

Chapter 15.1 – 15.3

Mutations

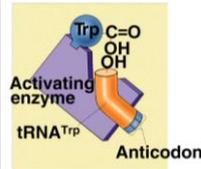
DNA → Transcription → mRNA → Translation → Protein

(b) Sickled red blood cells and the primary structure of sickle-cell hemoglobin

Val | His | Leu | Thr | Pro | Val | Glu | ...
1 2 3 4 5 6 7

Universal Code

- Code is redundant
 - several codons for each amino acid
 - “wobble” in the tRNA
 - “wobble” in the aminoacyl-tRNA synthetase enzyme that loads the tRNA



		Second base				
		U	C	A	G	
U	UUU	Phe	UCU	UAU	UGU	Cys
	UUC		UCC	UAC	UGC	
	UUA	Leu	UCA	UAA	UGA	Stop
C	CUU		CCU	CAU	CGU	
	CUC		CCC	CAC	CGC	
	CUA	Leu	CCA	CAA	CGA	Arg
A	AUU		ACU	AAU	AGU	Ser
	AUA	Ile	ACA	AAC	AGC	
	AUG	Met or start	ACG	AAG	AGG	Arg
G	GUU		GCU	GAU	GGU	
	GUC		GCC	GAC	GGC	
	GUA	Val	GCA	CAA	GGA	Gly
		GUG	GCG	GAG	GGG	

Mutations

Point mutations

- single base change
- base-pair substitution
 - silent mutation
 - no amino acid change
 - redundancy in code
 - missense
 - change amino acid
 - nonsense
 - change to stop codon

Wild type

mRNA: A U G A A G U U U G C U A A

Protein: Met - Lys - Phe - Gly - Stop

Base-pair substitution

No effect on amino acid sequence
U instead of C
A U G A A G U U U G U U A A

Missense
A instead of G
A U G A A G U U U A C U A A

Nonsense
U instead of A
A U G A A G U U U G C U A A

Mutations

Frameshift

- shift in the reading frame
 - changes everything “downstream”
- insertions
 - adding base(s)
- deletions
 - losing base(s)

Wild type

mRNA: A U G A A G U U U G C U A A

Protein: Met - Lys - Phe - Gly - Stop

Base-pair insertion or deletion

Frameshift causing extensive missense
Missing U
A U G A A G U U U G G C U A A ...
Met - Lys - Leu - Ala ...

Frameshift causing immediate nonsense
Extra U
A U G U A A G U U U G C U A A
Met - Stop

Insertion or deletion of 3 nucleotides: no frameshift; extra or missing amino acid
Missing A A G
A U G A A G U U U G C U A A
Met - Phe - Gly - Stop

A Mutation Leads to Sickle Cell Anemia

<p>Wild-type hemoglobin DNA</p> <p>3' C T T 5'</p> <p>mRNA: G A A</p> <p>Normal hemoglobin: Glu</p>	<p>Mutant hemoglobin DNA</p> <p>3' C A T 5'</p> <p>mRNA: G U A</p> <p>Sickle-cell hemoglobin: Val</p>
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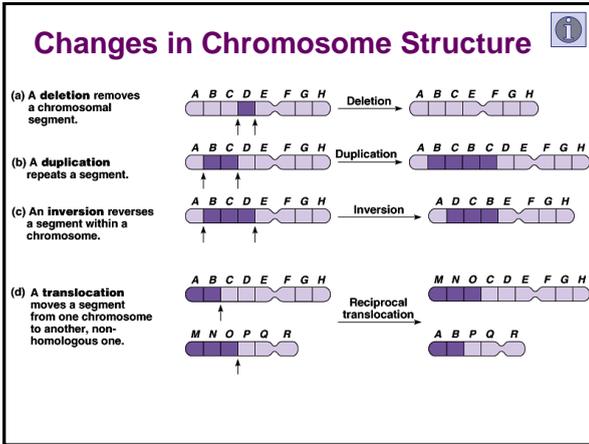
Sickle Cell Anemia

