



# IDENTIFICATION OF THE LIVING AND THE DEAD

Forensic Medicine Lecture series

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# Lecture outline

- Blood serology
  - Principles of BG inheritance
  - Applications of BG/BG serology in identification
- Identification by DNA profiling
  - Methods of DNA typing
  - DNA technologies used in forensic investigations
  - Disputed parentage
  - DNA forensics databases
  - Ethical, legal & social concerns; limitations

# BLOOD GROUP SEROLOGY.

- Blood groups: Classification of RBC by antigens on RBC membrane
- 41 formally registered blood group systems (ISBT) (e.g. ABO, Lewis, MNSs etc.)
- 45 genes responsible for these systems have been identified and sequenced; associated polymorphisms known
- ~339 BG Ag identified
- A person has the same blood group for life
- Very rarely an individual's BG may change through addition/suppression of an antigen by infection, malignancy, autoimmune disease, BMT/organ transplant

## Applications of BG and BG serology:

- To show whether blood stains (on weapon, clothing etc) are from a particular suspect
- Matching fragmented human remains
- Help resolve parentage/inheritance

# PRINCIPLES OF BG INHERITANCE.

- Autosomal inheritance\* BG genes passed from parents → offspring
- One gene/allelic pair derived from each parent.
- BG antigen in child must be present in at least one parent.
- If parent homozygous for allelic pair (AA, BB) one gene must be present in his/her offspring.

\*Except for XG blood group system (Xg (a); MIC2 gene (CD 99 ) on pseudoautosomal region of X and Y chromosome

## BLOOD GROUP INHERITANCE

Parent 1	Parent 2	Possible offspring	Impossible offspring *
O	O	O	A, B, AB
A	A	O, A	B, AB
A	AB	A, B, AB	O
B	AB	B, A, AB	O
O	AB	A, B	AB, O

# BG applications

- ABO major blood group system
- 80% of individuals secrete water soluble BG in body fluids
- Other BG systems useful – Rh, Lewis etc (Obst./BT)
- Disease setting/associations – eg AIHA
- MN blood group legal paternity cases

# BG application

- Red cell antigens that can be typed:
  - MNS, Rh, ABO, Duffy, Kidd, Kell, Lutheran
- Serum proteins systems that can be typed
  - Haptoglobulins, C3, Gm, Gc, ADA, AK and others
- Red cell enzymes that can be typed:
  - Erythrocyte acid phosphatase, glutamate pyruvate transaminase, phosphoglucomutase and others.



- Using above combined systems one can exclude in 93%. (i.e. the more systems used the better)
- BG systems can exclude a man from parentage but cannot totally confirm paternity
- DNA profiling has now become widely available and is more often used
- Initial serologic testing may be employed where DNA is not available although it has major limitations

# DNA FINGERPRINTING/PROFILING

- Jeffrey Glassberg (American biologist), first patented use of DNA variation for forensics in 1983
- In 1984 Sir Alec Jeffreys (British geneticist) with others used DNA to help solve the rape and murder of two teenagers in Leicestershire in 1983 and 1986.



Sir Alec Jeffreys

# DNA fingerprinting/profiling

## How forensic ID works:

- Identification of DNA sequences that are unique to a species can identify any organism
- DNA is present in nucleated cells
- DNA is extractable from body fluids
- 99.9% of the genome is the same in humans; the remaining 0.1% shows variations between individuals
- These variable DNA sequences - polymorphic markers - can be used to both differentiate and correlate individuals

# DNA forensics cont

- DNA regions that vary from person to person are scanned and used to create a DNA profile of that individual (DNA fingerprint). Usually 13 regions used
- Extremely small chance that another person has the same DNA profile for a particular set of regions
- Sufficient DNA is different to distinguish one individual from another, unless they are **monozygotic twins**

# DNA in identifying persons

- Portions of the DNA sequence that vary the most among humans must be used
- Portions must be large enough to overcome the fact that human mating is not absolutely random
- A match at all would then be rare enough for confidence “beyond reasonable doubt”

