

Entire Genetic Code of a Bacteria



**DNA** Fingerprinting



Cloning: Ethical Issues and Future Consequences



HC70A, SAS70A, & PLSS059 Winter 2018 Genetic Engineering in Medicine, Agriculture, and Law

Professors Bob Goldberg, John Harada, & Channapatna Prakash

Lecture 5 The Nuts & Bolts of Genetic Engineering: From Mutations to Pedigrees to Drug The Factor XIII Story





### THEMES

#### 1. PCR

- 2. What Causes Gene Mutations?
- 3. How Do Gene Mutations Lead to Genetic Variability?
- 4. How Can We Test For Gene Mutations at the DNA Level?
- 5. What is Hemophilia and How is it Inherited?
- 6. How Can a Disease Gene Be Found When It is Not Known Where the Gene is Expressed?
- 7. What Vectors Can Be Used For Cloning DNA?
- 8. What is the Advantage of Using a Virus Vector For Constructing Genome Libraries?
- 9. How To Make a Library of the Human Genome?
- 10. How Find a Gene With Only a Knowledge of the Protein Sequence?
- 11. How Use DNA Testing to Detect Factor VIII Disease Alleles?
- 12. How Isolate a Factor VIII cDNA Clone?
- 13. How Produce Factor VIII Protein For Use as a Drug?

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DNA Genetic Code of Life

Entire Genetic Code of a Bacteria

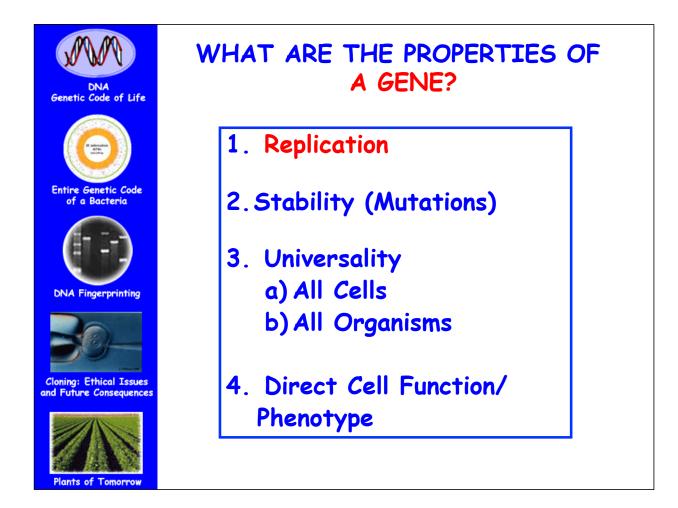


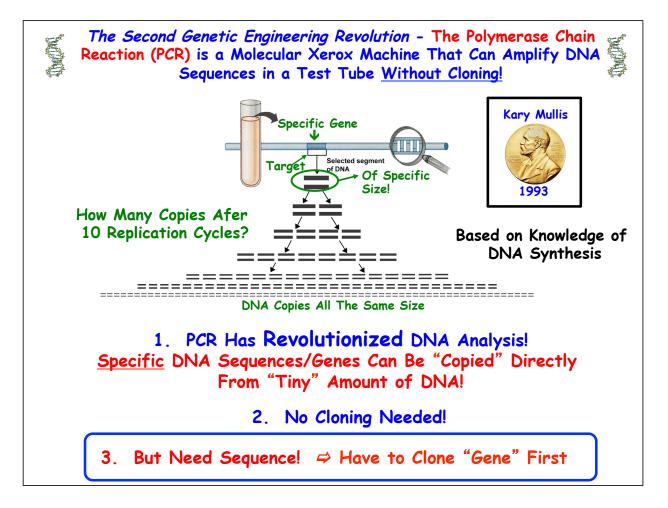
**DNA** Fingerprinting

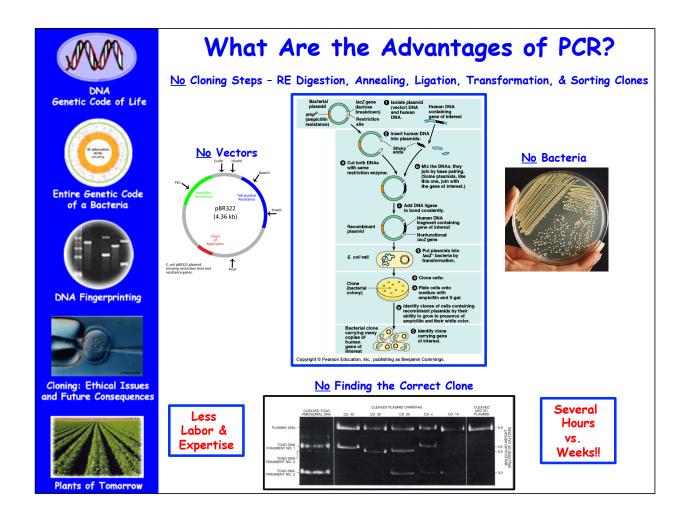


Cloning: Ethical Issues and Future Consequences





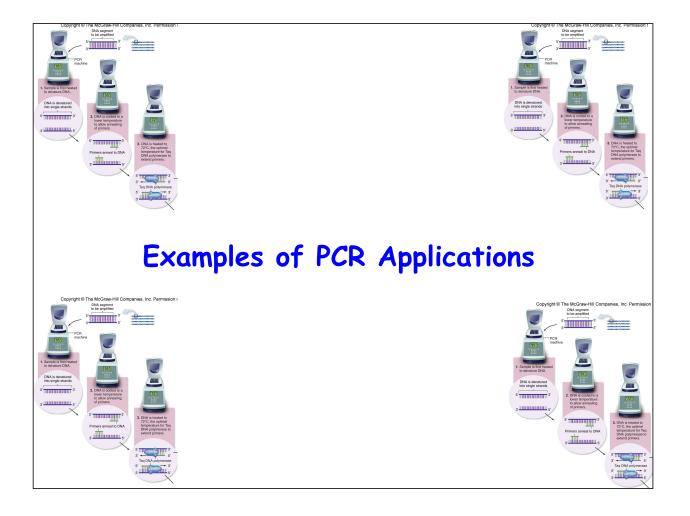


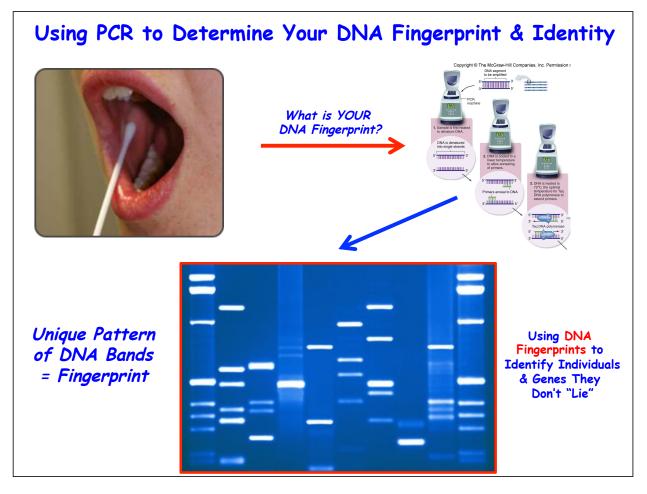


#### PCR Has Many Uses, Has Changed Many Fields, and Lead To New Ones That Have Had a Big Impact On Our Lives

- 1. <u>Amplify Any DNA Sequence</u>, or Gene, From "Tiny" Amounts of DNA or Biological Materials IF ORIGINAL SEQUENCE KNOWN
- 2. <u>Study DNA From Limited and/or Degraded Sources Such As:</u>
  - 1. A Single Human Hair or Cheek Cell
  - 2. An Ancient Fossil (e.g., Neanderthal Bone or Mammoth Hair)
  - 3. An Ancient Insect Trapped in Amber
  - 4. Human Remains (e.g., 9/11 Victims)
  - 5. A Single Human Embryo Cell
  - 6. Contaminated Meat To Determine the Causal Organism
- 3. Used In:
  - 1. DNA Fingerprinting-Individual Identification-Genetic Disease Screening
  - 2. Forensics (Crime Scenes, Mass Graves, Criminal Suspects, Wrongfully Convicted)
  - 3. Paternity & Family Relationships (e.g., Immigration, Tracing Lost Children)
  - 4. Disease Diagnosis & Pathogen Identification (Humans, Animals, & Plants)
  - 5. Human Origins & Migrations
  - 6. Ancient Genome Sequences & Evolutionary Studies
  - 7. Specific mRNA Detection
  - 8. "Cloning" Specific DNA Sequences
  - 9. Tracing Plant & Animal Sources (e.g., Poaching Stolen Cattle, Cactus)
- 4. <u>Need as Little as One Molecule of DNA</u> & Can Replicate an ∞ Amount of Specific Sequences

