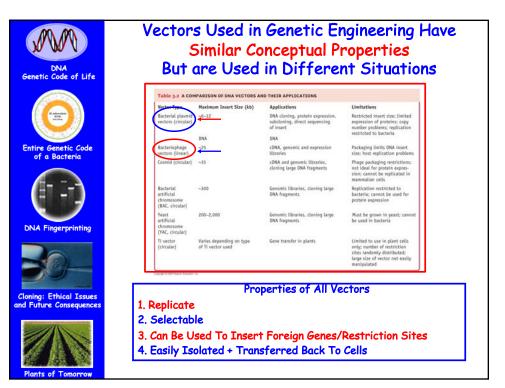


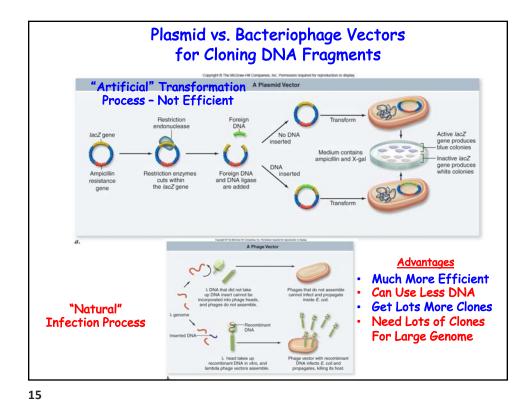


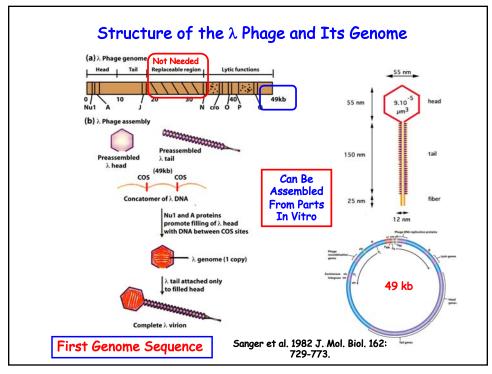


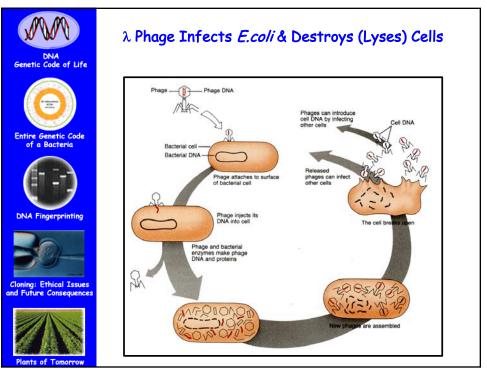


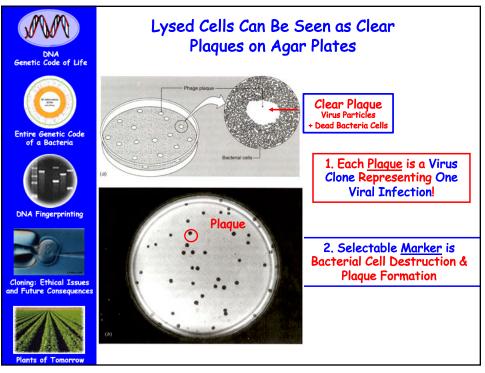


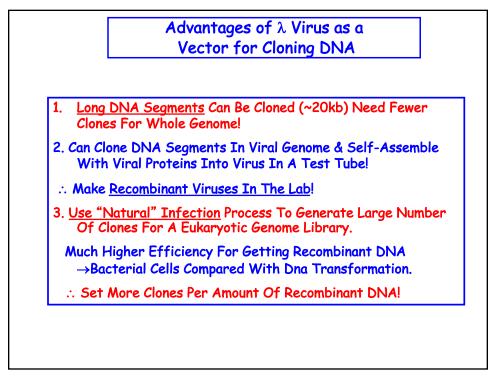




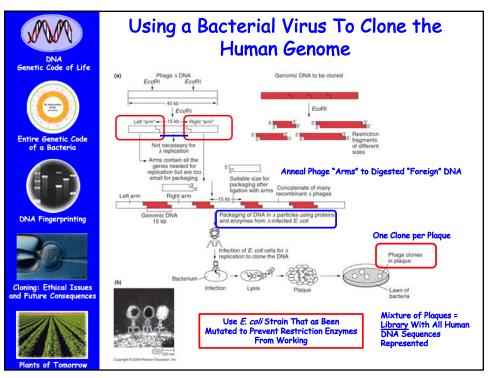


