

Washington State Patrol Crime Laboratory Division Biochemistry and STR Training Program Manual

Washington State Patrol – Crime Laboratory Division

Biochemical and STR Training Program Manual

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1. Introduction and Quality

Welcome to the Washington State Patrol Crime Laboratory Division (WSP CLD). This training manual is intended for candidates who have been successful in obtaining employment in a DNA Unit or CODIS Laboratory and for existing staff who are undergoing additional training in biochemistry and/or DNA STR analysis within the WSP CLD. Unless otherwise specified, the trainee's immediate supervisor is the trainer. The time needed to complete the training program will be determined by the trainer and DNA Technical Leader. (CODIS, HT, Tech)

1.1. Goals

- 1.1.1. The training manual is to guide the trainee to become sufficiently knowledgeable and proficient in forensic biochemistry and DNA analysis to perform the role for which they have been employed.
- 1.1.2. Depending on the trainee's prior education, experience, and background, demonstration of competency in each of the major areas may be all that is required to complete many of the modules.
 - 1.1.2.1. The DNA Technical Leader shall be responsible for assessing the previous training of analysts/technicians with outside experience and ensuring it is adequate and documented. Modification to the training program may be appropriate and shall be approved by the DNA Technical Leader.
- 1.1.3. CODIS DNA Analyst trainees are only required to complete the sections and readings that are relevant to their work duties. Introductory paragraphs and assessment of all assigned modules should be read.
 - 1.1.3.1. Modules 1, 2, 11, 12, 15, 19, 20, 21, and 23
 - 1.1.3.2. Portions of Modules 3, 13, 14, and 17 (marked with "CODIS")
 - 1.1.3.3. Associated readings (marked with "CODIS")
- 1.1.4. High Throughput Laboratory Casework DNA Analyst trainees are only required to complete the sections and readings that are relevant to their work duties. Introductory paragraphs and assessment of all assigned modules should be read.
 - 1.1.4.1. Modules 1, 2, 3, 5, 7, 8, 11, 12, 13, 14, 15, 16, 17, 19, 20, 21, and 23
 - 1.1.4.2. Associated readings (marked with "HT")
- 1.1.5. Laboratory technicians are only required to complete the sections and readings that are relevant to their work duties. Introductory paragraphs and assessment of all assigned modules should be read.
 - 1.1.5.1. Modules 1, 2, and 19
 - 1.1.5.2. Portions of Modules 3, 5, 7, 8, 12, 13, 18, and 20 (marked with "Tech")
 - 1.1.5.3. Associated readings (marked with "Tech")
- 1.1.6. Analysts serving as screeners may need to repeat portions of this training manual (with adjustments), depending on the goals of the training plan at the time.
- 1.1.7. At the completion of this module, the trainee should be able to:
 - 1.1.7.1. Understand the expectations of the training program.
 - 1.1.7.2. Understand the general operation and quality assurance of the laboratory.
 - 1.1.7.3. Be familiar with the laboratory facility.
 - 1.1.7.4. Understand the organizational structure, code of ethics, and chain of command.

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1.1.7.5. Understand security and confidentiality.