# DNA profiling

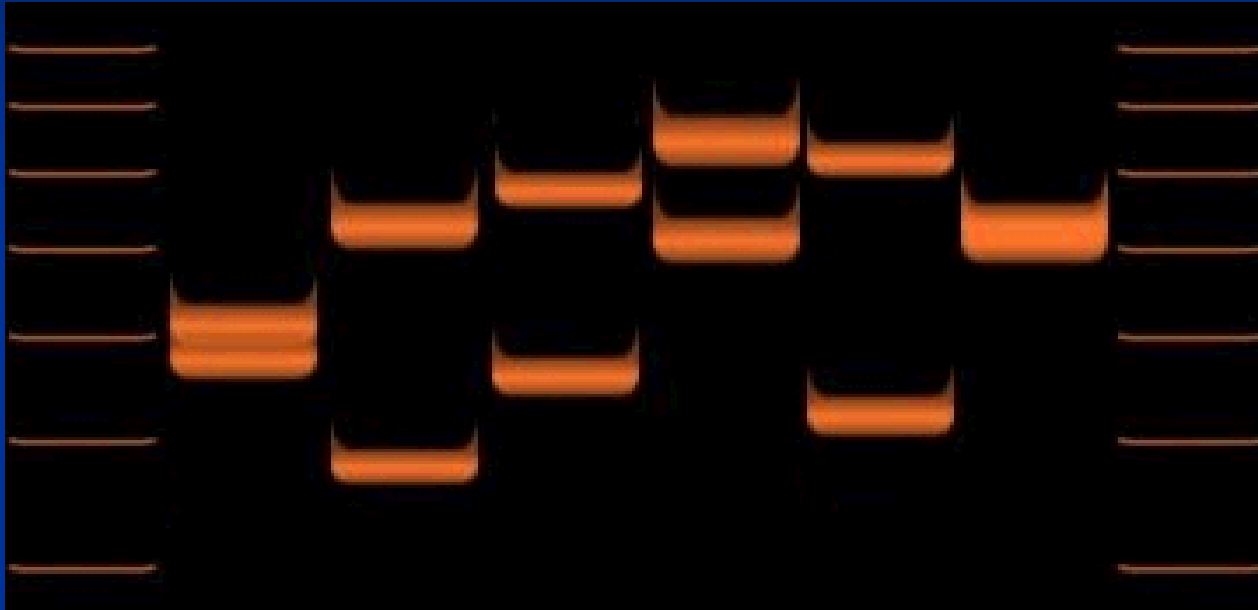
- Only 0.1 % of DNA (about 3 million bases) differs from one person to the next
- 10% DNA molecule used for genetic coding
- These variable regions are used to generate DNA profile of individual
- Chances of 2 unrelated individuals having same sequence 1:1 million billion
- Siblings (not identical twins) 1:10, 000 million

# DNA profiling cont

DNA profiling uses

- Variable number tandem repeats (VNTR)
  - Repetitive sequences that are highly variable,
  - VNTRs loci are very similar between closely related humans, but extremely variable in unrelated individuals
- Short tandem repeats (STR)
  - A type of VNTR also called microsatellite or minisattelites
  - consists of a unit of two to thirteen nucleotides **repeated** hundreds of times in a row on the DNA strand.





Variations of VNTR allele lengths in 6 individuals.

# Technologies used in DNA profiling

## RFLP analysis

(Restriction fragment length polymorphism)

- The first methods for finding out genetics used for DNA profiling involved restriction enzyme digestion, followed by Southern blot analysis.
- Southern blot technique is laborious, and requires large amounts of un-degraded sample DNA
- Method not much used now

# DNA profiling cont.

## PCR analysis

- DNA profiling using PCR analysis gives better discriminating power and the ability to use minute (or degraded) samples.
- PCR greatly amplifies the amounts of a specific region of DNA, using oligonucleotide primers and a thermostable DNA polymerase
- After amplification, DNA Gel electrophoresis is usually performed for analytical purposes
- DNA profiling used today is based on PCR and uses STR

