Advanced Topics in Forensic DNA Analysis New Jersey State Police DNA Laboratory Training Workshop December 5-6, 2006

Instructor: John M. Butler (NIST)

Material based largely on Butler, J.M. (2005) Forensic DNA Typing: Biology, Technology, and Genetics of STR Markers. Elsevier Academic Press: New York.

December 5, 2006 (Tuesday)

9:00 – 9:20 a.m. **Introductions** by instructor and participants

9:20 – 10:30 a.m. ABI 310/3100/3130 Capillary Electrophoresis Fundamentals

Advantages of capillary electrophoresis and historical perspectives ABI Genetic Analyzer instrument overview: differences between ABI 310/3100/3130xl Sample preparation, injection, separation, and detection

10:30 – 10:45 a.m. BREAK

10:45 a.m. – 12:15 p.m. **CE Troubleshooting**

Deciphering artifacts from true alleles: dye blobs, spikes, pull-up External factors: temperature shifts and impact on sizing precision Instrument problems: capillary clogging, syringe leaks Troubleshooting benchmarks/QC monitoring (current and syringe travel)

12:15 p.m. – 1:15 p.m. LUNCH

1:15 – 2:45 p.m. **Mixture Interpretation**

Sample interpretation overview NIST MIX05 interlab study results Discussion

2:45 - 3:00 p.m. BREAK

3:00 – 4:30 p.m. qPCR and Low-Copy Number DNA Testing

Real-time PCR and DNA quantitation
Issues and challenges with low-copy number DNA testing

4:30 - 5:00 p.m. Q&A as needed

December 6, 2006 (Wednesday)

9:00 – 9:30 a.m. **miniSTRs**

miniSTRs – tools for improving recovery from degraded DNA MiniFiler kit and concordance studies
Characterization of 26 new miniSTR loci at NIST
New European core STR loci – D2S441, D10S1248, D22S1045

9:30 – 10:30 a.m. Essentials of Validation

Urban Legends of Validation Validation definitions and requirements for documentation Some examples

10:30 – 10:45 a.m. BREAK

10:45 – 12:15 a.m. Fundamentals of Statistics and Population Genetics

How population databases are constructed and validated Calculating random match probabilities Issues of relatedness and mixture statistics Parentage and kinship analysis

12:15 – 1:15 p.m. LUNCH

1:15 – 3:00 p.m. **Mitochondrial DNA**

mtDNA background, structure, and function HV1 & HV2 details LINEAR ARRAYs as potential screening assays mtGenome sequencing to address most common types

3:00 - 3:15 p.m. BREAK

3:15 – 4:30 p.m. **Y-Chromosome Analysis**

Lineage markers Y-STR loci and commercial kits Duplications and deletions

Q&A as needed throughout the various presentations