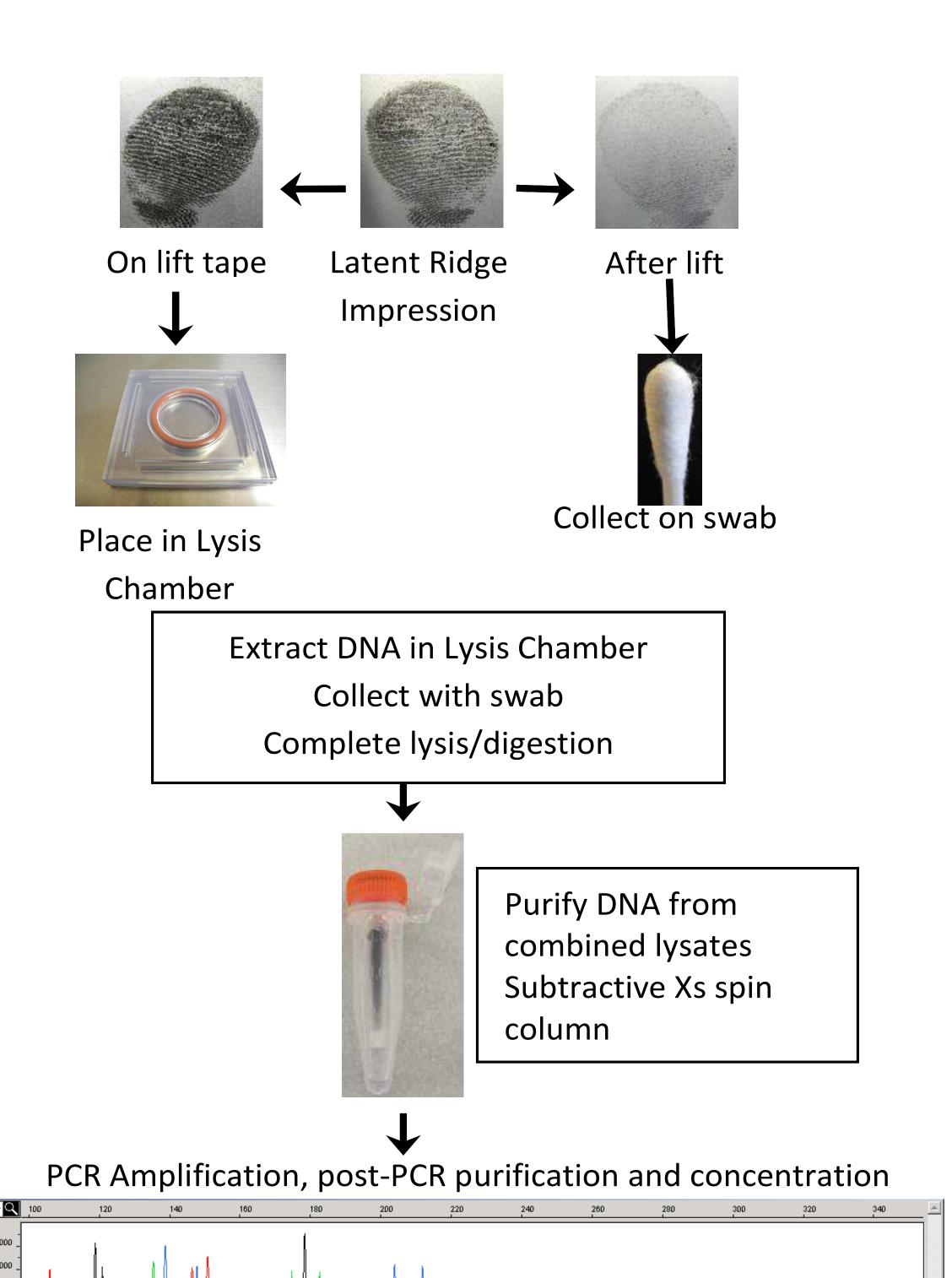


A New SOP: DNA-STR Profiles from Individual Fingerprints on Sticky Tape

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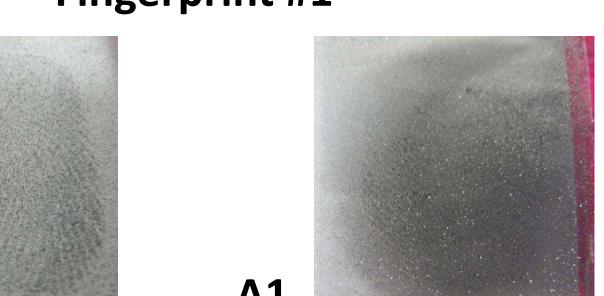
Workflow Overview