**1. The sample of DNA is placed in a solution containing free nucleotides and the appropriate enzyme**



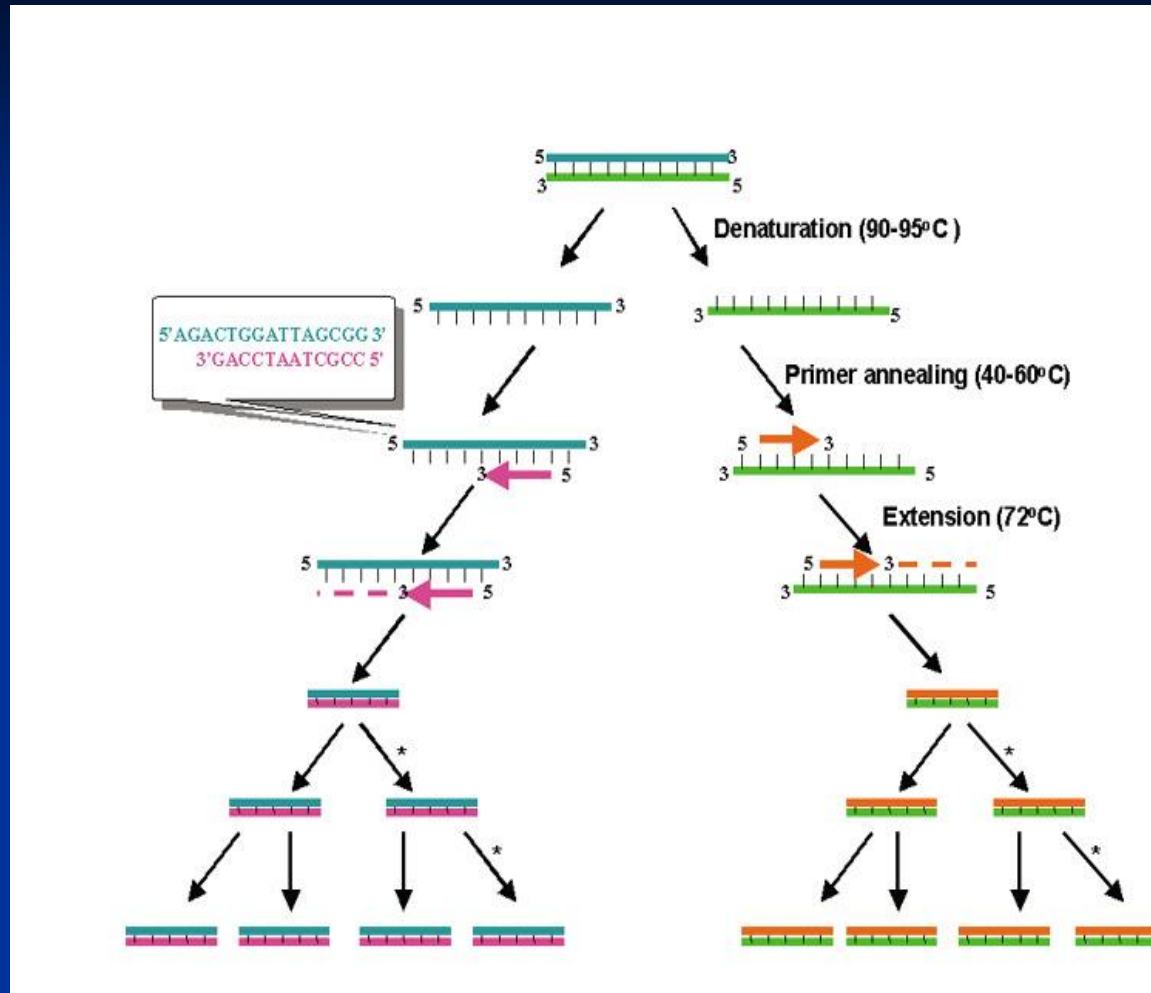
**2. The complementary DNA strands are separated**



**3. New complementary strands are formed using nucleotides from the solution**

**4. The cycle is repeated (usually 20-30 times)**





Copying DNA using polymerases: requires 2 primers, each complimentary to the opposite strand of the denatured DNA

# Polymerase chain Reaction

- PCR – copies small DNA fragments a million fold
- Can be done on very small degraded old samples e.g. licked envelope, stamp, single hair root.
- DNA fingerprinting 1st developed as ID technique 1985
- Useful in – identification of human remains  
comparing crime scene evidence with suspect
- Great care needed to prevent contamination with other biological materials

