# The Basis of Evidence: Transfer and Persistence

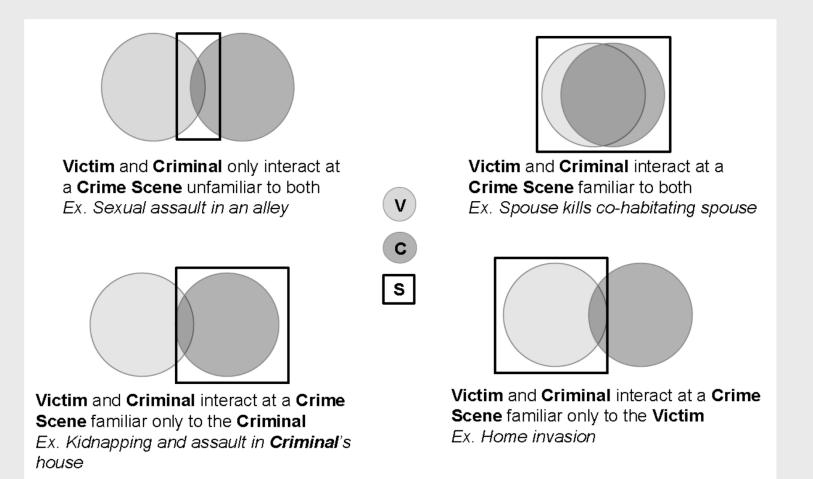
Developed by Edmund Locard, a French forensic microscopist in the early part of the 20th Century, it posits that this exchange of information occurs, even if the results are not identifiable or are too small to be found.

The results of such a transfer would be proxy data: Not the transfer itself, but the remnants of that transaction.



Because forensic science demonstrates associations between people, places, and things through the analysis of proxy data, essentially *all evidence is transfer evidence*.

#### **Relationships and Context**





# Hairs

# Basis for Method

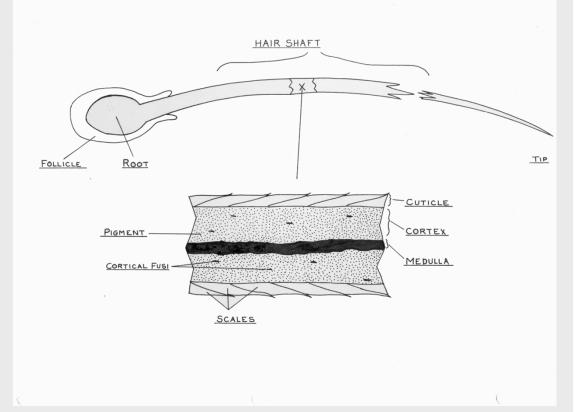
- Anthropology
  - Study of humans
- Zoology/Mammology
  - Study of animals and mammals
- Comparative biology
  - The study of similarities and differences between taxonomic groups
- Microscopy
- Histology

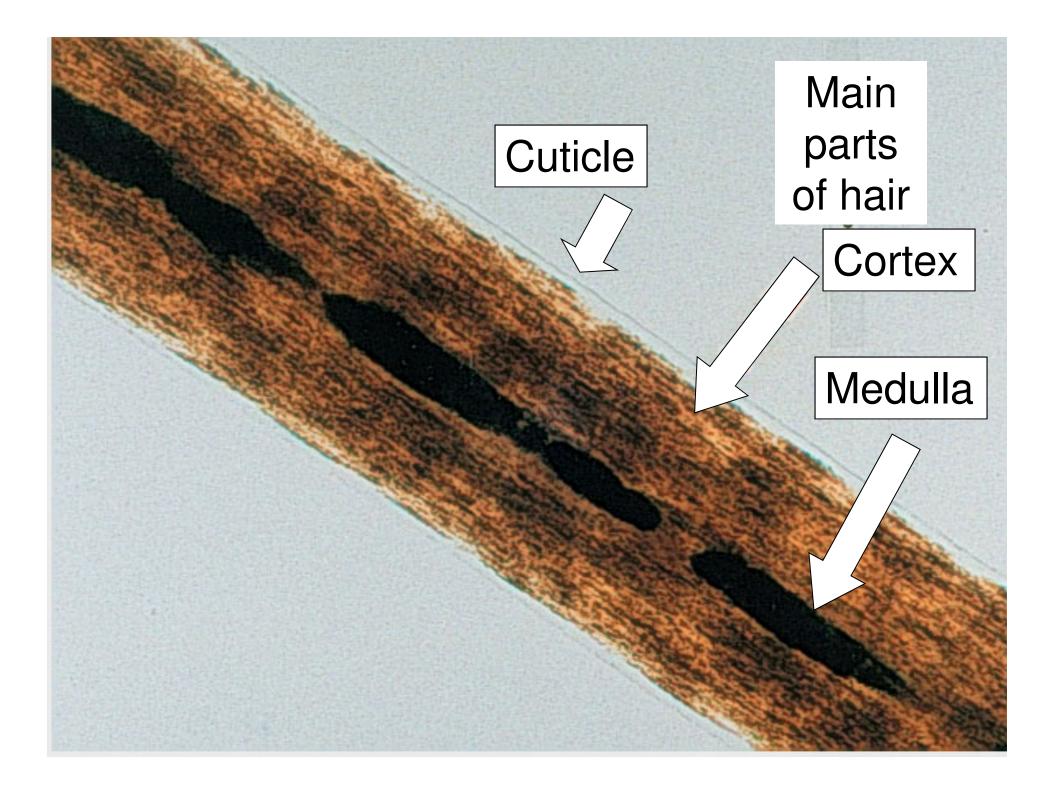
## What can be determined?