Revolutionized How To Study & Manipulate DNA





## Using PCR To Determine an Individual's Ancestry DNA Tribes Genetic Ancestry Analysis **Discover Your Past!** Determine if two pe Test your autosomal DNA inherited from maternal and paternal, lineal Determine if two ve test available World Regions sis with Add-Ons any t Our Premium Kit test now include 21 powerful STR marker systems PCR Started a New Industry Maternal & Paternal Testing Adopted? Find out about your ancestry DNA can reveal ancestors' lies and secrets LA Times, January 18, 2009

## Using PCR to Amplify Neanderthal Bone DNA & Sequence The <u>Entire</u> Genome!

# Analysis of one million base pairs of Neanderthal DNA From a 45,000 Year-Old Bone

Richard E. Green<sup>1</sup>, Johannes Krause<sup>1</sup>, Susan E. Ptak<sup>1</sup>, Adrian W. Briggs<sup>1</sup>, Michael T. Ronan<sup>2</sup>, Jan F. Simons<sup>2</sup>, Lei Du<sup>2</sup>, Michael Egholm<sup>2</sup>, Jonathan M. Rothberg<sup>2</sup>, Maja Paunovic<sup>3</sup><sup>‡</sup> & Svante Pääbo<sup>1</sup>

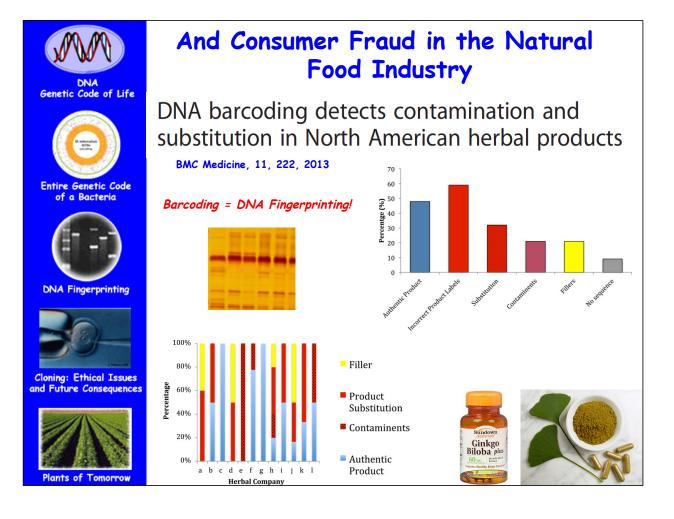


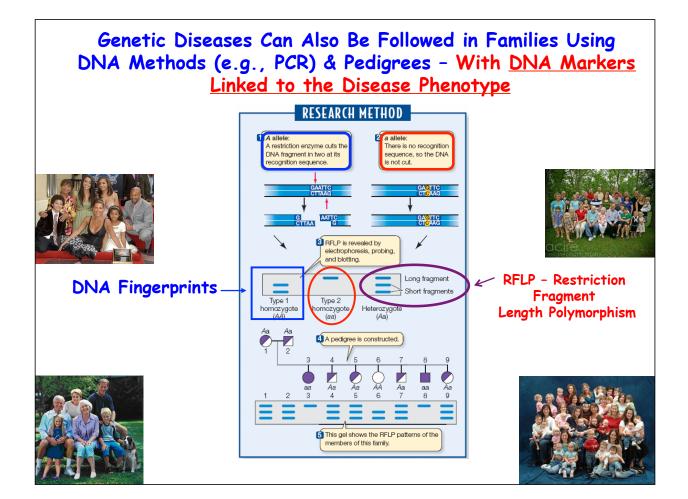
Nature, November, 2006

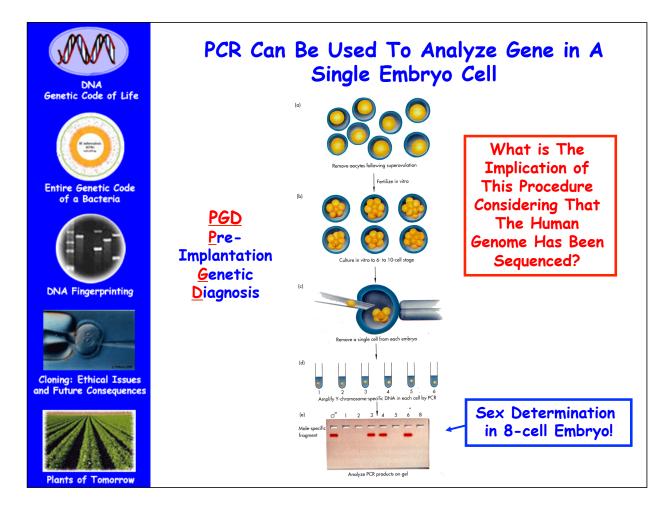


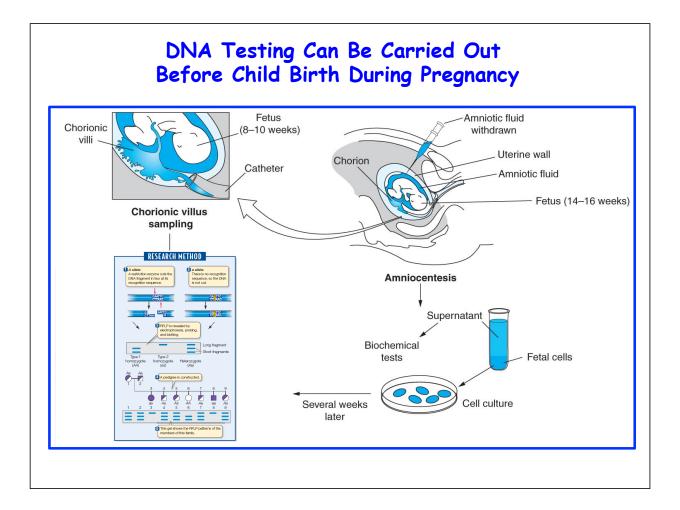
#### Using PCR To Detect Human Pathogens (Viruses, Fungi, Bacteria)