1.2. Tasks

- 1.2.1. The trainer will provide the trainee with the necessary instruction and reading materials to complete the training module. Not all trainees will be instructed in all modules.
- 1.2.2. The trainee will get instruction from a variety of secondary trainers, which may include spending time at another WSP laboratory. Alternatively, the training may be outsourced to an accredited vendor, such as the National Forensic Science Technology Center (NFSTC). All outsourced training will follow the guidelines set forth in this training manual with some modifications allowed. Any modifications to the training manual must be approved by the DNA Technical Leader and be documented in the trainee's training file.
 - 1.2.2.1. The method of instruction will include reading, lectures, discussions, demonstrations, observing others perform casework, and observation of others in court.
 - 1.2.2.2. The practical training will include assigned practice exercises and moot court.
- 1.2.3. Training will include written tests, oral tests, competency tests, and/or a final qualifying test prior to casework assignment. A single test and/or competency can be used to cover multiple modules. Refer to the CLD Quality and Operations Manual (QOM) for further specifications on competency testing.
- 1.2.4. The trainee shall keep a training record, which shall include at minimum: notes from the discussions and summary discussions, any completed supplemental practical exercises or readings, the results of competency tests, and documentation of court observations.
- 1.2.5. The trainer will consult with the DNA Technical Leader to plan, schedule, and report the progress of each trainee's program. At the conclusion of each module the primary trainer will assess the trainee's depth of understanding of the material covered and ensure the required readings have been read before documenting the trainee's qualification in that module. An interoffice communication (IOC) may be prepared, addressed to the trainee's supervisor, and forwarded through the appropriate chain of command to the Division Manager upon the employee's successful completion of various phases in the training program. The approval documentation shall also include the DNA Technical Leader. Alternatively, one IOC can be written upon the completion of all modules. Once all members of the appropriate chain of command have signed the IOC, the trainee's supervisor will make arrangements for the trainee to initially perform supervised casework with experienced, qualified forensic scientists for a period of time to be determined by the supervisor. At the end of the training period, the effectiveness of the training actions shall be evaluated and documented.
- 1.2.6. If required, the trainee will complete the new employee orientation modules on the training division iWSP website as required by the New Employee Orientation Supervisor Checklist.
- 1.2.7. The trainee will be introduced to quality assurance and quality control practices of the laboratory.

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1.3. Assessment

No practical exam or competency is provided for this module. The trainer will assess the trainee's knowledge of the subject areas through discussion and will document the training using the trainer's evaluation form.

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WSP DNA BIOCHEMISTRY AND STR TRAINING PROGRAM MODULE 1 – Introduction and Quality

Evaluation Form

	Security and confidentiality
	Introduction to quality assurance/quality control
The t	rainee has completed the above checked sections and is able to: Understand the expectations of the training program Explain the general operation of the laboratory
Comr	nents:
	Trainee Printed Name + Initials Date Trainer Printed Name + Initials Date

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MODULE 1 INTRODUCTION AND QUALITY READING ASSIGNMENTS

CODIS HT Tech

TRAINEE:	

Reference	INITIALS	DATE
WSP CLD Quality Operations Manual: Introduction		
WSP CLD Quality Operations Manual: Scope		
WSP CLD Quality Operations Manual: Definitions		
WSP CLD Quality Operations Manual: Quality Assurance Program		
WSP CLD Quality Operations Manual: Structure, Services and Functions		
WSP CLD Quality Operations Manual: CLD Management and Personnel		
WSP CLD Quality Operations Manual: Personnel Qualifications and Training		
WSP CLD Quality Operations Manual: Document Control Policy and Procedures		
WSP CLD Quality Operations Manual: Quality System Records: Access, Filing, Storage,		
Retention and Disposal		
WSP CLD Quality Operations Manual: Nonconforming Work and Corrective Actions		
WSP CLD Quality Operations Manual: Appendix 1: Root Cause Analysis Guidelines and Procedures		
WSP CLD Quality Operations Manual: Acquisition of Services, Supplies and Equipment		
WSP CLD Quality Operations Manual: Inventories and Reference Collections		
WSP CLD Quality Operations Manual: Audit Program and Management System Review		
WSP CLD Quality Operations Manual: Assuring the Quality of Test Results		
WSP CLD Quality Operations Manual: Technical Procedures and Methods		
WSP CLD Quality Operations Manual: Research Projects, Publications and Presentations		
WSP CLD Quality Operations Manual: Laboratory Facilities and Security		
WSP CLD DNA Analysis Quality Assurance Manual: Goals and Objectives		
WSP CLD DNA Analysis Quality Assurance Manual: Organization and Management		
WSP CLD DNA Analysis Quality Assurance Manual: Personnel Qualifications and		
Training		
WSP CLD DNA Analysis Quality Assurance Manual: Facilities		
WSP CLD DNA Analysis Quality Assurance Manual: Analytical Procedures		
Federal Bureau of Investigation. Quality assurance standards for forensic DNA testing laboratories. Identify significance as related to audits and accreditation. Current version.		
Federal Bureau of Investigation. Audit document for forensic DNA testing laboratories.		
Current version. Review the laboratory's most recent audit findings and responses.		
Federal Bureau of Investigation. Quality assurance standards for databasing		
laboratories. Identify significance as related to audits and accreditation. Current version. CODIS only		
Federal Bureau of Investigation. Audit document for DNA databasing laboratories.		
Current version. Review the laboratory's most recent audit findings and		
responses. CODIS only		