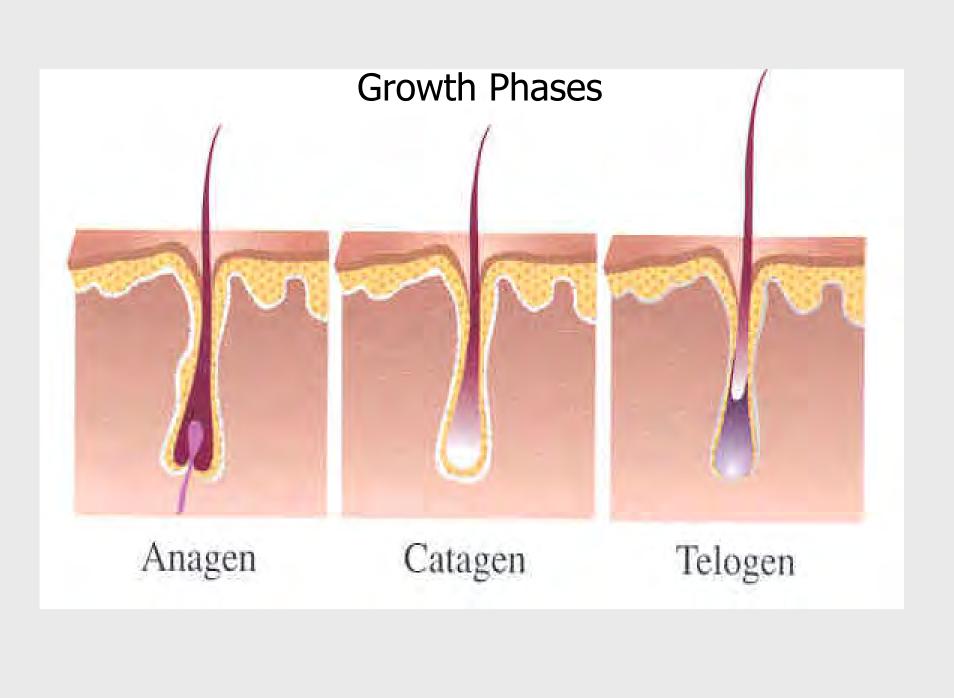
- Is it a hair?
- Is it human?
- What area of the body is it from?
- What is the person's ancestry?
- Is there damage, disease, treatment?
- Is it suitable for comparison?