(a) Normal red blood cells and the primary structure of normal hemoglobin

(b) Sickled red blood cells and the primary structure of sickle-cell hemoglobin

Val | His | Leu | Thr | Pro | Glu | Glu | ...
1 2 3 4 5 6 7

Val | His | Leu | Thr | Pro | Val | Glu | ...
1 2 3 4 5 6 7

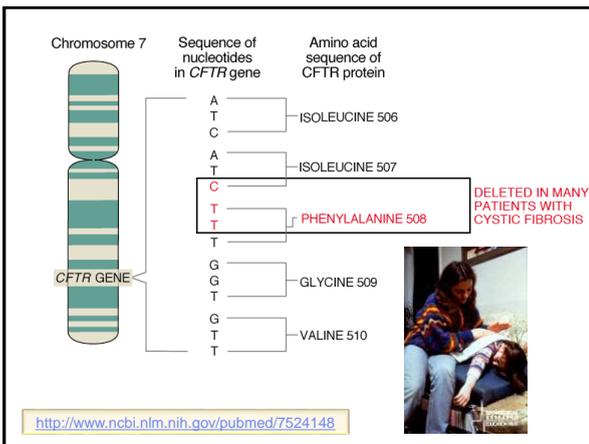
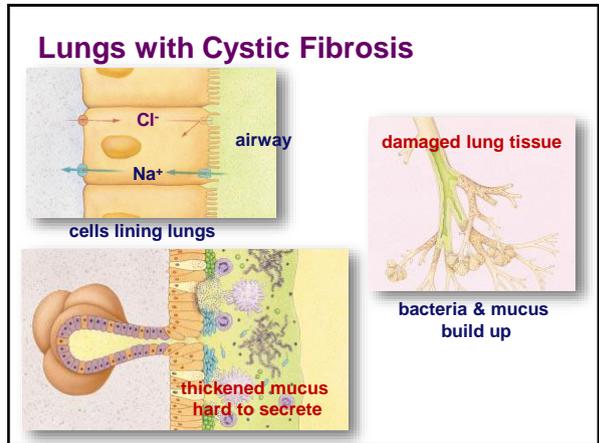
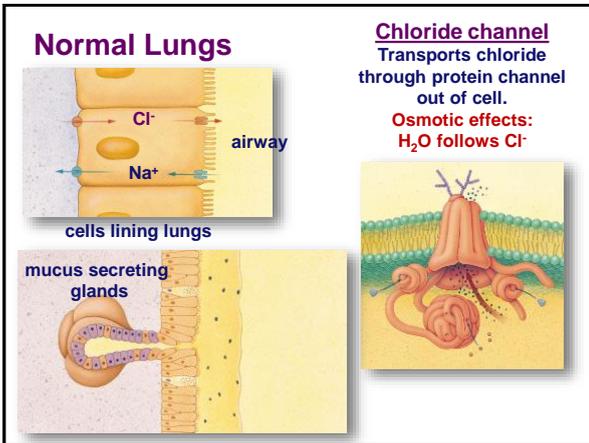


Cystic Fibrosis

■ Primarily Caucasians of European descent

- ◆ strikes 1 in **2500** births
 - 1 in 25 'whites' is a carrier (Aa)
- ◆ normal allele codes for a membrane protein that transports Cl⁻ across cell membrane
 - defective or absent channels fail to transport Cl⁻
 - thicker & stickier mucus coats around cells
 - mucus build-up in the pancreas, lungs, digestive tract & causes bacterial infections
- ◆ without treatment children die before 5; with treatment can live past their late 20s

normal lung tissue



Tay-Sachs

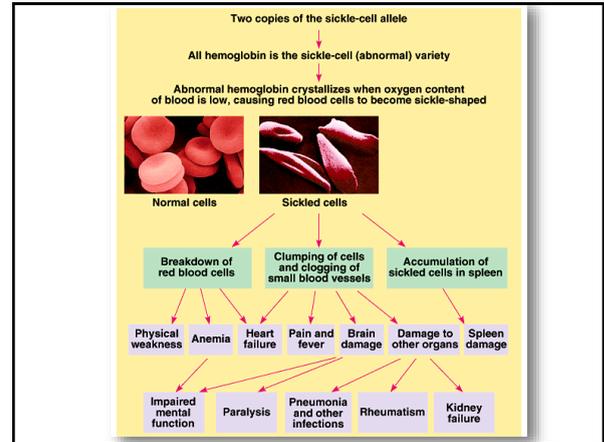
■ Primarily Jews of eastern European (Ashkenazi) descent & Cajuns

- ◆ strikes 1 in **3600** births
 - 100 times greater than incidence among non-Jews or Mediterranean (Sephardic) Jews
- ◆ non-functional enzyme fails to breakdown lipids in brain cells
 - symptoms begin few months after birth
 - seizures, blindness & degeneration of motor & mental performance
 - child dies before 5yo

Sickle Cell Anemia

■ **Primarily Africans**

- ◆ strikes 1 out of **400** African Americans
- ◆ (high frequency in sub-Saharan Africans)
- ◆ caused by substitution of a single amino acid in hemoglobin
- ◆ when oxygen levels are low, sickle-cell hemoglobin crystallizes into long rods
 - deforms red blood cells into sickle shape
 - sickling creates **pleiotropic** effects = cascade of other symptoms



