

DNA, Blood, & Toxicology Unit

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1. Understanding deoxyribonucleic acid (DNA)

↳ molecules carrying a body's genetic information. Shaped in a double-helix

A. Human genome (DNA) - 1st completely mapped in 1985.

- Alec Jeffries & colleagues @ Leicester University in England

- realized that portions of DNA structure contain genes that are unique to each individual

↳ only identical twins have the same DNA

↳ chimera - someone who absorbed their twin (not always identical) in utero & have 2 sets of dna. in one body.

- dna fingerprinting (typing/profiling) - technique for isolating & reading those individual dna markers.

- dna can be obtained from biological evidence: blood, semen, hair (follicular tag), tissue, & bone.

B. What is dna?

(1) - human body has 60 trillion cells containing chromosomes - threadlike structures in the cell nucleus containing genes.

- genes - arranged on the thread like beads. Each chromosome has nearly 85,000 genes

- fundamental unit of heredity - instructs body cells to make proteins that determine everything from hair color to susceptibility to diseases

- each gene is composed of dna designed to carry out a single body function

(2) structure of dna

- discovered by James Watson & Francis Crick in the 1950s.

- dna is a Polymer - molecule made of a large # of atoms, usually arranged in repeating units called monomers. In dna, the monomers are called nucleotides.

can be millions of bases long

↳ repeating dna unit made of 4 bases attached to a phosphate-sugar group.

see Pic 1

- adenine
- guanine

~~polys~~

- cytosine
- thymine

(3) complementary base pairing

A & T go together as do C & G on the double helix to make the ladder



the number of different sequences combining is staggering since average chromosome contains 1 million base pairs human has dna that

- all the human chromosomes taken together contain ~3 billion base pairs.

(4) dna at work

- Inheritable traits are controlled by dna's ability to direct the production of proteins - made by linking a combo of amino acids (~20)

pic 2

Ex) hemoglobin protein - (found in red blood cells, which carry O₂ to our cells & carry away CO₂, is made of 4 amino acid chains

Normal	valine	- histidine	- leucine	- threonine	- proline	- glutamate	- glutamate
Sickle cell	↓	↓	↓	↓	↓	- valine	- glutamate

- the genetic info that determines the amino acid sequence for every protein manufactured in the body is stored in dna. (the A, T, G, & C)

- the key for deciphering the genetic code for each amino acid is a sequence of 3 of those bases

- & alanine - C-G-T
- aspartate - C-T-A
- phenylalanine - A-A-A
- glutamate - G-A-G

so looking at a dna segment

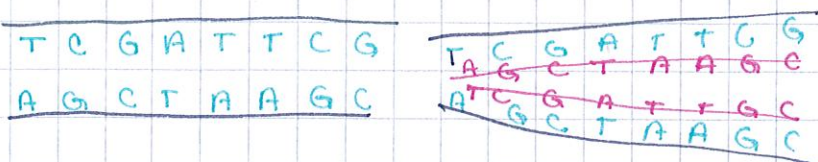
(C-G-T) (C-T-A) (A-A-A) (C-G-T)
 alanine aspartate phenylalanine alanine

2. Replication of DNA

A. The Process of Replication (copying)

(1) DNA replication - the synthesis of new DNA from existing DNA

- each strand is exposed to a collection of free nucleotides.
- Letter by letter, the double-helix is re-created as nucleotides are assembled in the proper order



See Pic 3

B. Polymerase Chain Reaction (PCR) - Lab technique for replicating (copying) a DNA strand outside the human body.

(1) small quantities of DNA found at a crime scene can be replicated for testing so that the sample is not lost by the testing.

(2) done by the aid of a DNA polymerase in a DNA thermal cycle.

a. each cycle doubles the amount of DNA. In a few hours 30 cycles can multiply DNA by a billion.

b. now DNA sample size is no longer a limitation in characterizing DNA from a crime scene.

c. process

- restriction enzymes cut DNA molecules from 2 different sources (more than 20 commercially available)
- insert DNA into a foreign DNA strand (usually bacterium) the bacteria multiply rapidly, passing on altered copies to descendants

3. DNA Typing

30% of human genome is made of repeating segments of DNA.

- acts as filler b/w coding regions of dna

RFLP - restriction fragment length polymorphisms

- typically 15-35 bases long ~~tho~~ repeats many times

1st scientifically accepted protocol for forensics but short lived (1985-90's) (took too long)

see pic p 380

- fragments separated by electrophoresis

- dna placed on a gel-coated plate

- electrical charge \therefore dna migrates across the plate

- smaller fragments move faster \therefore further across the plate

- transferred to a nylon membrane as is on gel, exposed to radioactive particles. X ray film is placed next to membrane \therefore the developed.

used in Clinton-Lewinsky case

PCR - Polymerase Chain Reaction

- used to duplicate a DNA strands millions of times

1st - short sequences of dna identified in pure form (primers)

2nd - add primers to separated dna strands \therefore allow them to combine w/ dna strands primer

G-T-C-T-A-G-C-T-T-C-C-A-G
C-A-G-A-G-T-C-G-A-A-G-G-T-C

G-T-C-T-A-G-C-T-T-C-C-A-G

C-A-G-A

C-C-A-G

C-A-G-A-T-C-G-A-A-G-G-T-C

28-32 cycles are done yielding more than 1B copies

3rd - add dna polymerases \therefore a mix of free nucleotides to separate strands \therefore molecules rebuild themselves.

each cycle takes 2 min.

- Advantages of PCR

- RFLP strands 1000s bases long, PCR only 100's

- shorter strands are more stable, less subject to degradation

- use tiny amounts of original substance

STR - short tandem repeats

- most successful & widely used.
- repeating sequences of 3-7 bases, usually less than 450 bases long, found in great abundance.
- ideal for PCR, overcoming limited sample size
(1,000,000,000 g of less of dna required
(1/50 - 1/100 the amount needed for RFLP))

look @ a common STR known as TH01 - has repeating A-A-T-G
7 variants of TH01 are found in humans

TH01 is extracted & amplified by PCR, then separated by electrophoresis. By examining the distance the STR has migrated, you can determine # of A-A-T-G repeats in the STR.

- every person has 2 STR types for TH01 - one from each parent.

so you might have a semen stain TH01 w/ 6 repeats & 8 repeats

Multiplexing - detecting more than 1 dna marker in a single analysis.

See table
10 p 388

In the US, the forensic sci community has standardized on 13 STRs for entry into CODIS.