## FORENSIC BIOLOGY PROTOCOLS FOR FORENSIC STR ANALYSIS

Chelex Extraction from Soft Tissue (e.g. Fetus Samples)		
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## **Chelex Extraction from Soft Tissue (e.g. Fetus Samples)**

Sample sizes for this Chelex extraction should be approximately a 3x3mm cutting of tissue.

- 1. Remove the extraction rack from the refrigerator. Extract either evidence or exemplars. Obtain tubes for the extraction negatives and label them. Have a witness confirm the order of the samples.
- 2. Have a witness confirm that the tube label and entire LIMS input sample ID match for each sample and that the samples are in the correct order.
- 3. Obtain reagents and record lot numbers.
- 4. Pipette 1 mL of sterile or UltraPure deionized water into each of the tubes in the extraction rack. Mix the tubes by inversion or vortexing.
- 5. Incubate at room temperature for 15 to 30 minutes. Mix occasionally by inversion or vortexing.
- 6. Spin in a microcentrifuge for 2 to 3 minutes at 10,000 to 15,000 x g (13,200 rpm).
- 7. Carefully remove supernatant (all but 30 to 50  $\mu$ L).
- To each tube add: 200 μL of 5% Chelex (from a well-resuspended Chelex solution).
  1 μLof 20 mg/mL Proteinase K
- 9. Mix using pipette tip.
- 10. Incubate at 56°C for 60 minutes.
- 11. Vortex at high speed for 5 to 10 seconds.
- 12. Incubate at 100°C for 8 minutes using a screw down rack.
- 13. Vortex at high speed for 5 to 10 seconds.
- 14. Spin in a microcentrifuge for 2 to 3 minutes at 10,000 to 15,000 x g (13,200 rpm).
- 15. Place the LIMS output sample labels on the proper tubes. Confirm that the tube label and entire LIMS output sample ID match for each sample.
- 16. As needed, pipette aliquots of a neat, 1/100 dilution and a 1/10,000 dilution (using TE<sup>-4</sup>) into microcentrifuge tubes for real-time PCR analysis to determine human DNA concentration (refer to Section 4 of the STR manual).
- 17. Store the extracts at 2 to 8°C or frozen.
- 18. In the LIMS system, navigate to the Data Entry page, assign the samples to a storage unit (cryobox), and indicate which samples are completed.