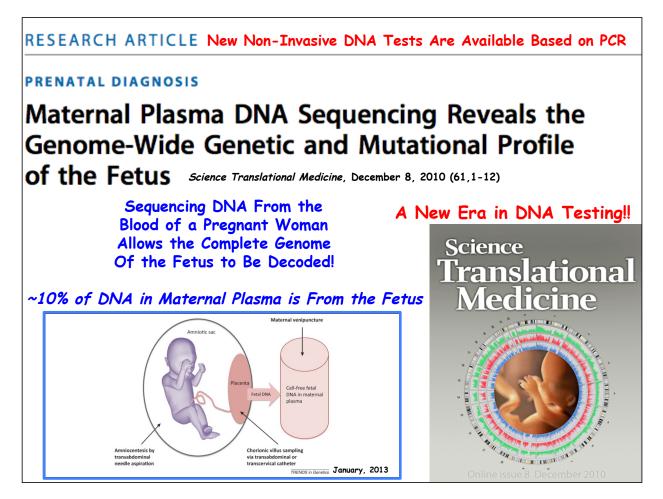


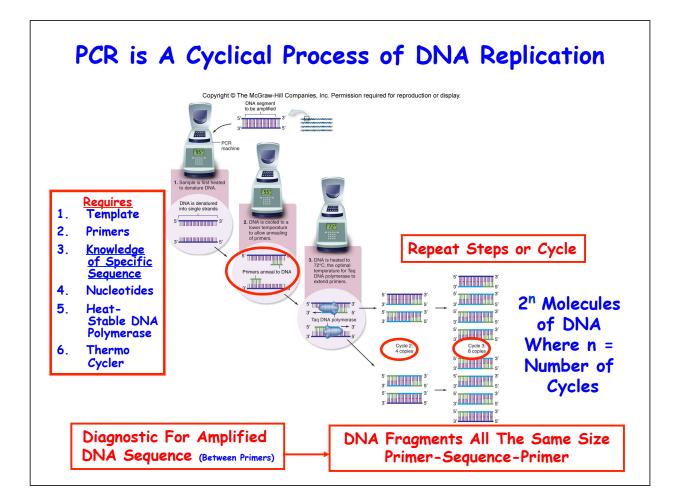


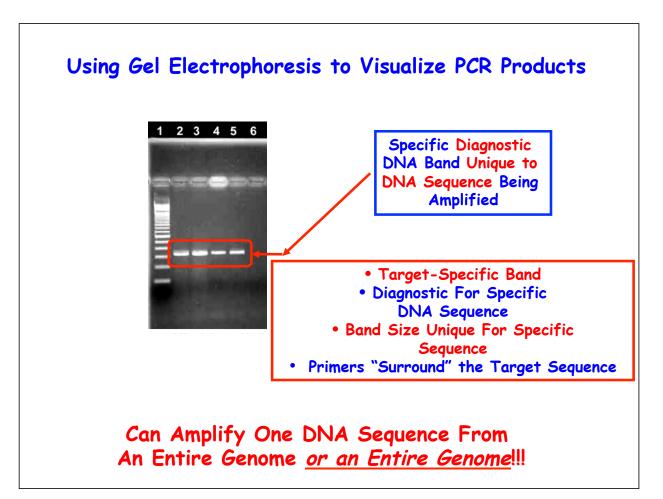








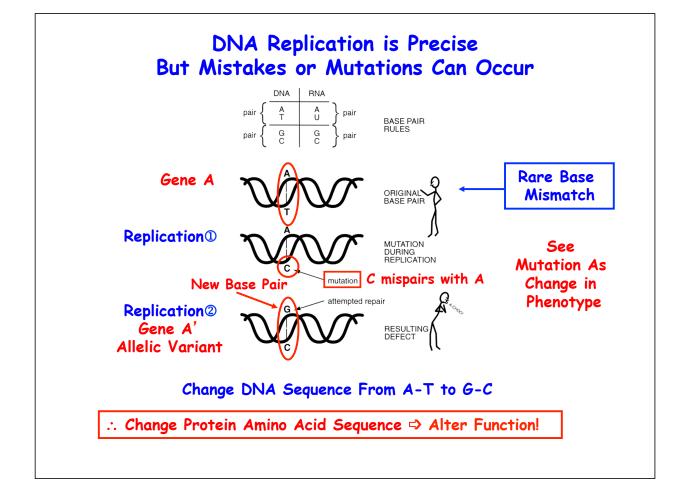


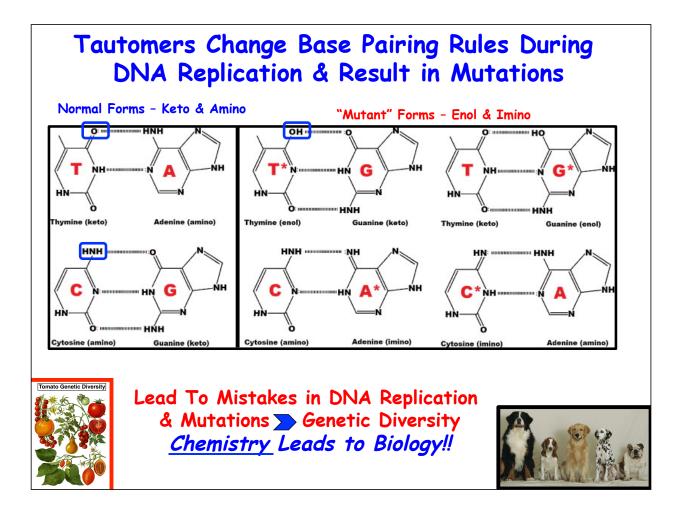


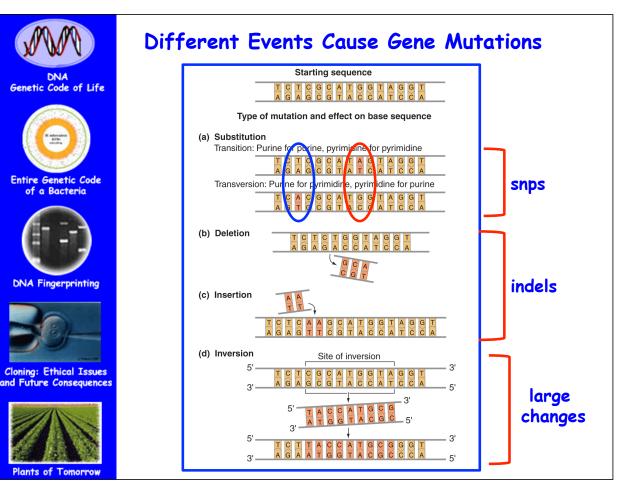


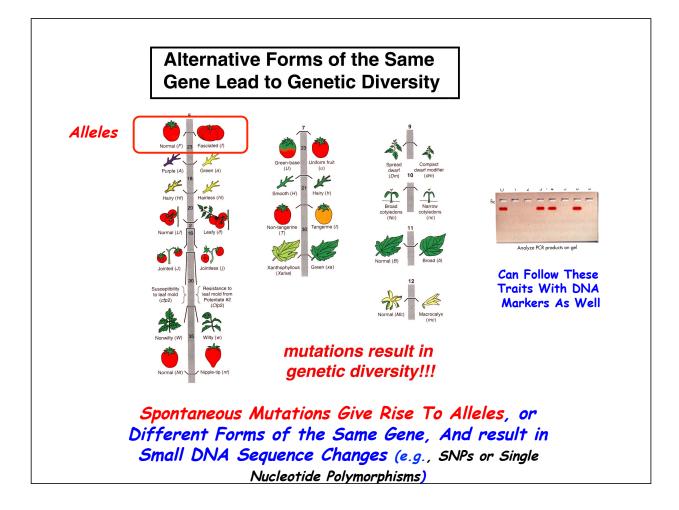
## WHAT ARE THE PROPERTIES OF A GENE?

- 1. Replication
- 2. Stability (Mutations)
- 3. Universalitya) All Cellsb) All Organisms
- 4. Direct Cell Function/ Phenotype





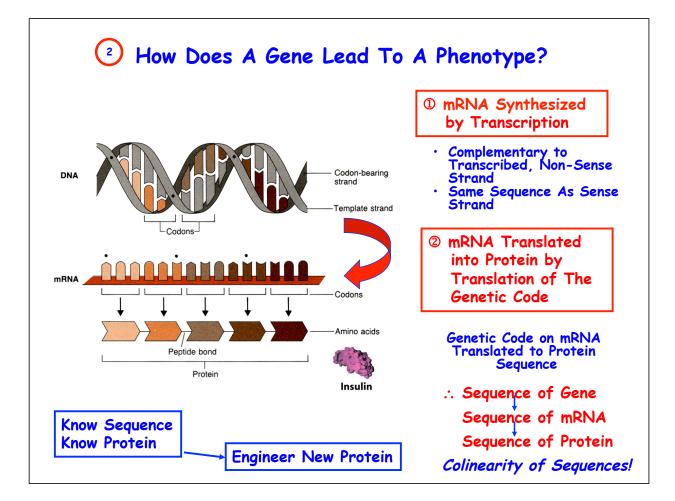


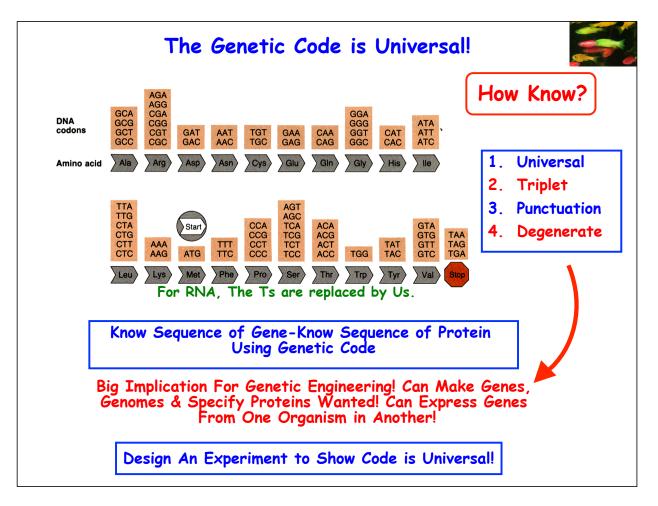




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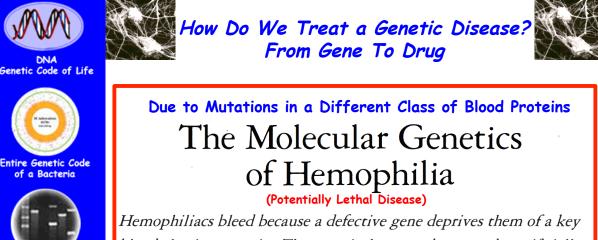


# Expression of Jellyfish Green Fluorescence Protein (GFP) in Pigs Shows That Genetic Code is Universal!!

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blood-clotting protein. The protein has now been made artificially by isolating the normal gene and then inserting it into cultured cells