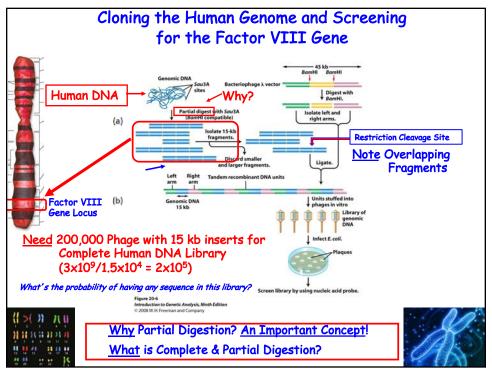


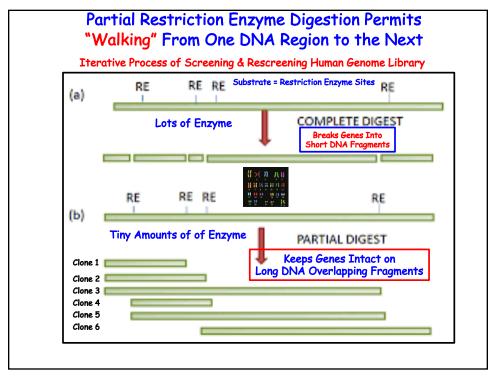


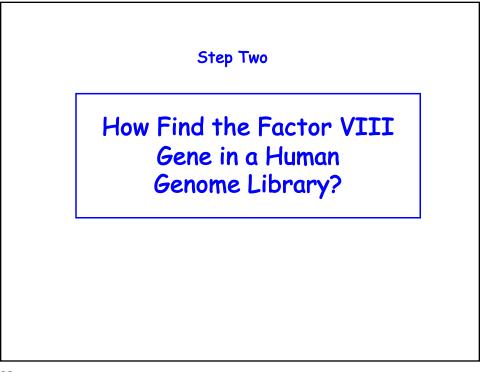


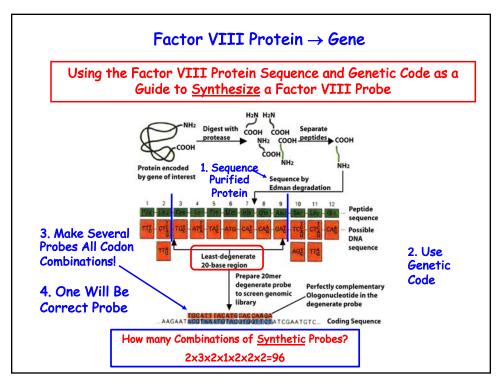


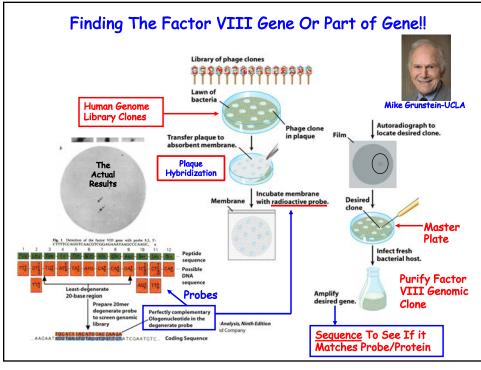


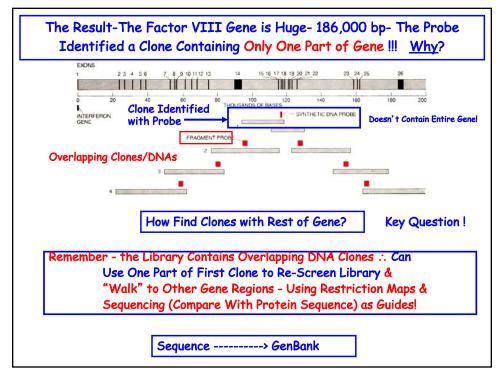


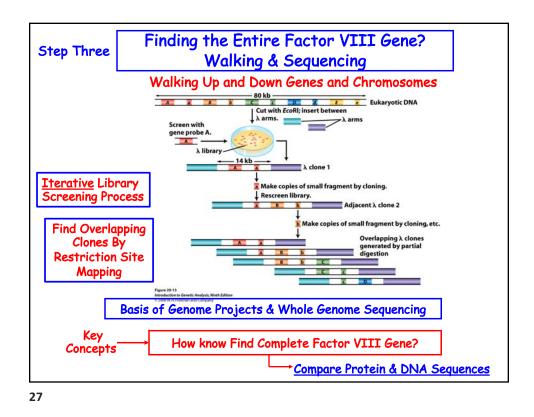


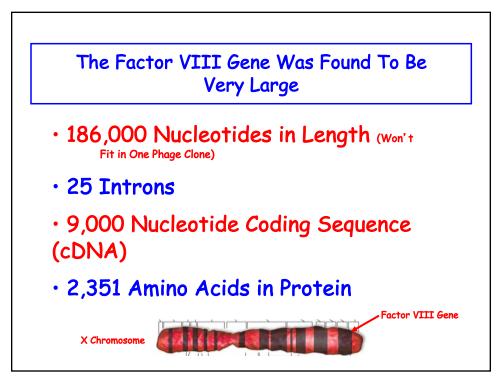