Sickle Cell Anemia

- Substitution of one amino acid for another in a polypeptide chain



Val His Leu Thr Pro Glu Glu ...
1 2 3 4 5 6 7

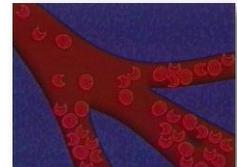
Val His Leu Thr Pro **Val** Glu ...
1 2 3 4 5 6 7

(a) Normal red blood cells and the primary structure of normal hemoglobin

(b) Sickled red blood cells and the primary structure of sickle-cell hemoglobin

Sickle Cell Phenotype

- Often a heterozygous individual
- The 2 alleles are both represented...
 - ◆ both normal & abnormal hemoglobins are synthesized in heterozygote (Aa)
 - ◆ carriers usually healthy, although some suffer some symptoms of sickle-cell disease under blood oxygen stress
 - exercise
 - high altitude...



Heterozygote Advantage

■ **Sickle cell frequency**

- ◆ high frequency of heterozygotes is unusual for allele with severe detrimental effects in homozygotes
 - 1 out of **400** African Americans
- Suggests some selective advantage of being heterozygous
 - ◆ sickle cell: resistance to malaria?
 - ◆ cystic fibrosis: resistance to cholera?

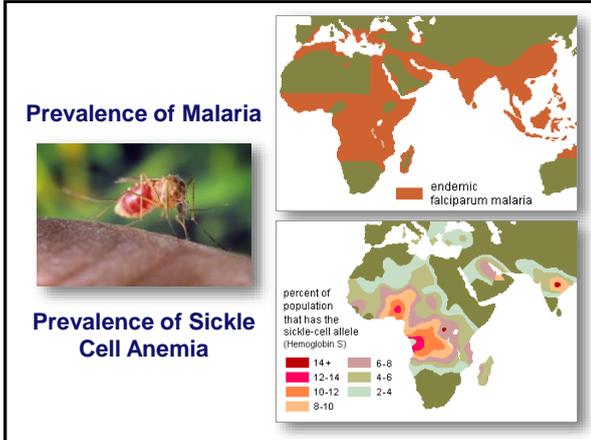


Heterozygote Advantage

■ **Malaria**

- ◆ single-celled eukaryote parasite spends part of its life cycle in red blood cells
- In tropical Africa, where malaria is common:
 - ◆ **homozygous normal** individuals die of malaria
 - ◆ **homozygous recessive** individuals have shorter median life span
 - ◆ **heterozygote carriers** are relatively free of both
- High frequency of sickle cell allele in African Americans is vestige of African roots





Genetics & Culture

- Why do all cultures have a taboo against incest?
 - ◆ laws or taboos forbidding marriages between close relatives are fairly universal
- Fairly unlikely that 2 carriers of same rare harmful recessive allele will meet & mate
 - ◆ but matings between close relatives increase risk
 - consanguineous matings
 - ◆ individuals who share a recent common ancestor are more likely to carry same recessive alleles

- How many genes?**
- Genes
 - ◆ only ~3% of human genome
 - ◆ protein-coding sequences
 - 1% of human genome
 - ◆ non-protein coding genes
 - 2% of human genome
 - ◆ tRNA
 - ◆ ribosomal RNAs
 - ◆ siRNAs
 - ◆ 'junk' DNA as part of the other 97%

- What about the rest of the DNA?**
- Non-coding DNA sequences
 - ◆ regulatory sequences
 - promoters, enhancers
 - terminators
 - ◆ "junk" DNA
 - introns
 - repetitive DNA
 - ◆ centromeres
 - ◆ telomeres
 - ◆ tandem & interspersed repeats
 - transposons & retrotransposons

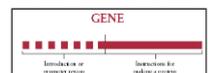
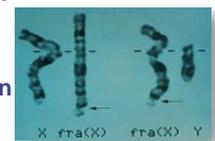
Repetitive DNA

Repetitive DNA & other non-coding sequences account for most of eukaryotic DNA

Table 19.1 Types of Repetitive DNA	
Tandemly Repetitive DNA (Satellite DNA)	
Repeated units at a site are usually identical	
Proportion of mammalian DNA:	10–15%
Length of each repeated unit:	1–10 base pairs
Total length of repetitive DNA per site, in base pairs:	
Regular satellite DNA	100,000–10 million
Minisatellite DNA	100–100,000
Microsatellite DNA	10–100
Interspersed Repetitive DNA	
"Copies" are very similar but not identical	
Proportion of mammalian DNA:	25–40%
Length of each repeated unit:	100–10,000 base pairs
Number of repetitions per genome:	10–1 million