# Is it a hair?

- Identifiable by its shape and structure
- Parts of a hair
  - Root
  - Shaft
    - Cuticle
    - Medulla
    - Cortex
  - Tip







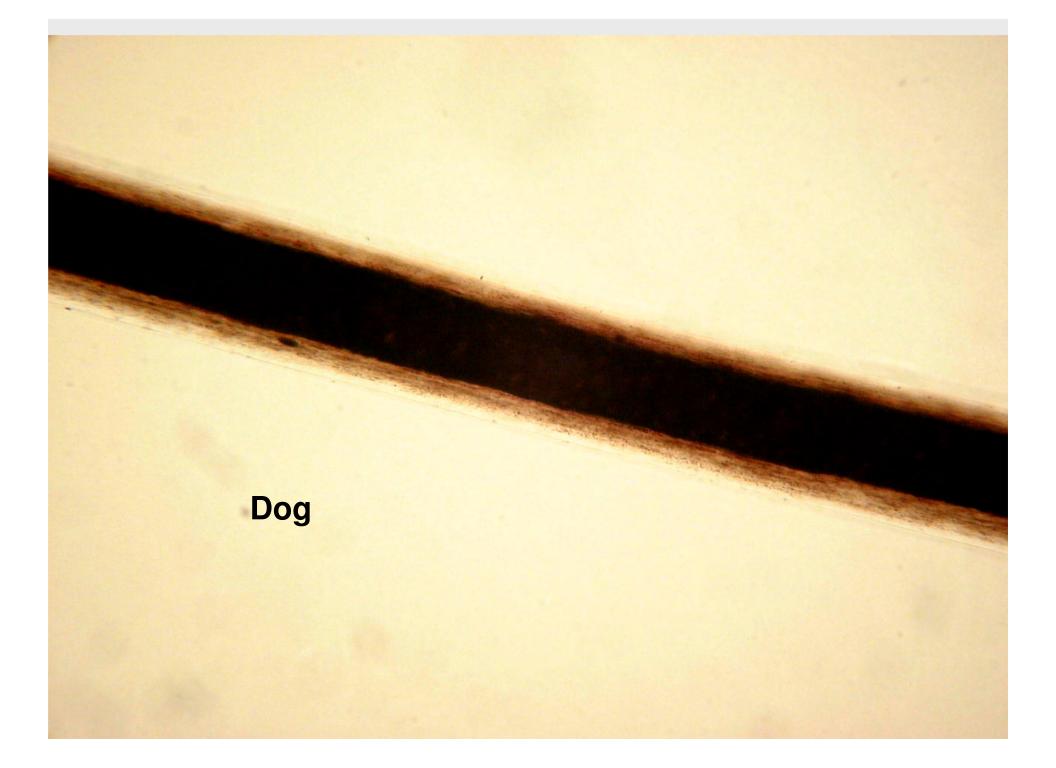
Is it human?