SOP Summary

- 1. Identify individual fingerprint on surface of questioned item using fingerprint powder;
- 2. Lift the identified ridge details on a sticky tape/hinge card and collect the 'leftover' of the fingerprint using a sterile cotton swab wet with 10 μ L collection buffer (retain cotton swab for further processing step 4);
- 3. After ridge impression details transferred to tape/hinge card are photographed, immobilize sticky tape in the sticky tape lysis chamber so that the fingerprint on the adhesive side of the tape is exposed; cover tape with adhesive taming material;
- 4.Add 120 µL lysis buffer to tamed tape, close chamber and incubate 1 hr at 56°C; collect lysate with the retained swab (step 2);
- 5. Using spin-basket technique, collect all of the biological material from the swab by centrifugation;
- 6. Incubate recovered material at 56°C for 1 hr to complete digestion of all material;
- 7. Purify total lysate on OneTouch Xs DNA purification column (subtractive DNA purification);
- 8. Assess concentration of DNA in the purified lysate and, if necessary, concentrate 3-fold in a vacuum concentrator;
- 9. Perform multiplex DNA-STR PCR of your choice; if necessary, increase RFU signal by post-PCR cleanup and concentration (AmpliconRx™).

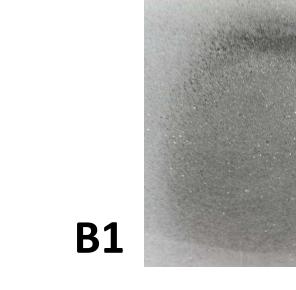
Fingerprint #1



Representative Data

Fingerprint #2

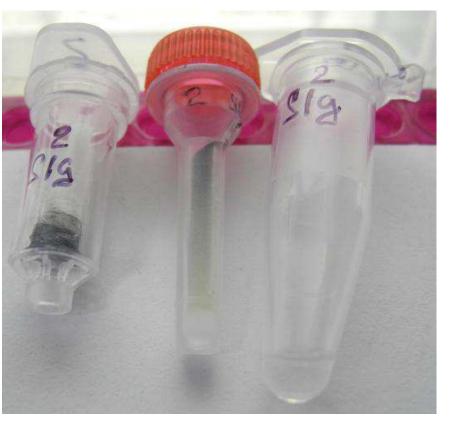




Latent ridge impressions from two fingerprints were identified with fingerprint powder (A and B) and individually transferred to two separate pieces of the lift tape (A1 and B1)