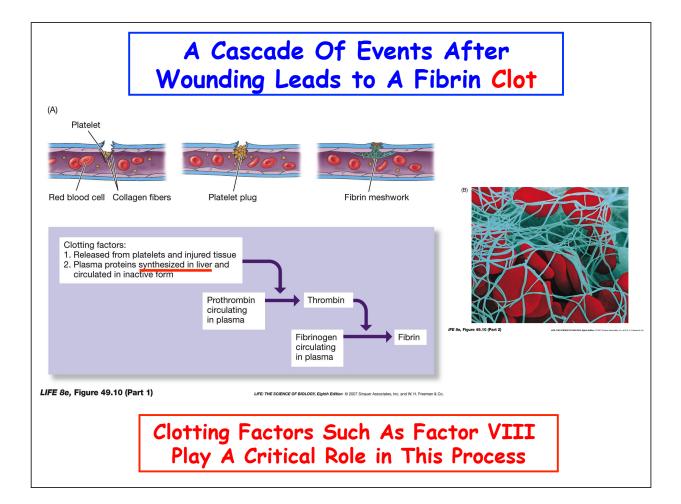
by Richard M. Lawn and Gordon A. Vehar

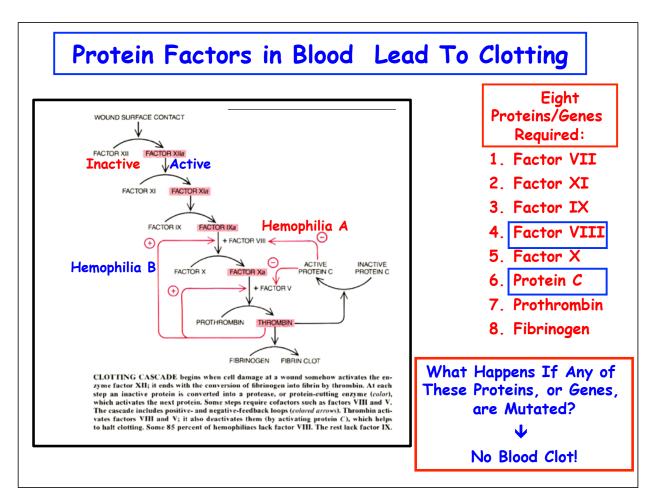


**DNA** Fingerprinting

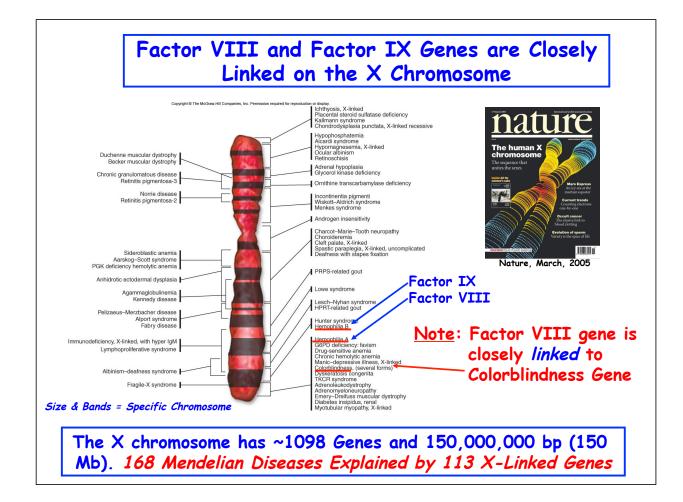
A Case Study of Cloning Genes and mRNAs Reference: Scientific American, March 1, 1986 (Pick Up After Class)

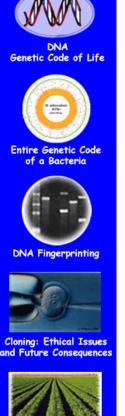






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TABLE 13.2	Some Important	Genetic Disorders			
Disorder	Symptom	Defect	Dominant/ Recessive	Frequency Among Human Births	
Cystic fibrosis	Mucus clogs lungs, liver, and pancreas	Failure of chloride ion transport mechanism	Recessive	1/2500 (Caucasians)	
Sickle cell anemia	Blood circulation is poor	Abnormal hemoglobin molecules	Recessive	1/600 (African Americans	
Tay–Sachs disease	Central nervous system deteriorates in infancy	Defective enzyme (hexosaminidase A)	Recessive	1/3500 (Ashkenazi Jews)	
Phenylketonuria	Brain fails to develop in infancy	Defective enzyme (phenylalanine hydroxylase)	Recessive	1/12,000	
Hemophilia	Blood fails to clot	Defective blood-clotting factor VIII	X-linked recessive	1/10,000 (Caucasian ma	
Huntington disease	Brain tissue gradually deteriorates in middle age	Production of an inhibitor of brain cell metabolism	Dominant	1/24,000	
Muscular dystrophy (Duchenne)	Muscles waste away	Degradation of myelin coating of nerves stimulating muscles	X-linked recessive	1/3700 (males)	
Hypercholesterolemia	Excessive cholesterol levels in blood lead to heart disease	Abnormal form of cholesterol cell surface receptor	Dominant	1/500	
Hemophilia A Hemophilia B	Prior to 1960s - Defective F	philia & 400 Babies/ Average Life Span actor VIII Gene actor IX Gene	Was 11 Year 1/10,00		

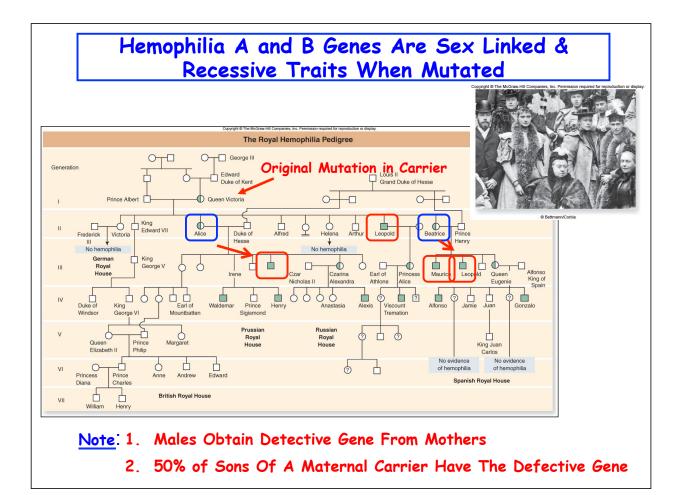


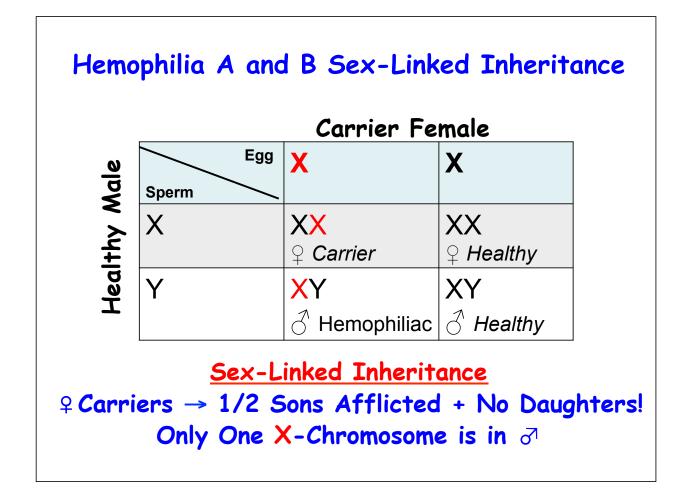


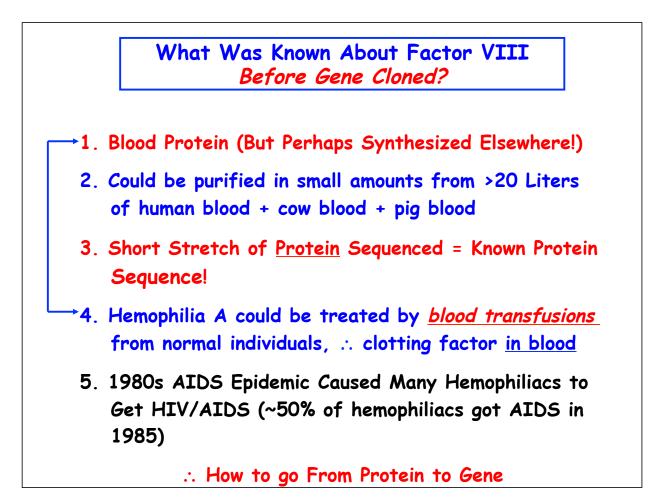


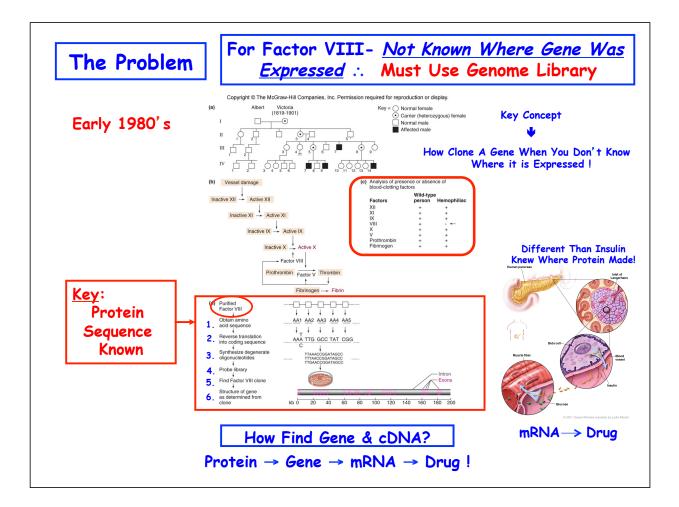
Pedigrees Can Be Used To Determine If a Trait is Dominant or Recessive

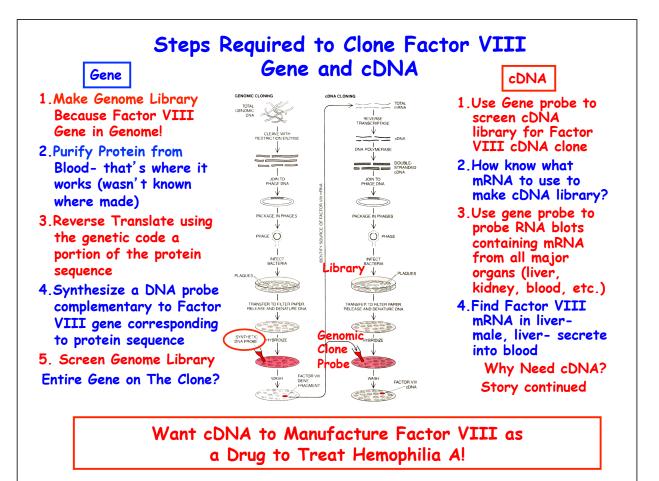
Each Type of Inheritance Predicts Specific Results in Each Generation











Step One

How to Construct a Human Genome Library to Find the Factor VIII Gene?