# Short tandem repeats (STR)

- Typically each STR allele will be shared by around 5 - 20% of individuals.
- Unrelated people have different numbers of repeat units so STRs can be used for discrimination
- STR loci are targeted with sequence-specific primers and amplified using PCR.
- Separation and identification of STRs is done using electrophoresis (capillary electrophoresis instrumentation is the gold standard for human identification)

# Next-generation sequencing

- Crime laboratories can analyze targeted and relevant forensic markers to generate investigative leads and help determine the number of contributors in a mixture analysis
- The application of NGS is particularly helpful with those degraded samples that may not provide a full profile using traditional capillary electrophoresis–based methods



# Short tandem repeat (STR) analysis

- STR used to evaluate specific regions (loci) within nuclear DNA
- High variability in STR regions can be used to distinguish one DNA profile from another
- Use in Criminal investigations
  - FBI uses standard set of 13 specific STR regions for CODIS (Combined DNA Index System)- software program operating in local, state, and national databases of DNA profiles from convicted offenders, unsolved crime scene evidence, and missing persons.

# Amplified fragment length polymorphism

- AmpFLP, started early 1990s.
- Uses VNTR polymorphisms
- Relatively low cost, easily set-up and operation makes it popular in lower income countries.

# IDENTIFICATION BY DNA PROFILING.

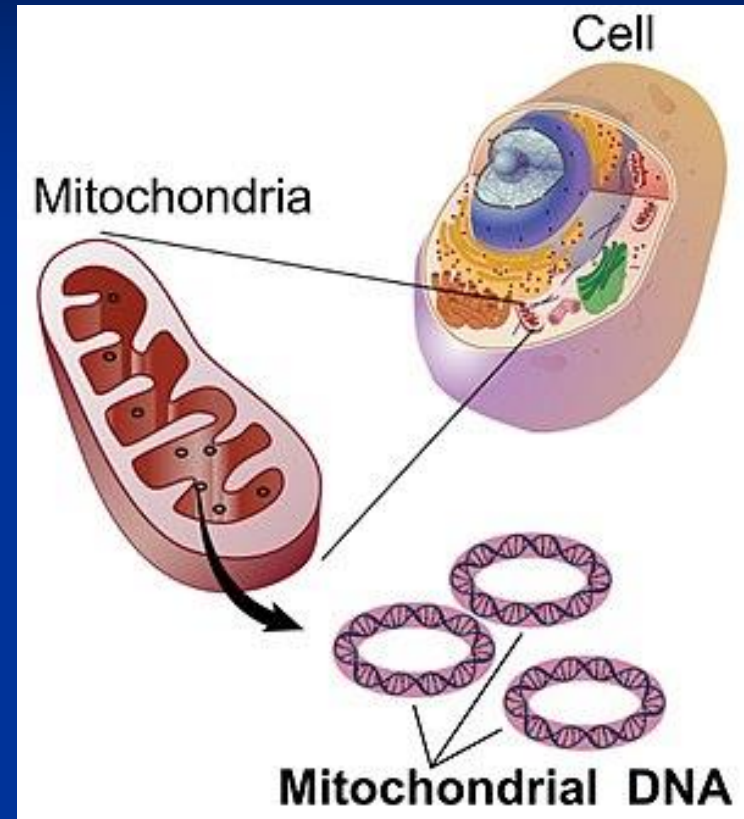
- DNA in hair root bulbs, nucleated hair can be profiled
- DNA from different sources can be matched (i.e. hair bulb/semen/ blood/ saliva etc)
- DNA data base can be created.

# Mitochondrial DNA Analysis (mtDNA)

- Mitochondrial DNA - small circular chromosome found inside mitochondria
- Passed from mother to offspring through egg
- mtDNA – useful in examination of DNA from samples that cannot be analyzed by RFLP or STR
- mtDNA analysis uses DNA extracted from mitochondria
- Older biological samples that lack nucleated cellular material – e.g. hair, bones, and teeth, can be analyzed with mtDNA

## Mitochondrial DNA

- ‘Powerhouse’ of the cell
- Human mtDNA includes 16,569 base pairs and encodes 13 proteins
- mtDNA permits an examination of the relatedness of populations
- important in anthropology, biogeography



- Mothers have the same mitochondrial DNA as their daughters (Embryo's mitochondria come from egg)
- Useful in missing-person investigations.
- mtDNA is extremely useful in the investigation of old unsolved cases