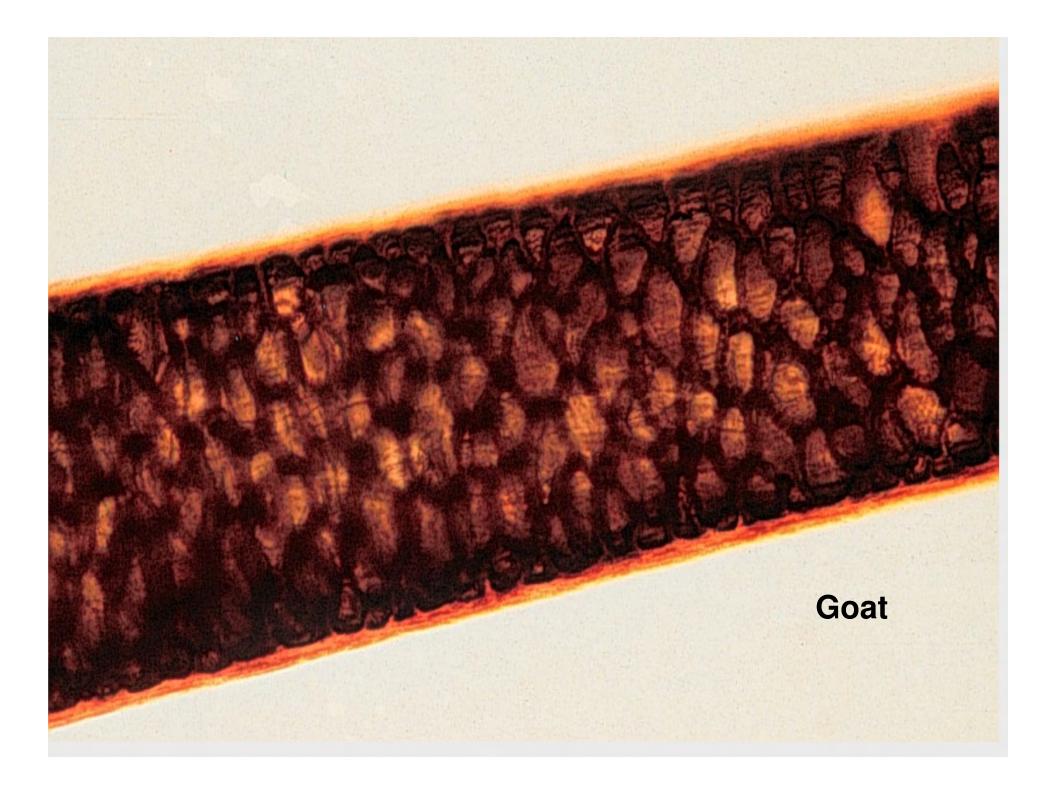
# Human

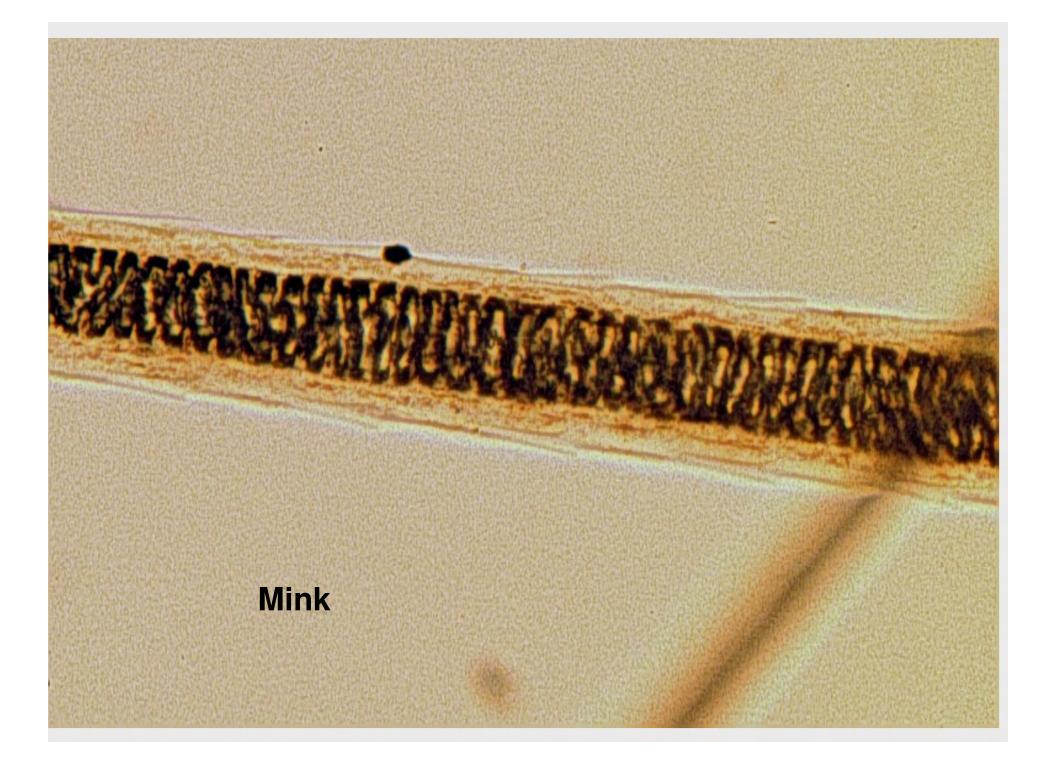
- Even shaft diameter
- Pigmentation even
- Color uniform
- Imbricate scales
- Amorphous medullation
- May be very long