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National Research Council Committee on DNA Forensic Science. 1992. DNA Technology in Forensic Science. Washington, D.C.: National Academies Press. Chapter 4 Ensuring High Standards: 97-110.	
National Research Council Committee on DNA Forensic Science. 1996. The Evaluation of Forensic DNA Evidence. Washington, D.C.: National Academies Press. Chapter 3 Ensuring High Standards of Laboratory Performance: 75-88.	
Butler JM. 2012. Advanced Topics in Forensic DNA Typing: Methodology. San Diego: Elsevier Academic Press. Chapter 7 Quality Assurance and Validation: 167-201.	
Epstein DM, et al. Eliminating sources of pipetting error in the forensic laboratory. Forensic Sci Communications 2003; 5(4): 1-7.	

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2. Safety

The purpose of this module is to familiarize the trainee with general safety precautions and procedures throughout the laboratory. Additional detailed safety precautions (e.g., specific chemical safety) will also be addressed in the applicable sections of the procedures manuals.

2.1. Goals

At the completion of this module, the trainee should be able to:

- 2.1.1. Understand general laboratory safety procedures.
- 2.1.2. Successfully explain the safety precautions that should be taken when handling biological evidence.

2.2. Tasks

Instruction, demonstration, and practical training will be provided. The trainee will be oriented to safety within the laboratory.

- 2.2.1. General laboratory safety topics
 - 2.2.1.1. Fire evacuation plan
 - 2.2.1.2. Earthquake evacuation plan
 - 2.2.1.3. Use of emergency equipment
 - 2.2.1.3.1. First aid kit
 - 2.2.1.3.2. Eye wash
 - 2.2.1.3.3. Emergency shower
 - 2.2.1.3.4. Fire blanket (if applicable)
 - 2.2.1.3.5. Fire extinguisher
 - 2.2.1.4. Use and cleaning of glassware and other equipment
 - 2.2.1.5. Use of electrical equipment
- 2.2.2. Personal protective equipment (PPE)
 - 2.2.2.1. Gloves
 - 2.2.2.2. Laboratory coat (general wear vs. PCR-dedicated)
 - 2.2.2.3. Eye wear
 - 2.2.2.4. Face masks
 - 2.2.2.5. Disposable sleeves
 - 2.2.2.6. Plastic shield
 - 2.2.2.7. Chemical fume hood
 - 2.2.2.8. Biological safety cabinet
- 2.2.3. Chemical safety topics
 - 2.2.3.1. Material Data Safety Sheets (MSDS)
 - 2.2.3.2. Universal safety measures for use of acids and bases
 - 2.2.3.3. Universal safety measures for use of carcinogenic and toxic materials
 - 2.2.3.4. Overview of hazards associated with specific chemicals used in reagents
 - 2.2.3.5. Chemical storage
 - 2.2.3.6. Spill kits
- 2.2.4. Biohazard safety

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Instruction and practical training will be provided on proper handling of liquid and/or wet samples (e.g., liquid blood) and on proper procedures to be used in the event of a biohazard spill (e.g., use of 10% bleach solution). Prevention of transmission of the following infectious diseases during evidence handling, and availability of vaccines, will also be discussed:

- 2.2.4.1. Hepatitis (vaccine available for HBV)
- 2.2.4.2. HIV
- 2.2.4.3. Tuberculosis
- 2.2.5. Hazardous waste materials and other lab-generated waste
 - 2.2.5.1. Chemical
 - 2.2.5.2. Biological
 - 2.2.5.3. Sharps

2.3. Assessment

No practical exam or competency is provided for this module. The trainer will assess through discussion the trainee's knowledge of the subject areas as per the goals stated above and will document the training using the trainer's evaluation form.