# Y-Chromosome Analysis

- Y chromosome is passed directly from father to son
- Analysis of genetic markers on the Y chromosome useful for:
  - Tracing relationships among males
  - Analyzing biological evidence involving multiple male contributors

# Examples of DNA Uses for Forensic ID

- DNA probes designed that will each seek out and bind to the complementary DNA sequence
- A series of probes create a distinctive pattern
- In criminal cases involves obtaining samples from crime-scene evidence and a suspect, extracting the DNA, and analyzing it for the presence of a set of specific DNA regions (markers)



## DNA ANALYSIS cont...

- Identify potential suspects whose DNA may match evidence left at crime scenes
- Exonerate persons wrongly accused of crimes
- Identify crime and catastrophe victims
- Establish paternity and other family relationships
- Identify endangered and protected species as an aid to wildlife officials (could be used for prosecuting poachers)

## DNA ANALYSIS cont...

- Detect bacteria and other organisms that may pollute air, water, soil, and food
- Match organ donors with recipients in transplant programs
- Determine pedigree for seed or livestock breeds
- Authenticate consumables such as caviar, champagne and wine

## DNA ANALYSIS cont...

- Monitoring bone marrow transplants
- Detecting fetal cells in a mother's blood
- Tracing human history
- Development of newer instrumentation and techniques making DNA profiling easier, faster, cheaper and more widely available

# Some Interesting Uses of DNA Forensic ID

- Identifying the victims of the September 11, 2001
- Various air crashes: Ethiopian Airlines in 2019 etc
- Disappeared Argentina children 1976-1983
- Migration Patterns (genetic anthropology, migrations, human lineage)
- Wine heritage (18 of world's great wines are closely related)
- Poached animals
- Burns victims in various tragedies
- Tracing ones roots – African Americans

# Disputed parentage

- In practice this means disputed paternity as maternity rarely in doubt.
- Exceptions exist: Where claims of swapped or stolen babies (Bishop Deya saga!)
  - A married man claiming wife has committed adultery and disputes being father of child/children
  - A woman with illegitimate offspring may allege that a certain man is the father to obtain an affiliation order

## Paternity testing cont...

- Court may request paternity determination
- Submission to blood testing agreed upon (court orders needed otherwise)
- Child's consent necessary (over 16/18 years)
- DNA profiling is used as more positively discriminating than BG serology
- Blood taken from child and putative father and mother
- No blood transfusion should have been administered within 3 months of sampling

# Disputed parentage analysis

<u>LOCUS</u>	<u>CHILD</u>	<u>FATHER</u>	<u>MATCH</u>
DSS1358	14/16	16/16	YES
VW3	18/18	17/18	YES
FGA	22/23	22/23	YES
AMELOGENIN	X/X	XY	YES
D881179	14/15	14/15	YES

- Use of several loci (depending on race) e.g. 16
- If all match 99.999% probability of paternity
- A calculated probability of  $> 99.9$  is taken as proof of paternity

# CONROVERSIES OF DNA FINGERPRINTING.

- Accuracy
  - DNA segments not complete strands (whole genome sequencing can be done)
  - Standards of the procedure
  - Human error
- Costs
- Misuse
  - Use of DNA information by unauthorized persons for unauthorized purposes



# Ethical, legal & social concerns of DNA data banking

- The primary concern is privacy:
  - Susceptibility to particular diseases, legitimacy of birth, and predispositions to certain behaviors and sexual orientation
  - Potential for genetic discrimination
  - Who is chosen for sampling also is a concern:
    - Police officers provide the state with intimate evidence that could lead to "investigative arrests."
  - Retention of possible innocent persons DNA

# Conclusion

- BG may be used to identify the living and the dead
- Currently identification is by DNA profiling
- Only  $\sim 0.1$  % of DNA differs from one person to the next
- Portions of DNA sequences that vary the most among humans must be used for identification
- DNA profiling uses VNTR and STR
- mtDNA - examination of DNA from samples that cannot be analyzed by RFLP or STR
- DNA technologies are used in various forensic investigations, disputed parentage
- There are ethical, legal & social concerns and limitations in DNA profiling



Questions?