DNA Mixture Interpretations and Statistics – To Include or Exclude

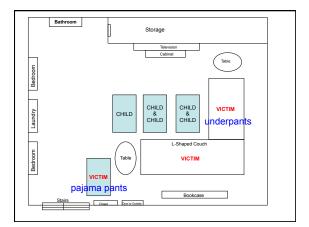
Mark W Perlin, PhD, MD, PhD Cybergenetics, Pittsburgh, PA

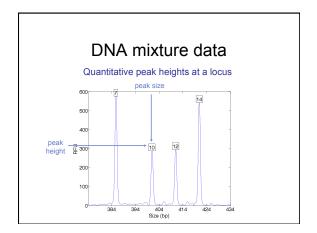
Prescription for Criminal Justice Forensics
ABA Criminal Justice Section

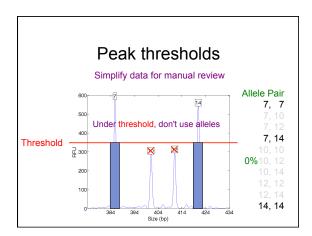


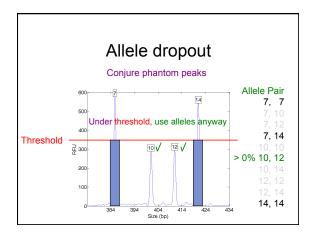
New York, NY June, 2013

Cybergenetics © 2003-2013









Manual interpretation issues

- Evidence call good data inconclusive
- peaks are too low for methods
- too many contributors to handle
- potential examination bias

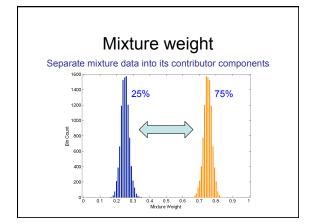
Database

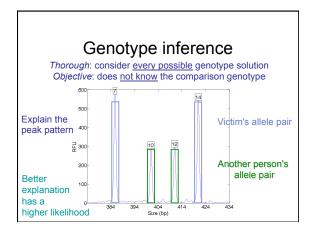
- hit by association, not by match
- comparison: make false hits
- restrict upload: lose true hits

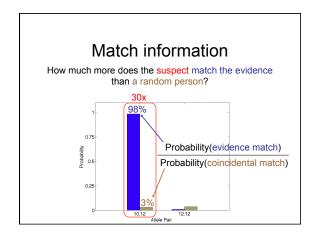
TrueAllele® genotype model

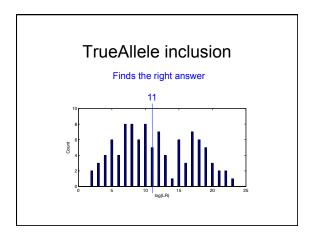
- Evidence preserve data information
- use all peaks, high or low
- any number of contributors
- entirely objective, no bias

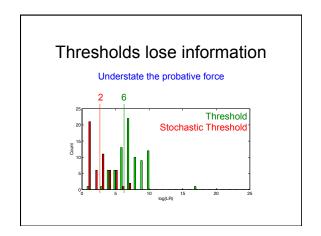
- <u>Database</u>
 hit based on LR match statistic
- sensitive: find true hits
- specific: only true hits

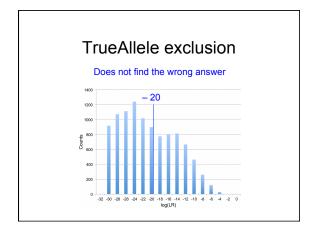












TrueAllele usage

- child molester got: 20 quadrillion match, 22 years
 validation study: 4 peer-reviewed scientific papers
 admissibility: Pennsylvania, California, N. Ireland
 precedent: Pennsylvania appellate ruling
 approval: NYS Commission on Forensic Science
 cases: over 100 reports in criminal cases
 trials: over 15 (rape, murder, terror, weapons, ...)

TrueAllele deployment Casework system Interpretation services

DNA truth, criminal justice

This truth is incontrovertible. Panic may resent it, ignorance may deride it, malice may distort it, but there it is. - Winston Churchill

Genotype models are more accurate. Who cares?

Police Defender Court Citizen

effective database Prosecutor convict the guilty exonerate innocent the whole truth prevent crime

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