Cracking the DNA mixture code – computer analysis of UK crime cases

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Cybergenetics © 2003-2014

Leicestershire bank robbery



Black balaclava



Brown parka





Brown parka

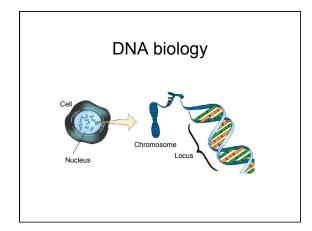


Black balaclava

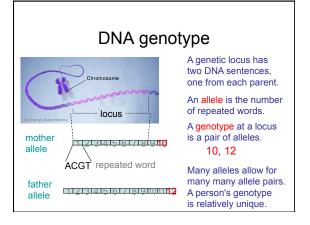


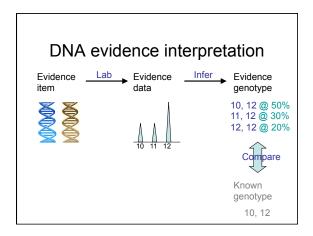
Black shoes

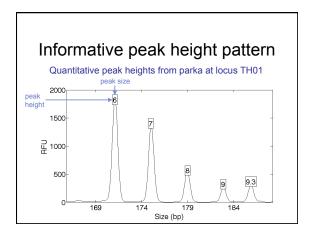


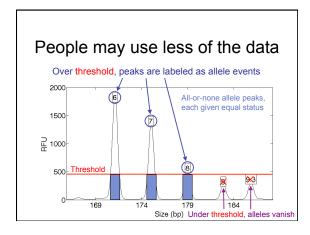


Short tandem repeat DNA locus paragraph It was the best of times, it was the worst of times, it was the epoch of locificity, it was the epoch of belief, it was the epoch of locificity, it was the epoch of locificity, it was the epoch of locificity, it was the epoch of Light, it was the season of Darkness, it was the spring of hope, it was the vinter of despair, we had everyfring before us, we were all going direct the other way - in the epoch of the epoc









DNA lab report: Balaclava

A sample from the inside crown area of the balaclava (item JAS/3) was submitted for DNA profiling tests.

A complex mixed DNA results which appeared to have originated from at least four people was obtained from biological material on the crown.

In my opinion, this result is not suitable for meaningful comparison.

DNA lab report: Shoes

The inside heel area and the toe area of the left shoe (item PAC/2) were submitted separately for DNA profiling tests.

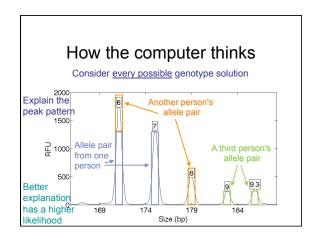
Mixed DNA results which appeared to have originated from at least three people were obtained from biological material on each of the sampled areas. In my opinion, Leroy Williams could have contributed DNA to these results in that all of the components that make up his DNA profile are represented in the result;

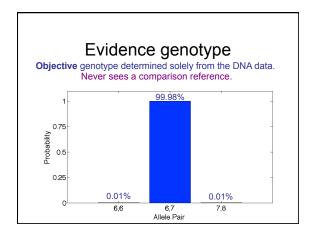
however, due to the overall complexity of the results and the number of contributors to them, a statistical evaluation is not possible.

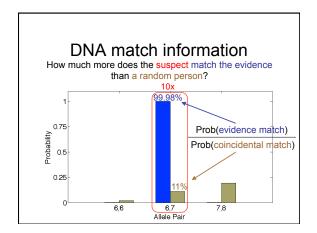
DNA lab report: Parka

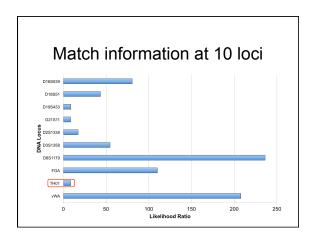
The left and right cuff areas of the brown parka (item CEG/2) were submitted separately for DNA profiling tests. A complex mixed DNA results which appeared to have originated from at least four people was obtained from biological material on the right cuff. In my opinion, this result is not suitable for meaningful comparison.

A mixed DNA result, which appeared to have originated from at least three people was obtained from biological material recovered from the left cuff of the brown parka (CEG/2). In my opinion Leroy Williams could have contributed DNA to this result; however, the finding is not suitable for statistical evaluation.









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Is the suspect in the evidence?

A match between the **parka** and **Leroy Williams** is: **10.2 quadrillion** times more probable than a coincidental match to an unrelated Black person

A match between the **shoe** and **Leroy Williams** is: **13.9 quadrillion** times more probable than a coincidental match to an unrelated Black person

A match between the balaclava and Leroy Williams is: 15.7 quadrillion times more probable than a coincidental match to an unrelated Black person



The TrueAllele option

Objective, reliable truth-seeking tool

- solves the DNA mixture problem
- relatives & up to six contributors
- handles low-copy and degraded DNA
- provides accurate DNA match statistics
- easy to understand, easy to explain
- automates DNA evidence interpretation

When a report says "a meaningful comparison" or "a statistical evaluation" is not possible, TrueAllele is often highly effective.

Reliable: validation studies

Perlin MW, Sinelnikov A. An information gap in DNA evidence interpretation. *PLoS ONE*. 2009;4(12):e8327.

Perlin MW, Legler MM, Spencer CE, Smith JL, Allan WP, Belrose JL, Duceman BW. Validating TrueAllele® DNA mixture interpretation. *Journal of Forensic Sciences*. 2011;56(6):1430-47.

Ballantyne J, Hanson EK, Perlin MW. DNA mixture genotyping by probabilistic computer interpretation of binomially-sampled laser captured cell populations: Combining quantitative data for greater identification information. *Science & Justice*. 2013;53(2):103-14.

Perlin MW, Belrose JL, Duceman BW. New York State TrueAllele® Casework validation study. *Journal of Forensic Sciences*. 2013;58(6):1458-66.

Perlin MW, Dormer K, Hornyak J, Schiermeier-Wood L, Greenspoon S. TrueAllele® Casework on Virginia DNA mixture evidence: computer and manual interpretation in 72 reported criminal cases. *PLOS ONE*. 2014;9(3)e92837.

TrueAllele in criminal trials

About 200 case reports filed on DNA evidence

Court testimony:

- state
- federal
- military
- international

Crimes:

- armed robbery
- child abduction
- child molestation
- murder
- rape
- terrorism
- weapons

United Kingdom & Commonweath countries

TrueAllele has analyzed 22 cases

Country

Crime

Australia (1) Canada (3) England (8) Northern Ireland (10) Armed robbery (1) Identify theft (1) Murder (14) Rape (2) Terror (4)

TrueAllele today

Invented math & algorithms 20 years Developed computer systems 15 years Support users and workflow 10 laboratories Used routinely in casework 3 labs Validate system reliability 20 studies Educate the community 50 talks Train & certify analysts 200 students Go to court for admissibility 5 hearings Testify about LR results 20 trials Educate lawyers and laymen 1,000 people Make the ideas understandable 200 reports

All the DNA, all the time

TrueAllele applications:

- eliminate DNA backlogs
- reduce forensic costs
- solve crimes
- find criminals
- · convict the guilty
- free the innocent
- · create a safer society

More TrueAllele information http://www.cybgen.com/information Courses Newsletters Newsroom Presentations Publications http://www.youtube.com/user/TrueAllele TrueAllele YouTube channel YouTube