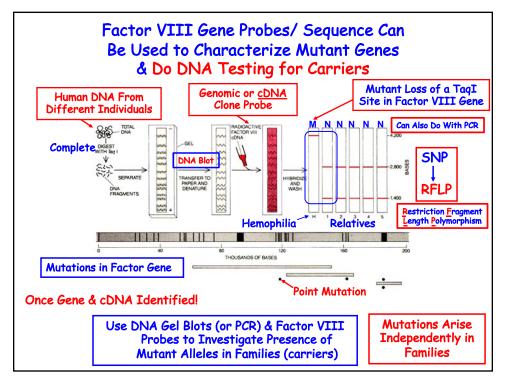


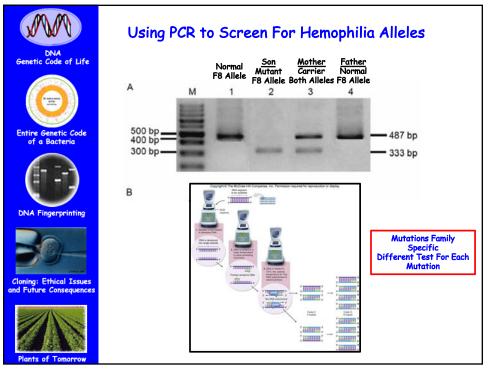


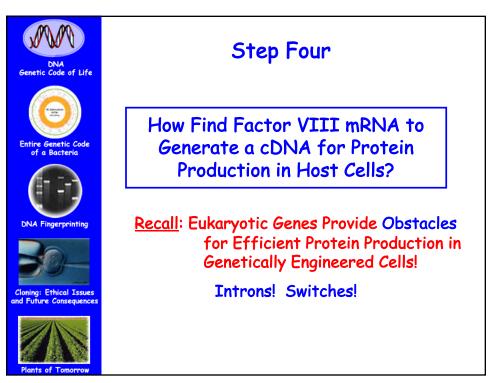
Factor VIII SNP Mutations Occur Throughout the Gene

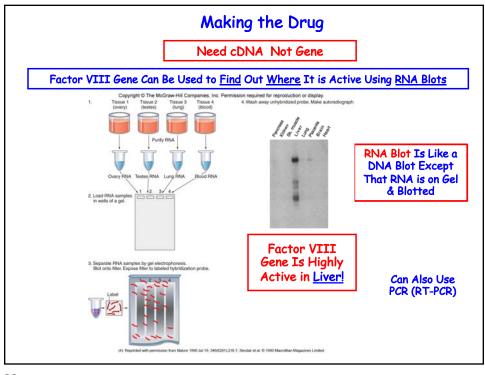
[Haemophilia 11, 481-491 (2005)] Larger the Gene - Larger Number of Mutations!