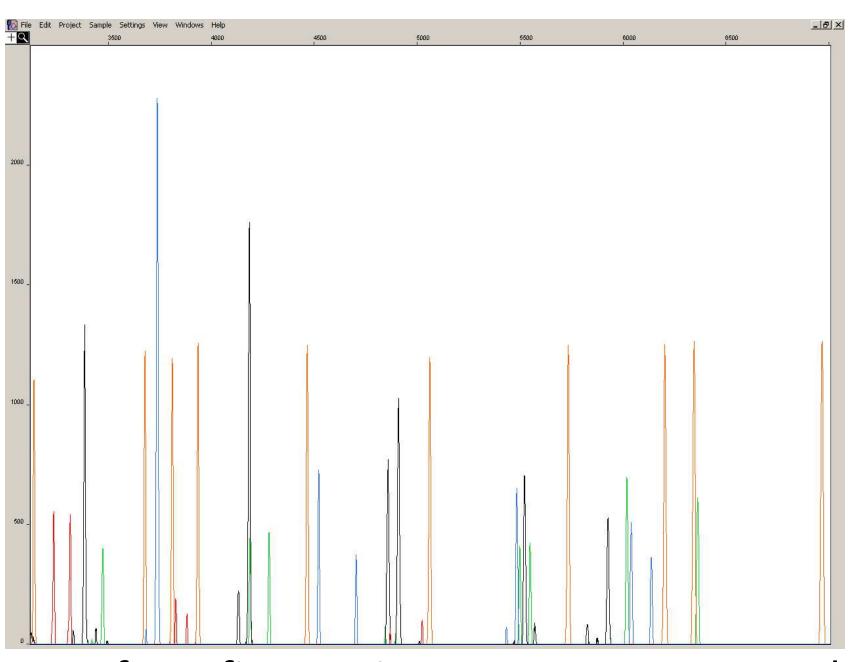


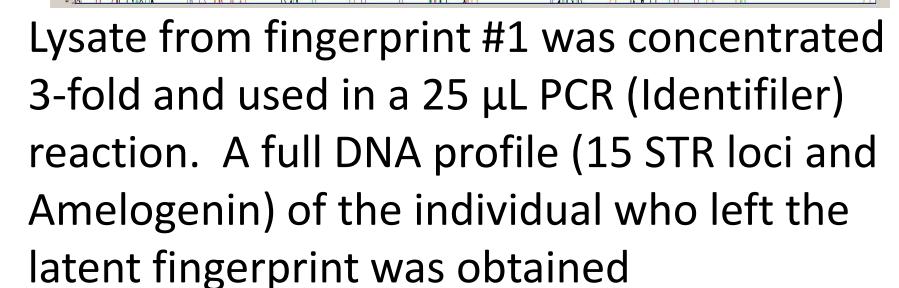
A2

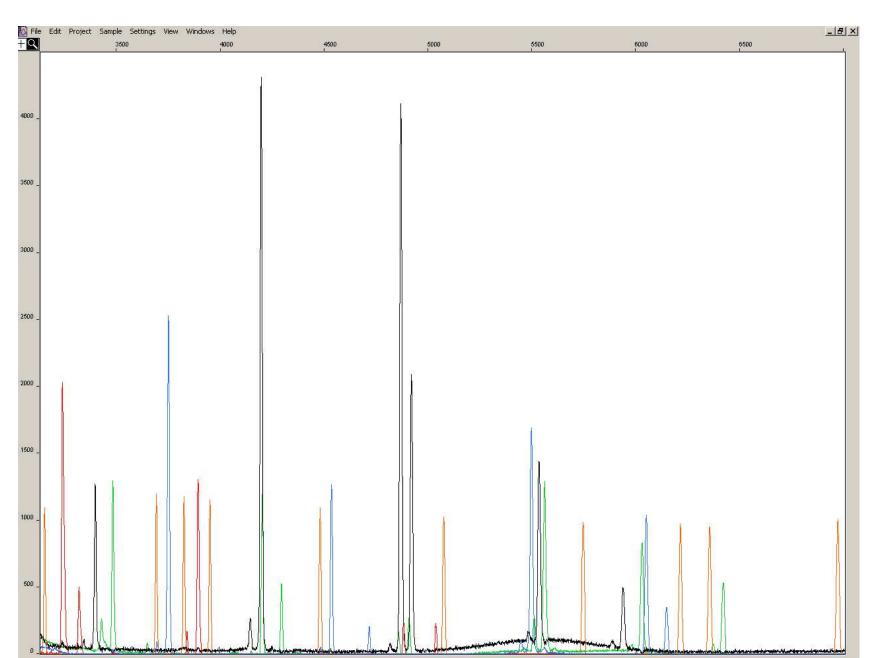


Each fingerprint was processed separately. Figures A2 and B2 show collection swab in a spin basket, DNA purification column, and final purified lysate.

B2







Lysate from fingerprint #2 was concentrated 3-fold and used in a 25 μ L PCR (Identifiler) reaction. After post-PCR purification (Amplicon Rx), a full DNA profile (15 STR loci and Amelogenin) of the individual who left the latent fingerprint was obtained