# Animal

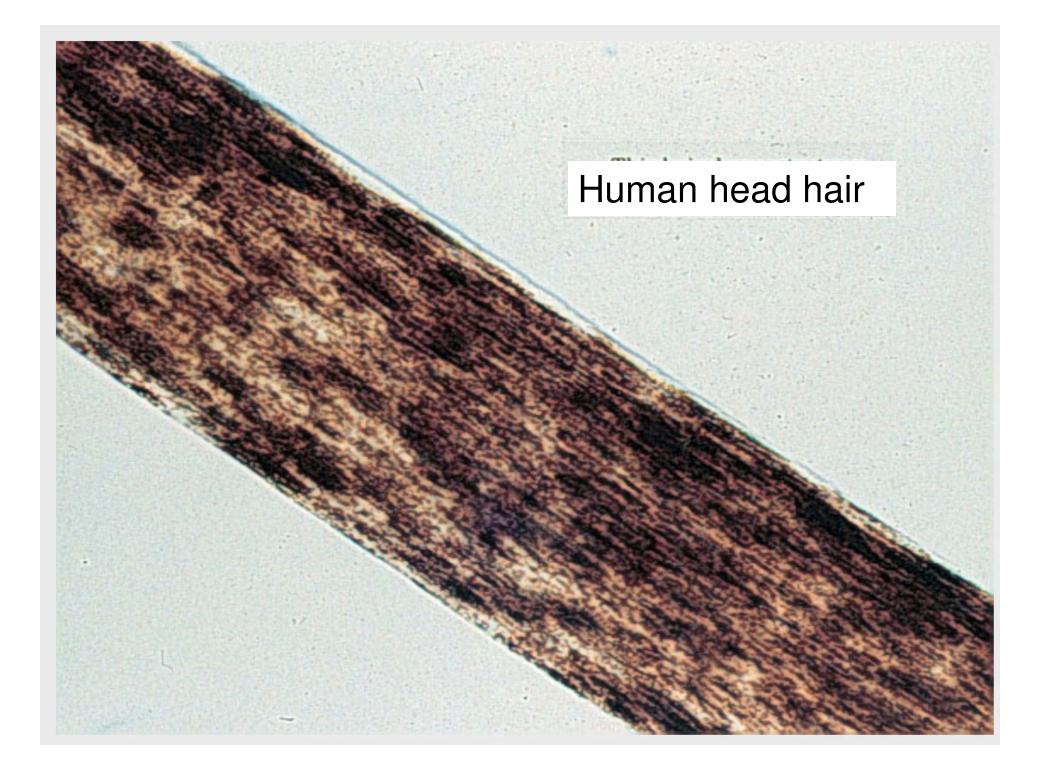
- Shaft varies
- Pigment varies
- Color banding
- Varies by species
- Medullation varies with species
- Most are less than 3"







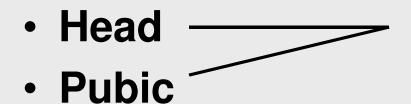




What is the person's ancestry?

- African ancestry
- Asian ancestry
- European ancestry
- Estimation of ancestry based on hair characteristics may not correlate with genetic or self-identity

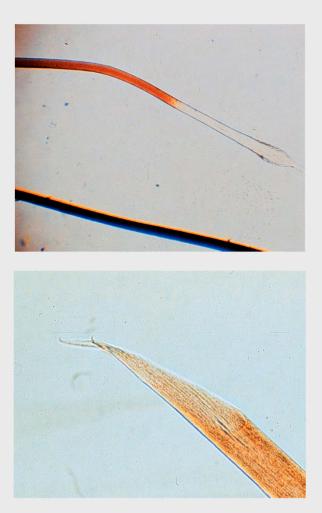
What area of the body is it from?



These carry the most information for microscopic comparisons

- Facial
- Limb
- Chest
- Axial (armpit)

# Damage/disease/treament







# Is it suitable for comparison?

- Unsuitable hairs
  - Damaged
  - Too short
  - Too light in color
  - Fragment
  - Extreme treatment
- Suitable hairs may be compared with suitable known hair samples of the same type
  - Head to head; pubic to pubic

## Comparison process

- Uses a comparison microscope
  - Two microscopes optically joined
  - Split-screen view
- Two samples side-by-side simultaneously
- Use all characteristics available
- Questioned hair must fall within variation established by the Known sample