If It is Not Known Where Gene is Active Can "Look" to Genome Instead of mRNA to Find + Clone Gene!

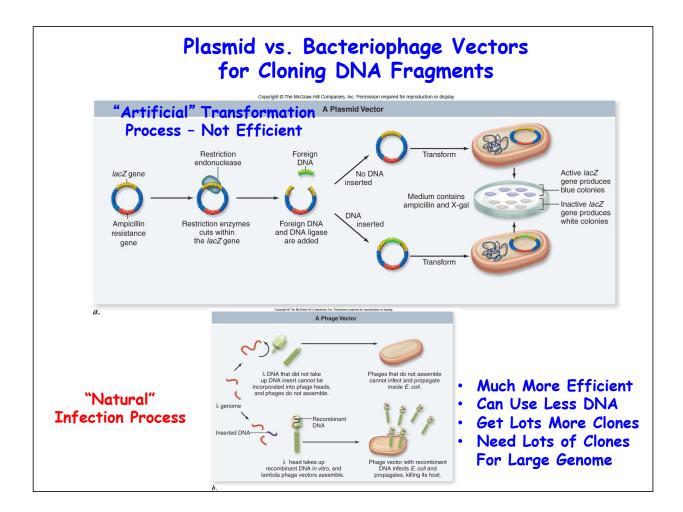


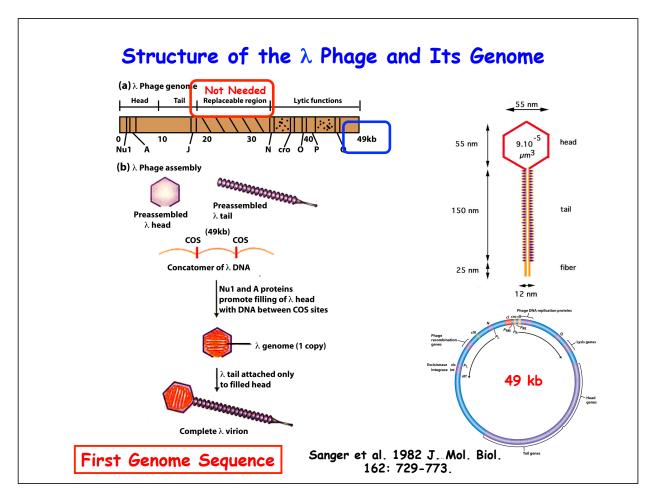
Vectors Used in Genetic Engineering Have Similar Conceptual Properties But are Used in Different Situations

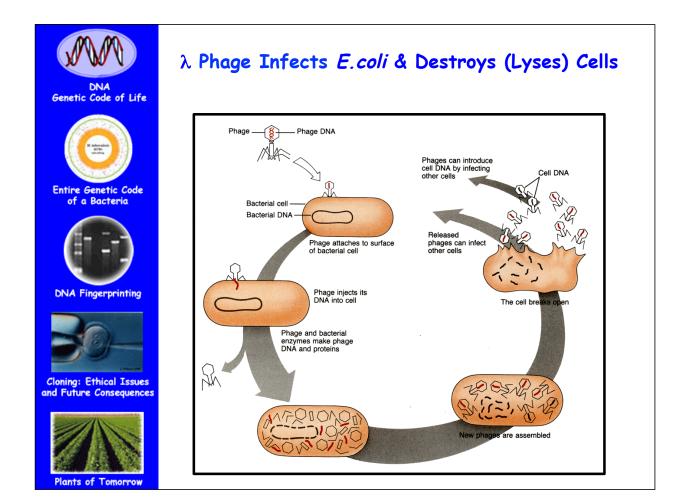
Vector Type	Maximum Insert Size (kb)	Applications	Limitations		
Bacterial plasmid vectors (circular)	₹6-12	DNA cloning, protein expression, subcloning, direct sequencing of insert	Restricted insert size; limited expression of proteins; copy number problems; replication restricted to bacteria		
	DNA	DNA	restricted to batteria		
Bacteriophage vectors (linear)	~25	cDNA, genomic and expression libraries	Packaging limits DNA insert size; host replication problems		
Cosmid (circular)	~35	cDNA and genomic libraries, cloning large DNA fragments	Phage packaging restrictions; not ideal for protein expres- sion; cannot be replicated in mammalian cells		
Bacterial artificial chromosome (BAC, circular)	~300	Genomic libraries, cloning large DNA fragments	Replication restricted to bacteria; cannot be used for protein expression		
Yeast artificial chromosome (YAC, circular)	200–2,000	Genomic libraries, cloning large DNA fragments	Must be grown in yeast; cannot be used in bacteria		
Ti vector (circular)	Varies depending on type of Ti vector used	Gene transfer in plants	Limited to use in plant cells only; number of restriction sites randomly distributed; large size of vector not easily manipulated		

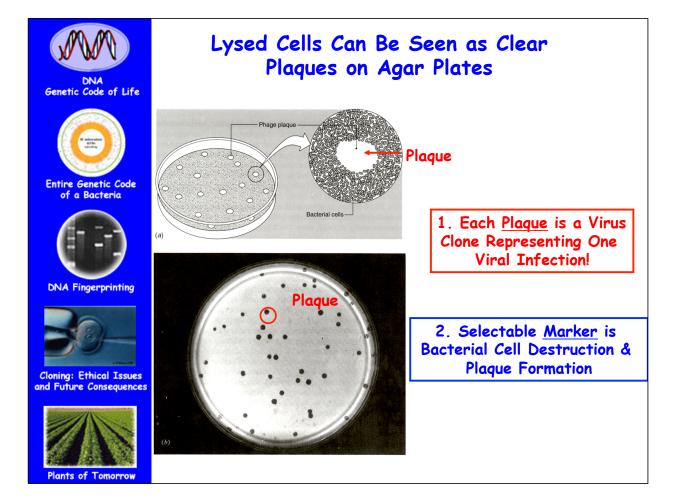
Plasmids vs. Bacteriophage Vectors

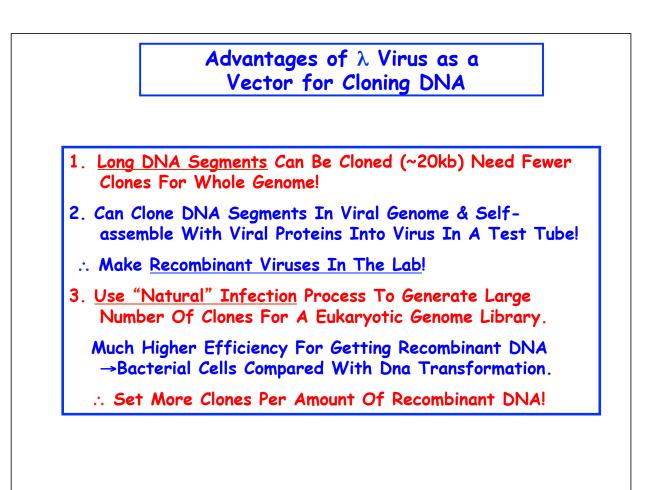
- 2. Selectable
- 3. Can be used to insert foreign genes/restriction sites
- 4. Easily isolated + transferred back to cells