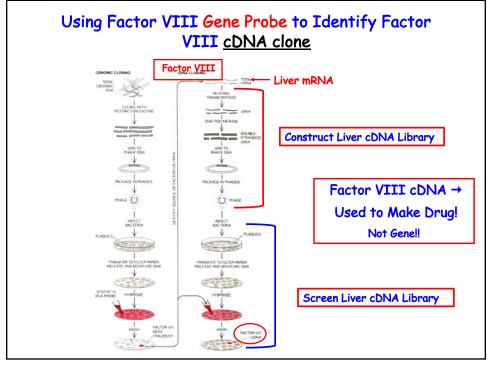
VIII:C (%)	Family history	Consanguinity*	Inversion	Codon†	Mutation	Amino acid change	Exon	Conservation:
1	Sporadic	NC	Normal	- 51	$TTT \rightarrow TCT_3$	Phe -> Ser	2	FFFF, identical
1.20	Sporadic	NC	Normal	80	$GTT \rightarrow GAT$	$Val \rightarrow Asp$	3	VVVV, identical
1	Sporadic	NC	Normal	102	GGT → GTTS	$Gly \rightarrow Val$	3	GGGG, identical
2	Sporadic	NC	Normal	104	TCC → CCCS	Ser -> Pro	3	SSSS, identical
6	Sporadic	NC	Normal	143	GAG -> AAGS	Glu → Lvs	4	EEEE, identical
1	Sporadic	NC	Normal	233	delCA§	Thr \rightarrow fs (TGA-264)	6	
2.70	Inherited	NC	Normal	32.1	$GAA \rightarrow AAA$	Glu -> Lys	8	EEEE, identical
0	Sporadic	NC	Normal	372	$CGC \rightarrow CAC$	Arg -> His	8	RRRR, identical
1	Inherited	NC	Normal	527	$CGG \rightarrow TGG$	Arg -> Trp	11	RRRR, identical
1	Sporadic	NC	Normal	52.8	TGC → TAC§	Cys -> Tyr	11	CCCC, identical
1	Inherited	NC	Normal	592	$CAA \rightarrow TAA$	Glu -> Stop	12	QQQQ, identical
1	Inherited	NC	Normal	864	delGACA	Gly → fs [TAA-867]	14	
					insCAATTAAATGAGAAS	and the formation of		
1	Sporadic	NC	Normal	948	insA§	Lys \rightarrow fs (TGA-984)	14	
1	Sporadic	NC	Intron 1	1107	AGG → TGG§	Arg -> Trp	14	RGKK, dissimilar
	Sporadic	NC	Normal	1107	$AGG \rightarrow TGGS$	$Arg \rightarrow Trp$	1.4	RGKK, dissimilar
1	Inherited	NC	Normal	1191-1194	delA	$llc \rightarrow fs$ (TAG-1198)	14	restored assessment
.40	Sporadic	NC	Normal	1191-1194	insA	Ile \rightarrow fs (TAA-1220)	14	
	Sporadic	C	Normal	1227	delCN	Leu \rightarrow fs (TGA-1231)	14	
2.10	Sporadic	NC	Normal	1241	$GAC \rightarrow GAG$	Asp → Glu	14	DGGE, similar
1	Sporadic	NC	Normal	1392	1392dcl14185	Pro → fs (TAG-1446)	14	to cover of another
1	Incrited	C	Normal	1392	1392del14185	$Pro \rightarrow fs (TAG-1446)$	14	
	Sporadic	NC	Normal	1441	insAS	nu - u(mo mu)	14	
1	Inerited	C	Normal	1441	insAS			
1	Inherited	NC	Normal	1502	CAG → TAG§	Gln → Stop	14	QREQ, dissimilar
1	Inherited	NC	Normal	1.504	delGTS	Val \rightarrow fs (TGA-1517)	14	dand annum
19	Sporadic	NC	Normal	1.535	$TGG \rightarrow TGA$	Trp → Stop	14	WLWM, dissimilar
hibitor 96 BU	oporane	140	14010man	1333	100 - 104	rip - Jobp	1.4	with with a dissimilar
1	Sporadic	NC	Normal	1571	$TAT \rightarrow TAAS$	Tyr -> Stop	14	Y-YY, dissimilar
	Sporadic	NC	Normal	1581	$AAA \rightarrow TAAS$	Lys \rightarrow Stop	14	KEKK, dissimilar
1.20	Sporadic	NC	Normal	1696	$CGA \rightarrow GGA$	$Arg \rightarrow Gly$	14	RRRR, identical
1.80	Sporadic	NC	Normal	1729	delAs	$Gln \rightarrow fs (TAA-1752)$	15	KINNS, MURLAN
1.00	Inherited	NC	Normal	1751	GAA -> AAAS	$Glu \rightarrow Lys$	15	EEEE, identical
	Sporadic	NC	Normal	1775	TTC -> TCCS	Phe \rightarrow Pro	16	FFFF, identical
2 -	Sporadic	NC	Normal	1835	$TGG \rightarrow TGAS$	$Trp \rightarrow Stop$	16	WWWW, identical
7.60	Sporadic	C	Normal	1882	$ATC \rightarrow ATAS$	$lip \rightarrow lip$	17	IIII, identical
2	Inherited	c	Normal	1966	$CGA \rightarrow CAA$	$Arg \rightarrow Glu$	18	RRRR, identical
	Sporadic	NC	Normal	1966	$CGA \rightarrow TGA$	$Arg \rightarrow Gau$ $Arg \rightarrow Stop$	18	RRRR, identical
1.7	sporadic	NG	Normal	1966	$\cos a \rightarrow 16A$	wik -> arob	15	RARA, stentical