Genetic Disorders of Repeats

- Fragile X syndrome
 - ◆ most common form of inherited mental retardation
 - ◆ defect in X chromosome
 - mutation of FMR1 gene causing many repeats of CGG triplet in promoter region
 - ◆ 200+ copies
 - ◆ normal = 6-40 CGG repeats
 - FMR1 gene not expressed & protein (FMRP) not produced
 - ◆ function of FMR1 protein unknown
 - ◆ binds RNA



Fragile X Syndrome

- The more triplet repeats there are on the X chromosome, the more severely affected the individual will be
 - ◆ mutation causes increased number of repeats (expansion) with each generation

Normal

FMR-1 gene

Premutation

FMR-1 gene

Full Mutation

FMR-1 gene

Huntington's Disease

- Rare autosomal dominant degenerative neurological disease
 - ◆ 1st described in 1872 by Dr. Huntington
 - ◆ most common in white Europeans
 - ◆ 1st symptoms at age 30-50
 - death comes ~12 years after onset
- Mutation on chromosome 4
 - ◆ CAG repeats
 - 40-100+ copies
 - normal = 11-30 CAG repeats
 - CAG codes for glutamine amino acid

Huntington's Disease

- Abnormal (*huntingtin*) protein produced
 - ◆ chain of charged glutamines in protein
 - ◆ bonds tightly to brain protein, HAP-1

Woody Guthrie

Interspersed Repetitive DNA

- Repetitive DNA is spread throughout genome
 - ◆ interspersed repetitive DNA (SINES Short **IN**terspersed **E**lements) make up 25-40% of mammalian genome
 - ◆ in humans, at least 5% of genome is made of a family of similar sequences called, *Alu* elements (PV92 anyone?!)
 - 300 bases long
 - *Alu* is an example of a "jumping gene" called a transposon; a DNA sequence that "reproduces" by copying itself & inserting into new chromosome locations

Restriction Enzymes

- Action of enzyme
 - ◆ cut DNA at specific sequences
 - restriction site
 - ◆ symmetrical "palindrome"
 - ◆ produces "ends"
 - sticky ends
 - blunt ends
- Many different enzymes
 - ◆ named after organism they are found in
 - EcoRI, HindIII, BamHI, SmaI

Madam I'm Adam

CTGAATTCGG
GACTTAAGGC

CTG | AATTC GG
GACTTAA | GGC

Discovery of Restriction Enzymes

1960s | 1978

Werner Arber Daniel Nathans Hamilton O. Smith

Restriction enzymes are named for the organism they come from:

EcoRI = 1st restriction enzyme found in *E. coli*

Gel Electrophoresis

- Separation of DNA fragments by size
 - ◆ DNA is negatively charged
 - moves toward + charge in electrical field
 - ◆ agarose gel
 - “swimming through Jello”
 - smaller fragments move faster

cut DNA 1st with restriction enzymes

Gel Electrophoresis

Gel Electrophoresis

Measuring Fragment Size

- compare bands to a known “standard”
 - ◆ usually lambda phage virus cut with HindIII
 - nice range of sizes with a distinct pattern

RFLP

- Restriction Fragment Length Polymorphism
 - ◆ differences in DNA between individuals

◆ if change in DNA sequence affects restriction enzyme “cut” site...
 ◆ ...it will create a different band pattern

Polymorphisms in Populations

- Differences between individuals at the DNA level (SNPs vs. STRs)

RFLP Use in Forensics

- DNA evidence

The diagram illustrates the RFLP process in three steps: 1. DNA from a crime scene, Suspect 1, and Suspect 2 is digested with enzymes to create fragments. 2. The fragments are separated by gel electrophoresis. 3. The resulting banding patterns are compared to identify the source of the DNA.

RFLP Use in Forensics

- 1st case successfully using DNA evidence
 - ◆ 1987 rape case convicting Tommie Lee Andrews

The top image shows a comparison of DNA profiles. The 'semen sample from rapist' (blue arrow) and 'blood sample from suspect' (red arrow) both show a unique banding pattern that matches the 'Victim Rapist's semen' and 'Suspect's blood' profiles. The 'standard' lanes provide a reference for the expected banding patterns.

The bottom image shows a similar comparison, with the 'semen sample from rapist' (blue arrow) and 'blood sample from suspect' (red arrow) matching the 'Victim Rapist's semen' and 'Suspect's blood' profiles.

RFLP Use in Forensics

- Evidence from murder trial
 - ◆ Do you think suspect is guilty?

The gel electrophoresis image shows the following lanes from top to bottom: blood sample 1 from crime scene, blood sample 2 from crime scene, blood sample 3 from crime scene, blood sample from suspect, blood sample from victim 1, and blood sample from victim 2. Standards are shown on the left and right. Red circles highlight bands in the crime scene and suspect lanes that match the victim 1 and victim 2 profiles. Blue ovals highlight bands in the victim 1 and victim 2 lanes that match the suspect profile.