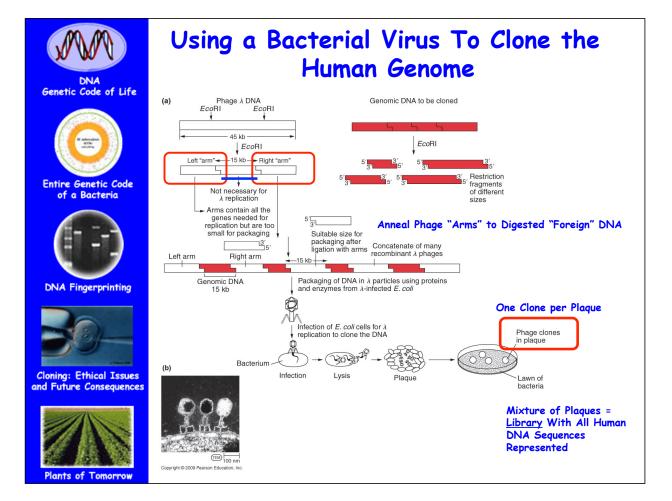


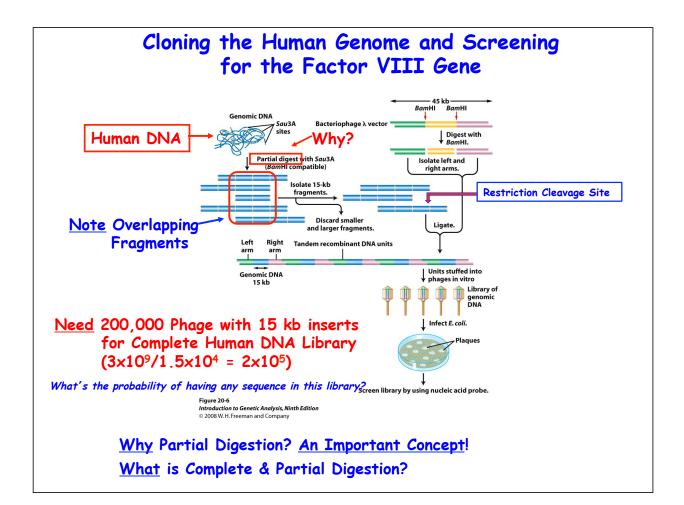


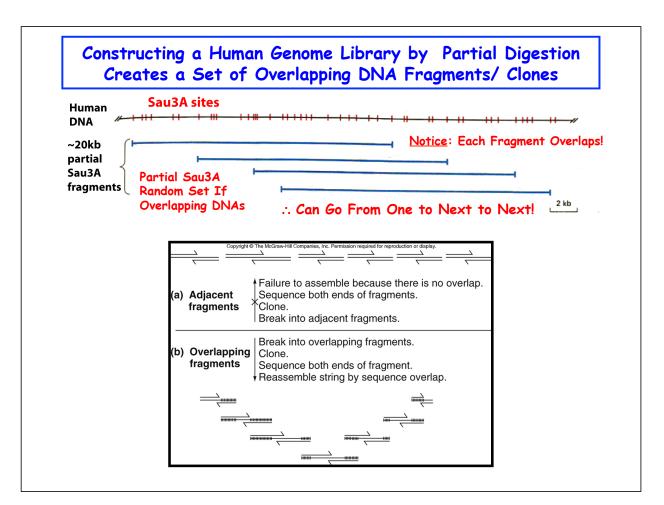








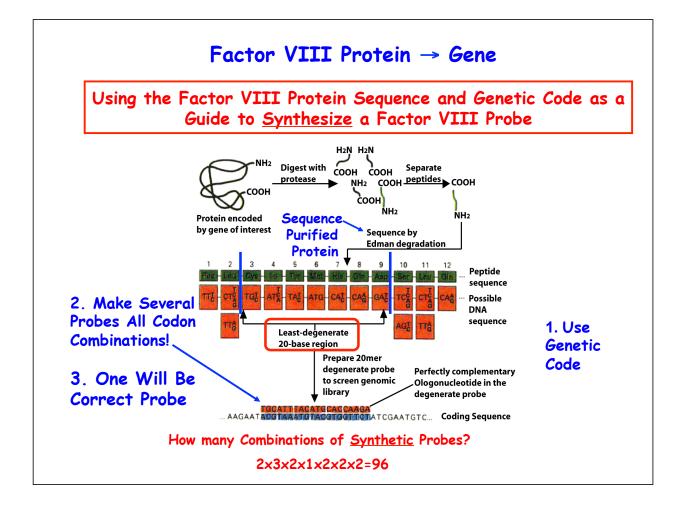


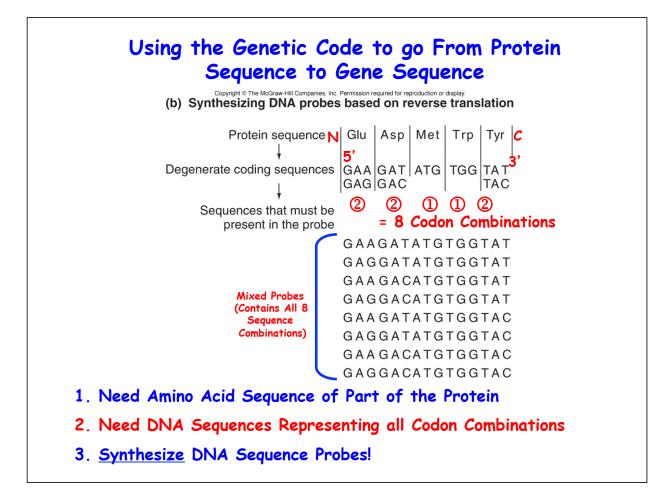


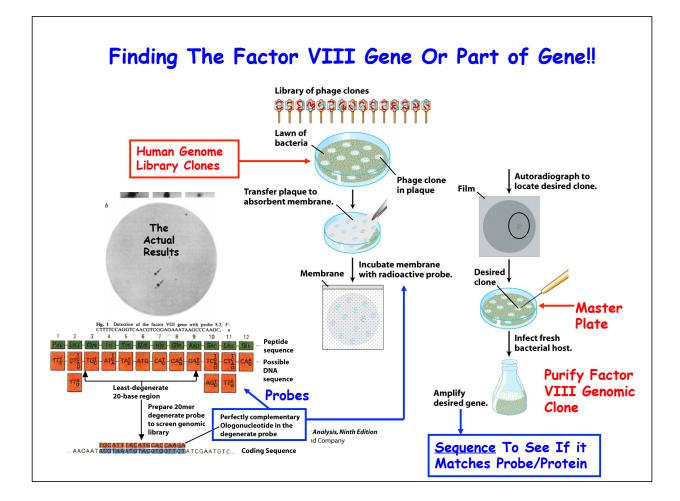
Step Two

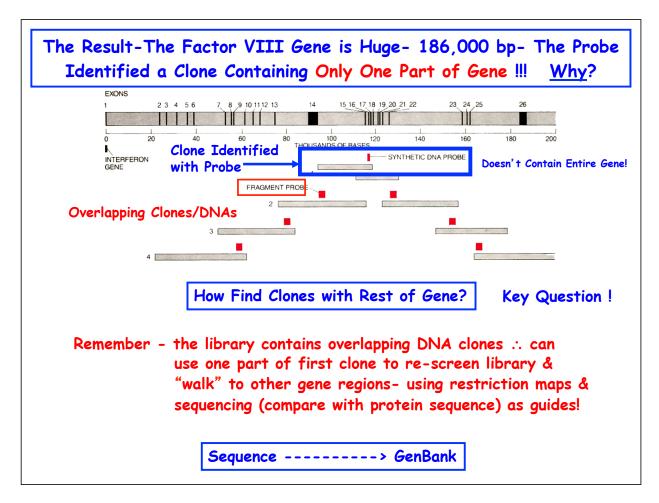
How Find the Factor VIII Gene in a Human Genome Library?

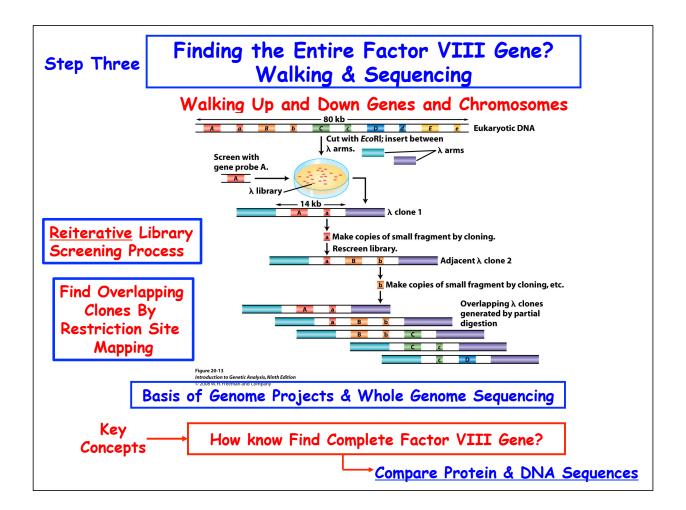
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		U		с		A		G			_
	U	UUU UUC UUA UUG	Phe Leu	UCU UCC UCA UCG	Ser	UAU UAC UAA UAG	Tyr Stop Stop	UGU UGC UGA UGG	Cys Stop Trp	U C A G	
1st	с	CUU CUC CUA CUG	Leu	CCU CCC CCA CCG	Pro	CAU CAC CAA CAG	His Gln	CGU CGC CGA CGG	Arg	U C A G	3rd
letter	A	AUU AUC AUA AUG	lle Start Met	ACU ACC ACA ACG	Thr	AAU AAC AAA AAG	Asn Lys	AGU AGC AGA AGG	Ser Arg	U C A G	letter
	G	GUU GUC GUA GUG	Val	GCU GCC GCA GCG	Ala	GAU GAC GAA GAG	Asp Glu	GGU GGC GGA GGG	Gly	U C A G	
			ſ	<u>Properties</u> • Universal • Three Nucleotides • Punctuation • Degenerate							