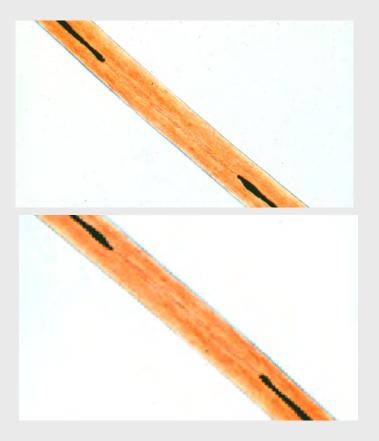
## **Ensemble of Class Traits**

• • • • •	Root	abundant fusi telogen anagen decomp stretched follicular tag cut	• ler • • •	ngth	short medium long thickness	thin	<ul> <li>Cortex</li> <li>Pigment</li> <li></li></ul>	cells prominent cells obscured size of granules shape of granules density local distribution
•	·	broken split pointed round	• • • Me	edulla	fluctuation absent	medium thick	• •	patchy streaky chaining
• • • •	Width Cuticle	coarse fine variation along shaft variation w/in sample thickness variation in thickness clarity	• • • • • •		translucent fragmented transparent discontinuous opaque continuous cell shape thick		• • • • • •	distribution w/in shaft gapping shallow short medium deep long
• • • •	Scales	color protrusion slight medium great	•		thin medium		<ul> <li>Cosmetic</li> <li>Special</li> <li></li></ul>	pigment in cuticle bleached dyed length of time since treatment cracked cuticle ovoid bodies double medulla diseases

vermin

# Inclusion

# Exclusion





# Conclusions

- Positive Association
  - The Q1 head hair exhibits the same microscopic characteristics as the K1 head hairs and, accordingly, could have come from the same source.
    - It should be noted that the microscopic comparison of hairs is not a method of positive identification.
- Exclusion
  - The Q1 head hair exhibits similarities to and differences from the K1 head hair sample. Accordingly, no conclusion could be made as to whether Q1 and K1 could have had a common source.
- Inconclusive
  - The Q1 head hair is microscopically dissimilar to the K1 head hair sample and, accordingly, could not have come from the same source.

# Mitochondrial DNA

- Provides additional information about hairs
- Sequences mtDNA; compared against data bank
  - Maternally inherited
  - Genotype
- Microscopy = phenotype
  - Genotype + environment
- Complementary

Microscopic hair examinations and DNA examinations are complementary techniques—both are more powerful together than separate.

Microscopic examinations evaluate the observed physical characteristics (**phenotype**) of a hair. Molecular examinations provide information about the **genotype** of a hair.



#### Houck and Budowle, 2002

	Microscopic	Mitochondrial
Association	80	97
Inconclusive	37	3
Exclusion	19	64
No Exam	34	6

