

The Problem

Current Procedures are Not Designed for Limiting Amounts of DNA

Significant losses (>75%) in collection & purification steps

Assaying only ~5% of the PCR reaction products

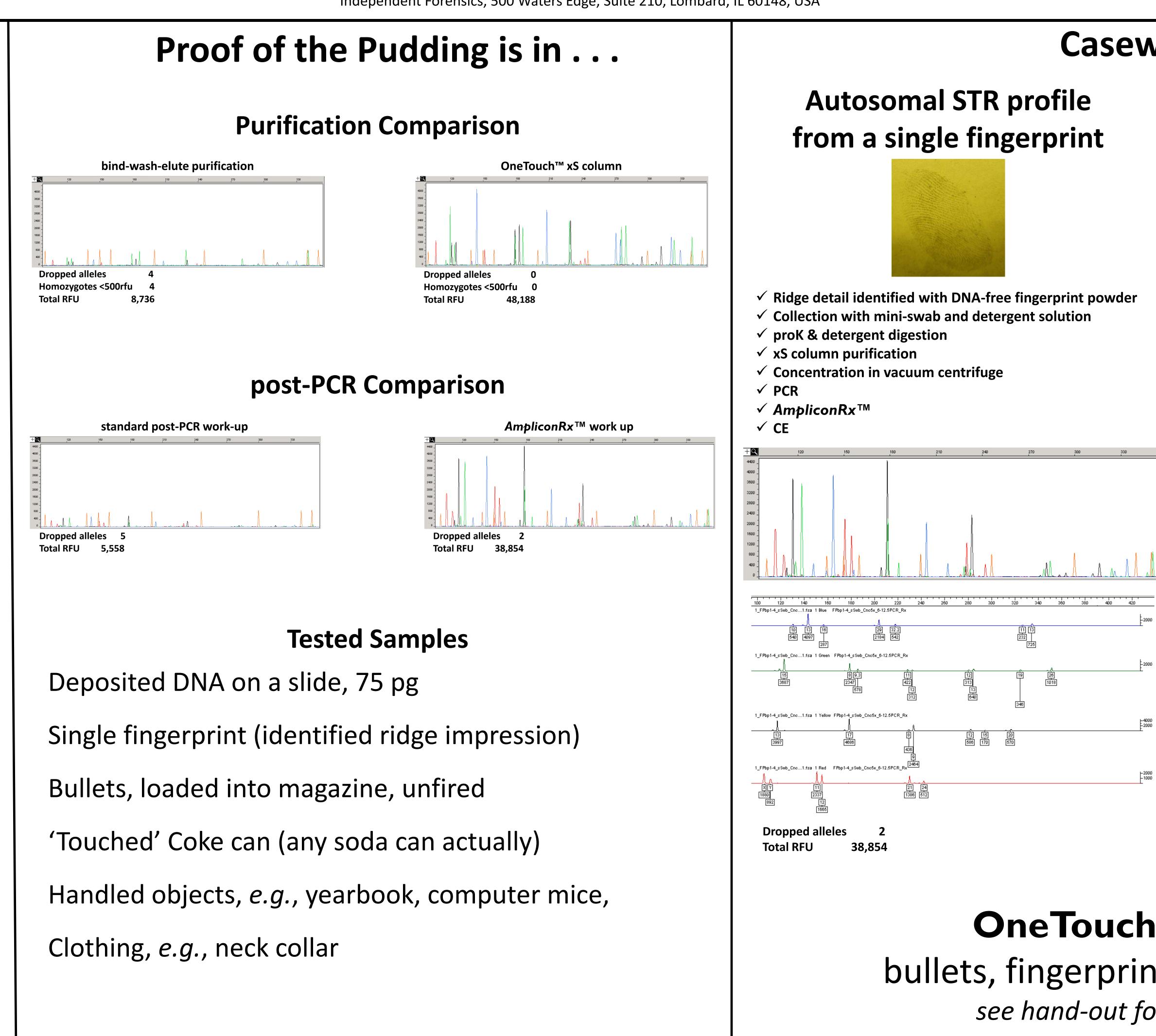
The Solution

- **a**) collecting the biological material: mini-swabs and detergent-based buffer* **secret sauce#1: better swabs and detergent*
- **b**) recovery of collected material: centrifugation at high speed
- c) release of DNA: proK & detergent at elevated °C
- **d**) purification of DNA: removal of inhibitors with 80-90% recovery* *secret sauce#2: NOT bind-wash-elute
- e) PCR amplification: modest changes, 29 v. 28 cycles, 2x TAQ* *better amp with no PCR drop-ins
- **f**) assay 100% of the PCR reaction: post-PCR processing and purification* *secret sauce#3: use all of the PCR reaction, Amplicon Rx[™]