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WSP DNA BIOCHEMISTRY AND STR TRAINING PROGRAM MODULE 2 – Safety

Evaluation Form

	Laboratory safety orientation		
	Personal Protective Equipment – PPE		
	Chemical safety topics		
	Biohazard safety		
	Hazardous waste materials and other lab generated waste		
The tra	ainee has completed the above checked sections and is able to: Understand the lab-specific emergency and safety procedures Successfully explain the safety precautions that should be taken when handling biological evidence		
Comm	nents:		
_			
I	Trainee Printed Name + Initials Date Trainer Printed Name + Initials Date		

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MODULE 2 SAFETY READING ASSIGNMENTS

CODIS HT Tech

REFERENCE	INITIALS	DATE
WSP CLD Quality Operations Manual: Health and Safety		
WSP CLD Safety Manual		
WSP Safety and Wellness Manual		
Emergency Procedures (lab specific)		
WSP CLD Biochemical Analysis Procedures: Glossary		
WSP CLD Biochemical Analysis Procedures: Safety		
WSP CLD Biochemical Analysis Procedures: Reagent Preparation		
WSP CLD DNA Casework STR Analysis Procedures: Reagent Preparation		

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3. Evidence Control, Preservation, and Examination

The purpose of this module is to familiarize the trainee with procedures to preserve the integrity of submitted evidence items as well as the use of a documented chain of custody. General principles and standard practices of examining evidence for the presence of biological material and other types of evidence will be explained to the trainee. The trainee will also be introduced to the ideas of contamination and DNA transfer.

3.1. Goals

At the completion of this module, the trainee should be able to:

- 3.1.1. Describe the precautions that must be taken when handling and preserving evidence within the DNA section and for evidence that will be shared between any of the following sections: Biochemistry/DNA, Materials Analysis, Firearms, Toxicology, Questioned Documents, Latent Fingerprints, and Crime Scene Reconstruction.
- 3.1.2. Describe the order of examinations between the DNA Units, Materials Analysis Section, Firearms Section, Toxicology Section, and Latent Fingerprints.
- Explain the administrative process for evidence receipt and maintaining Chain of Custody. (Tech)
- 3.1.4. Determine the relevance of an examination given supporting documentation and the characteristics of the evidence itself.
- 3.1.5. Recognize and minimize any potential for evidence to be compromised or contaminated during examination. (CODIS) (Tech)
- 3.1.6. Adopt standards of case management and documentation of examinations.
- 3.1.7. Successfully explain the administrative process for convicted offender sample receipt as well as their handling and preservation. (CODIS only)

3.2. Tasks

Instruction, demonstration and practical training will be provided.

- 3.2.1. Storage of biological evidence and preventative steps to minimize degradation (Tech)
 - 3.2.1.1. Refrigeration of liquid blood
 - 3.2.1.2. Preparation and storage of dried reference bloodstains
 - 3.2.1.3. Body fluid stains dried and frozen (including special circumstances such as knives, rocks, etc.)
 - 3.2.1.4. Sexual assault evidence dried and frozen
 - 3.2.1.5. Other biological evidence (e.g., hairs, condoms, etc.)
 - 3.2.1.6. Safety
- 3.2.2. Sample collection, examination, and contamination prevention
 - 3.2.2.1. Sample collection for biological trace evidence in conjunction with other laboratory analytical services, crime scene reconstruction and latent fingerprint analysis.
 - 3.2.2.2. Cleanliness of work area and examination tools (Tech)