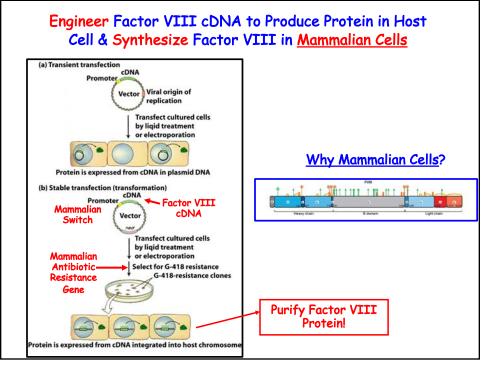


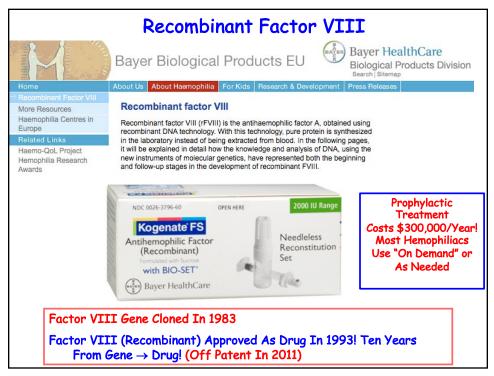












The Future: Gene Therapy - A Permanent "Cure"

December 10, 2011

Treatment for Blood Disease Is Gene Therapy Landmark

Partners with

Gene Therapy Shows Promise for Treating Hemophilia

The First Ever In-Human Gene Editing Will Try and Combat Factor IX - Hemoglobin B Hemophilia FDA-Approved Clinical Trial 2016

MM		The Factor VIII Story - A Summary
	1.	Purify Small Amounts of Factor VIII
DNA metic Code of Life	2.	Obtain Partial or Complete Amino Acid Sequence
	З.	Use the Genetic Code to Synthesize Degenerate DNA Probes
	4.	Isolate Factor VIII DNA Clones Complementary to Probe in Genome Library
	5.	Determine if Factor VIII Clones Contain the Complete Gene By Sequencing and Comparing With Protein Sequence
tire Genetic Code of a Bacteria	6.	If Not, "Walk" to Obtain Overlapping DNA Clones That Collectively Contain the Factor VIII Gene
(+ <u>-</u> -	7.	Sequence Clones To Determine Where the Factor VIII Gene Starts and Stops
	8.	Use Factor VIII Genome Probe to Find Out What Body Organ/Tissue Expresses the Factor VIII Gene
NA Fingerprinting	9.	Make a cDNA Library From the Target Organ/Tissue and Isolate a Factor VIII cDNA Clone
<u> 90</u> -	10.	Sequence the Factor VIII cDNA Clone and Compare With Factor VIII Gene Sequence to Map its Anatomy (I.e., introns, exons, swtiches) and Ensure That it Contains the Complete Protein Coding Sequence
ning: Ethical Issues Future Consequences	11.	Use Factor VIII cDNA and/or Genome Fragments as a Probe to Find RFLP Markers For Disease Alleles Or Sequence Disease Alleles to Find Relevant RFLP Markers By Comparison With Wild-Type Sequence
	12.	Insert Factor VIII cDNA Into an Expression Vector and Synthesize Factor VIII Protein in Host Cells (e.g., Mammalian Cells)