Σ : the most sensitive technique, ever!

Obtaining DNA-Short Tandem Repeat (STR) Profiles from Evidentiary Samples with Extremely Limited Amounts of DNA

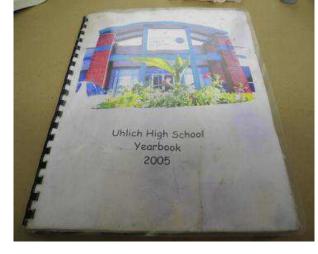
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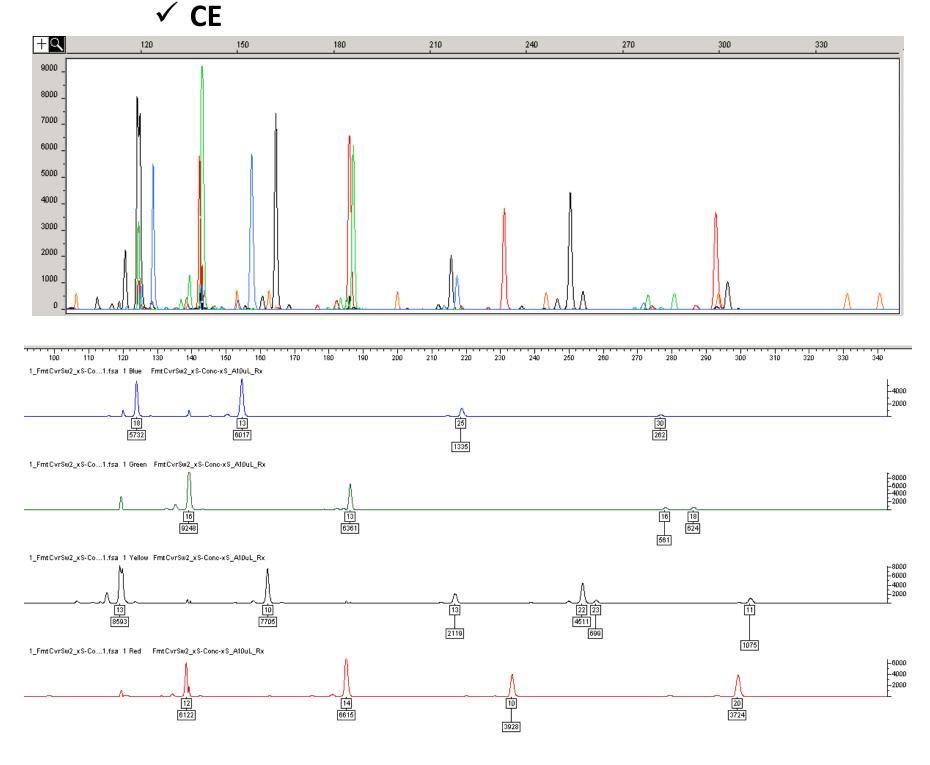
Poster #B9

Casework Examples

Y STR profile from a 'yearbook' cover



- ✓ Entire cover sampled with full-size swab & detergent solution (wet/dry technique)
- ✓ proK & detergent digestion
- \checkmark xS column purification [1st]
- ✓ Concentration in vacuum centrifuge
- \checkmark xS column purification [2nd]
- ✓ PCR
- ✓ AmpliconRx[™]



Dropped alleles Total RFU 77,532

OneTouch DNA Method bullets, fingerprints, handled objects... see hand-out for additional examples