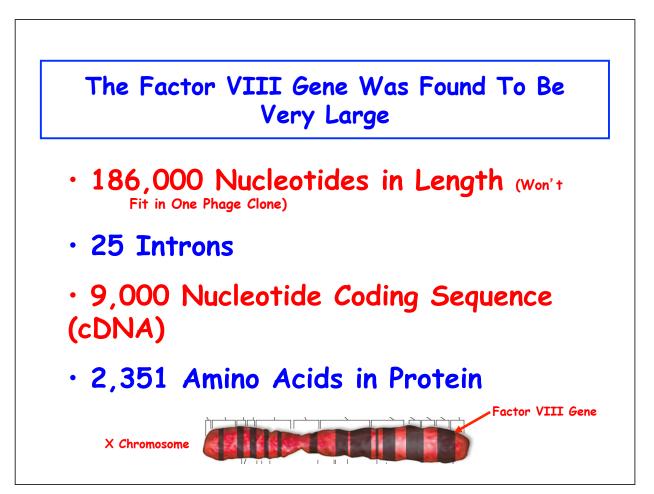








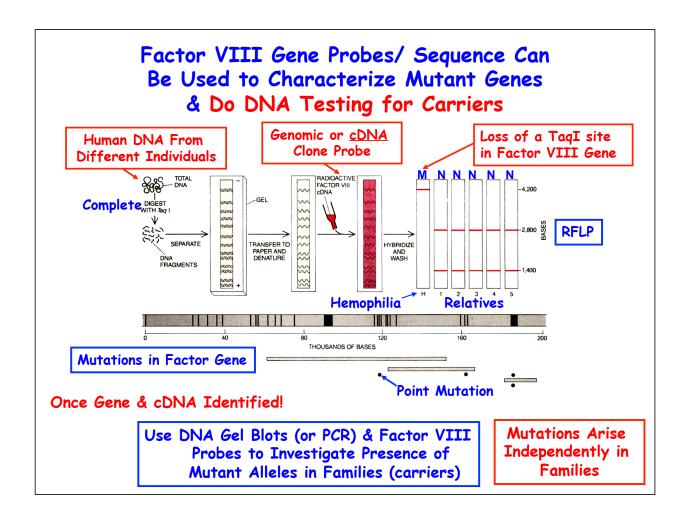


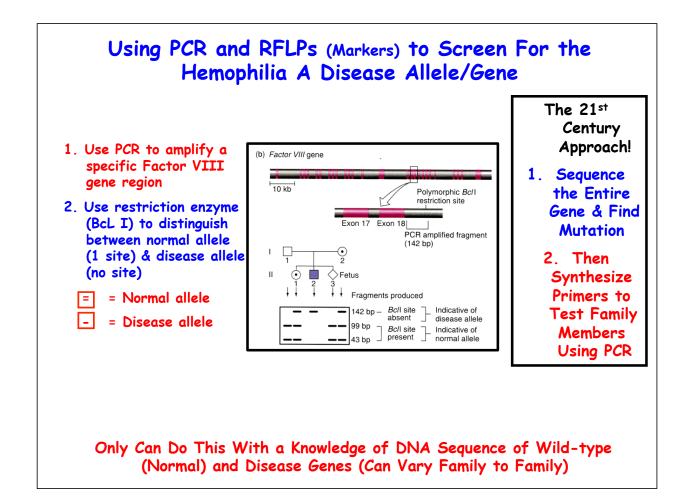


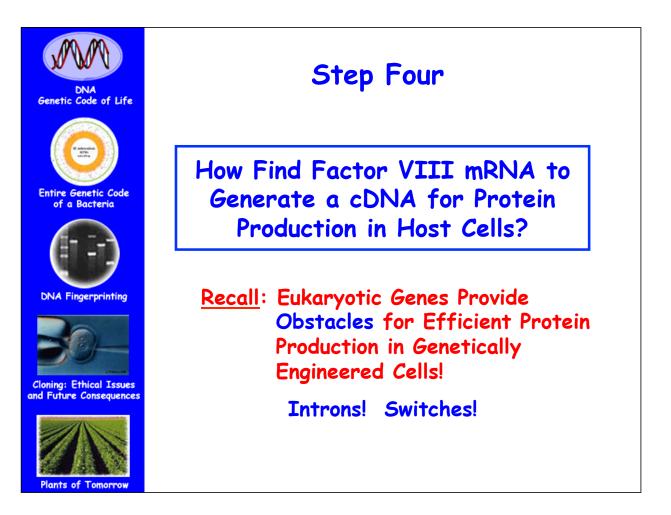
## Factor VIII Mutations Occur Throughout the Gene

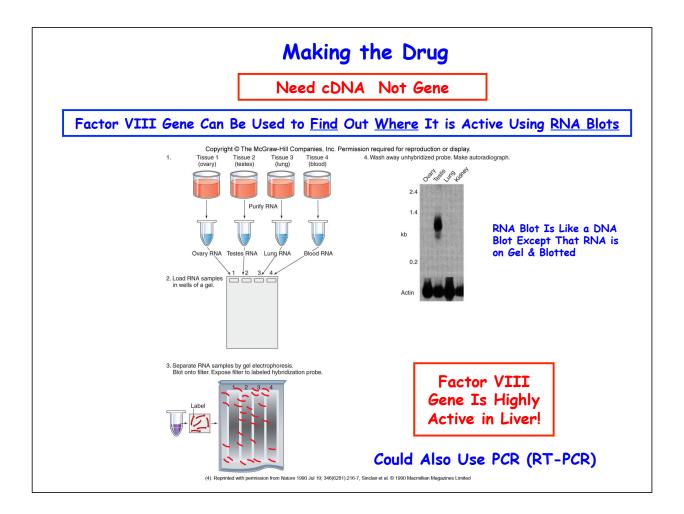
[Haemophilia 11, 481-491 (2005)]