Results of Microscopic and Mitochondrial DNA Analyses

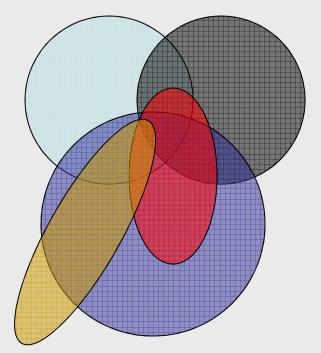
Houck and Budowle, 2000

#### **Mitochondrial Results**



Results of Microscopic and Mitochondrial DNA Analyses by Method

#### Did the hair examiners make a mistake?



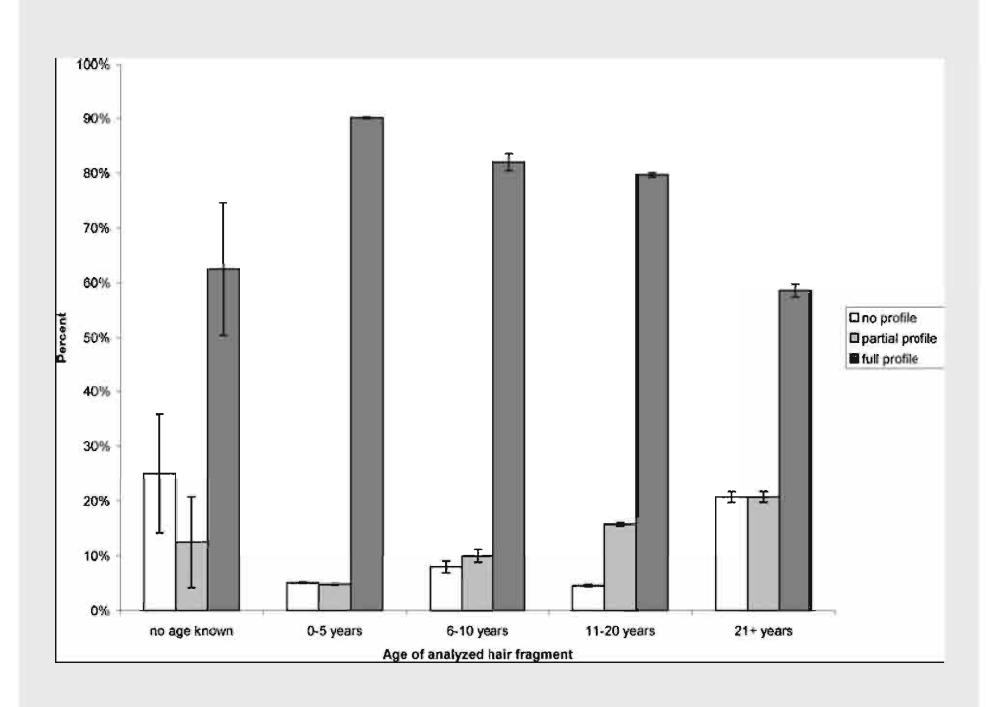
Hairs exhibit a multivariate phenotype

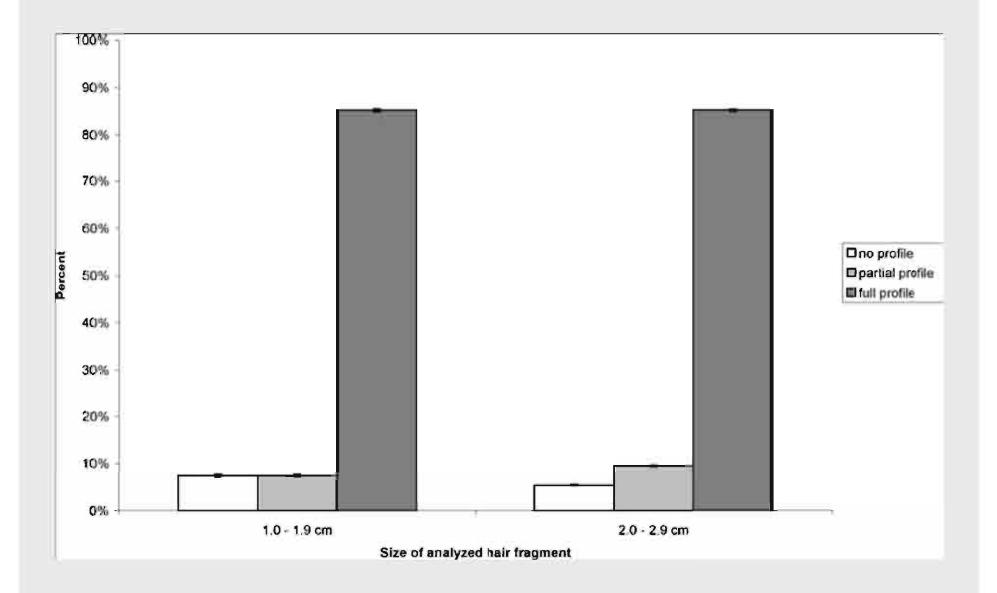
Many traits that overlap to a greater or lesser degree; some may not overlap with others

It's possible but rare for more than one individual to exhibit sufficient characteristics so as to be indistinguishable from another

#### Melton, et al. JFS (2005) 50; 1

- A five year retrospective review of mitochondrial DNA (mtDNA) analysis on 691 casework hairs was carried out.
- A full or partial mtDNA profile was obtained for >92% of hairs.
  - With increasing age of the hair, the likelihood of obtaining a full profile decreased
  - With increasing color and diameter of the hair, the likelihood of obtaining a profile increased.
  - Full or partial profiles were obtained on more than 80% of 114 hairs ≤1.0 cm.
     Mixtures were observed in 8.7% of hairs tested.
    - mixtures increased with the age of the hair and were presumed to be due to exterior surface contamination that could not be sufficiently cleaned prior to extraction, since the overall level of laboratory contamination was low.
- The frequency of sequence heteroplasmy was 11.4%, and both hot-spot and novel sites were observed.





# MtDNA and hair

- About 93% of hairs provide mtDNA info<sup>1</sup>
  - 1.0 cm of hair is typically sufficient
  - Hairs up to 30 years with no significant environmental damage still work
  - Rate of heteroplasmy is about 9-14%
- With telogen roots, hair success rate is independent of
  - cosmetic hair treatments; medulla structure; shaft length, diameter, and volume; and scalp origin.<sup>2</sup>

<sup>1</sup>Melton, et al., *JFS* V50, N1, 2005 <sup>2</sup>Roberts and Calloway, *JFS* V52, N1, 2007