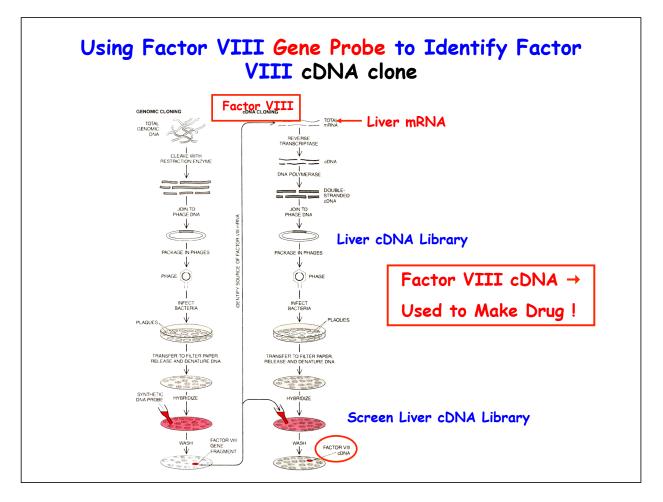
VIII:C (%)	Family history	Consanguinity*	Inversion	Codon <sup>†</sup>	Mutation	Amino acid change	Exon	Conservation <sup>‡</sup>
1	Sporadic	NC	Normal	51	$TTT \rightarrow TCTS$	Phe -> Ser	2	FFFF, identical
1.20	Sporadic	NC	Normal	80	$GTT \rightarrow GAT$	$Val \rightarrow Asp$	3	VVVV, identical
1	Sporadic	NC	Normal	102	$GGT \rightarrow GTT_{3}$	$Gly \rightarrow Val$	3	GGGG, identical
2	Sporadic	NC	Normal	104	$TCC \rightarrow CCCS$	Ser $\rightarrow$ Pro	3	SSSS, identical
6	Sporadic	NC	Normal	143	$GAG \rightarrow AAGS$	$Ghu \rightarrow Lys$	4	EEEE, identical
1	Sporadic	NC	Normal	233	delCA§	Thr $\rightarrow$ fs (TGA-264)	6	
2.70	Inherited	NC	Normal	321	$GAA \rightarrow AAA$	$Glu \rightarrow Lys$	8	EEEE, identical
0	Sporadic	NC	Normal	372	$CGC \rightarrow CAC$	$Arg \rightarrow His$	8	RRRR, identical
3	Inherited	NC	Normal	527	$CGG \rightarrow TGG$	$Arg \rightarrow Trp$	11	RRRR, identical
1	Sporadic	NC	Normal	52.8	$TGC \rightarrow TACS$	Cys → Tyr	11	CCCC, identical
1	Inherited	NC	Normal	592	$CAA \rightarrow TAA$	$Gln \rightarrow Stop$	12	QQQQ, identical
1	Inherited	NC	Normal	864	delGACA	Gly $\rightarrow$ fs [TAA-867]	14	
					insCAATTAAATGAGAA§			
1	Sporadic	NC	Normal	948	insA§	Lys $\rightarrow$ fs (TGA-984)	14	
1	Sporadic	NC	Intron 1	1107	$AGG \rightarrow TGGS$	$Arg \rightarrow Trp$	14	RGKK, dissimilar
1	Sporadic	NC	Normal	1107	$AGG \rightarrow TGG$	$Arg \rightarrow Trp$	14	RGKK, dissimilar
1	Inherited	NC	Normal	1191-1194	delA	Ile $\rightarrow$ fs (TAG-1198)	14	
1.40	Sporadic	NC	Normal	1191-1194	insA	Ile $\rightarrow$ fs (TAA-1220)	14	
1	Sporadic	C	Normal	1227	delC§	Leu $\rightarrow$ fs (TGA-1231)	14	
2.10	Sporadic	NC	Normal	1241	$GAC \rightarrow GAG$	$Asp \rightarrow Glu$	14	DGGE, similar
1	Sporadic	NC	Normal	1392	1392dcl14185	Pro $\rightarrow$ fs (TAG-1446)	14	
1	Incrited	C	Normal	1392	1392del14185	Pro $\rightarrow$ fs (TAG-1446)	14	
1	Sporadic	NC	Normal	1441	insA§		14	
1	Incrited	С	Normal	1441	insA§			
1	Inherited	NC	Normal	1.502	$CAG \rightarrow TAGS$	$Gln \rightarrow Stop$	14	QREQ, dissimilar
1	Inherited	NC	Normal	1504	delGT§	Val $\rightarrow$ fs (TGA-1517)	14	
1	Sporadic	NC	Normal	1535	$TGG \rightarrow TGA$	$Trp \rightarrow Stop$	14	WLWM, dissimilar
hibitor 96 BU								
1	Sporadic	NC	Normal	1571	$TAT \rightarrow TAAS$	$Tyr \rightarrow Stop$	14	Y-YY, dissimilar
1	Sporadic	NC	Normal	1581	$AAA \rightarrow TAAS$	Lys $\rightarrow$ Stop	14	KEKK, dissimilar
0.20	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	
1	Inherited	NC	Normal	1751	$GAA \rightarrow AAAS$	$Glu \rightarrow Lys$	15	EEEE, identical
1	Sporadic	NC	Normal	1775	$TTC \rightarrow TCC$	Phe $\rightarrow$ Pro	16	FFFF, identical
1	Sporadic	NC	Normal	1835	$TGG \rightarrow TGAS$	$Trp \rightarrow Stop$	16	WWWW, identical
7.60	Sporadic	C	Normal	1882	$ATC \rightarrow ATAS$	$Ile \rightarrow Ile$	17	IIII, identical
3	Inherited	С	Normal	1966	$CGA \rightarrow CAA$	Arg → Glu	18	RRRR, identical
1	Sporadic	NC	Normal	1966	$CGA \rightarrow TGA$	$Arg \rightarrow Stop$	18	RRRR, identical

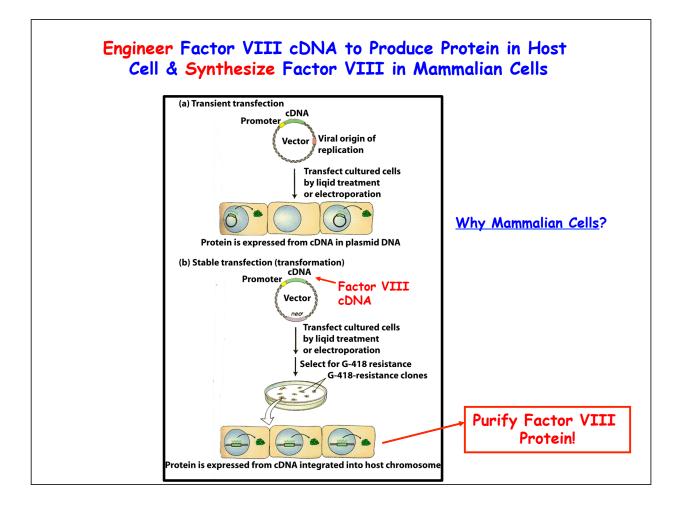


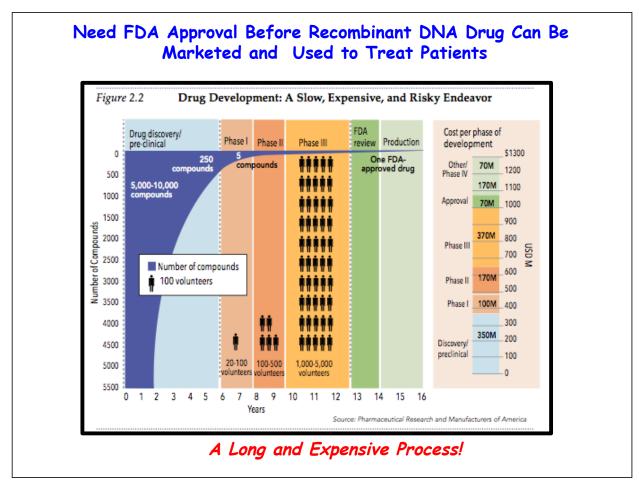


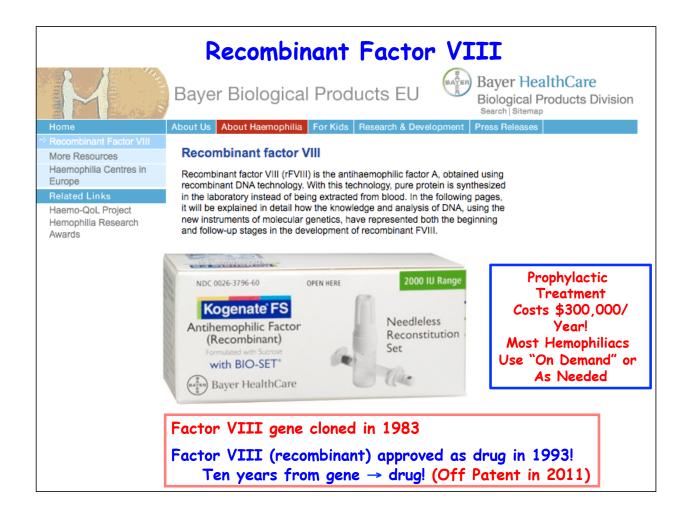


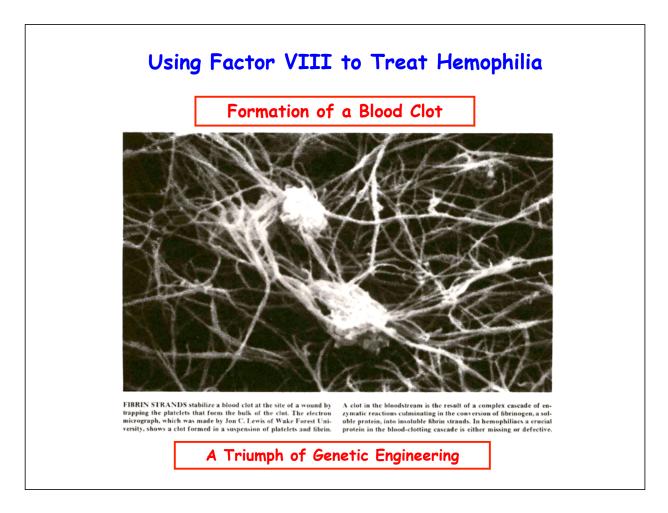












## The Future: Gene Therapy – A Permanent "Cure"

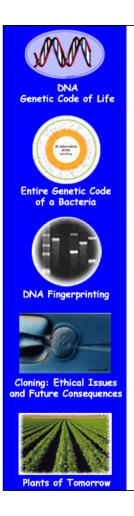
December 10, 2011

# **Treatment for Blood Disease Is Gene Therapy Landmark**

**By NICHOLAS WADE** 

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



#### The Factor VIII Story -- A Summary

- 1. Purify Small Amounts of Factor VIII
- 2. Obtain Partial or Complete Amino Acid Sequence
- 3. Use the Genetic Code to Synthesize Degenerate DNA Probes
- 4. Isolate Factor VIII DNA Clones Complementary to Probe in Genome Library
- Determine if Factor VIII Clones Contain the Complete Gene By 5. Sequencing and Comparing With Protein Sequence
- If Not, "Walk" to Obtain Overlapping DNA Clones That Collectively 6. Contain the Factor VIII Gene
- 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
- 9. Make a cDNA Library From the Target Organ/Tissue and Isolate a Factor VIII cDNA Clone
- 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
- 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)