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- 3.2.2.3. Preserving the integrity of the evidence (prevention of contamination and sample loss) (Tech)
- 3.2.2.4. Casework approach relevance and thoroughness of examinations
- 3.2.2.5. Documentation of examinations
 - 3.2.2.5.1. Notetaking
 - 3.2.2.5.2. Digital photos
 - 3.2.2.5.3. Sketches
 - 3.2.2.5.4. Scanning
- 3.2.2.6. Minimizing the risk of contamination at a PCR level of sensitivity for detection. (CODIS) (Tech)
- 3.2.3. Maintaining the chain of custody (Tech)
 - 3.2.3.1. LIMS
 - 3.2.3.2. Request for Laboratory Examination (RFLE) completion and discrepancies on RFLE
 - 3.2.3.3. Accepting and releasing evidence
 - 3.2.3.4. Creating a new item of evidence
 - 3.2.3.5. Marking and sealing evidence
 - 3.2.3.6. Evidence retained in the laboratory
 - 3.2.3.7. Interlaboratory evidence transfers
 - 3.2.3.8. Convicted offender sample receipt, handling, and LIMS data entry. (CODIS only)
- 3.2.4. Preservation of evidentiary value of items shared between sections
 - 3.2.4.1. Using magnification to identify evidence (stereomicroscope and compound microscope)
 - 3.2.4.2. Collection of trace evidence
- 3.2.5. Conservation of sample
 - 3.2.5.1. Stain collection and substrate control collection
 - 3.2.5.2. Sample collection using the M-Vac® System (if applicable)
 - 3.2.5.3. Saving half the sample
 - 3.2.5.4. Requirement of a letter of consumption of a sample
- 3.2.6. Evidence storage during analysis (Tech)
 - 3.2.6.1. Temperature of storage during analysis
 - 3.2.6.2. Items stored at laboratory

3.3. Assessment

Casework Analysts are responsible for sample collection and evidence handling to prevent contamination and cross-contamination. CODIS DNA Analysts are responsible for sample handling to prevent contamination and cross-contamination, Convicted Offender Form entry, and receipt and handling of convicted offender samples. All material in this module should be reviewed by experienced staff training in this area to ensure their knowledge is current. No practical exam or competency is provided for this module. The trainer will assess the trainee's knowledge of the subject areas through discussion and will document the training using the trainer's evaluation form.

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WSP DNA BIOCHEMISTRY AND STR TRAINING PROGRAM MODULE 3 – Evidence Control, Preservation, and Examination Evaluation Form

analytical services and latent fingerprint analysis
Storage of biological evidence and preventative steps to minimize degradation (Tech)
Maintaining chain of custody (Tech)
Conservation of sample
Evidence storage during analysis (Tech)
Minimizing the risk of contamination at a PCR level of sensitivity for detection (CODIS) (Tech)
Cleanliness of work area and examination tools (Tech)
Documentation and thoroughness of examinations
Casework approach
Convicted offender administrative process including: sample receipt, handling, and Convicted Offender Form data entry (CODIS only)

The trainee has completed the above checked sections and is able to:

Successfully explain the proper procedures and precautions to be taken when handling and preserving evidence for DNA and latent fingerprint analysis

Describe the order of examinations between the DNA Units, Materials Analysis Section, Firearms Section, Toxicology Laboratory, Latent Fingerprints, and Crime Scene Reconstruction

Explain the administrative process for evidence receipt and maintaining chain of custody

Determine the relevancy of an examination given characteristics of the evidence and supporting documentation

Recognize and minimize potential for evidence to be compromised during examination Adopt standards of case management and documentation of examinations Understand Convicted Offender Form data entry (CODIS only)

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Comments:			
Trainee Printed Name + Initials	Date	Trainer Printed Name + Initials	Date

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MODULE 3 EVIDENCE CONTROL, PRESERVATION, AND EXAMINATION READING ASSIGNMENTS

REFERENCE	INITIALS	DATE
WSP FLSB Forensic Services Guide HT Tech		
WSP CLD Quality Operations Manual: Evidence Management HT		
WSP CLD Quality Operations Manual: Sampling and Sample Selection HT		
WSP CLD Laboratory Information Management System Operations Manual: Section 10, 14-17, 18-18.05, 18.07-18.09, 19, 22, 23.05, 24-24.05, 24.07-24.08, 26 HT Tech	ns 1-	
ASTM International Committee E-30.11. Standard Guide for Physical Evidence		
Labeling and Related Documentation. 2018: E1459-13. HT		
ASTM International Committee E-30.11. Standard Practice for Receiving,		
Documenting, Storing, and Retrieving Evidence in a Forensic Laboratory. 2 E1492-11. HT	2017:	
Lee HC, Ladd C. Preservation and collection of biological evidence. Croatian Med 2001; 42(3): 225-228. HT		
National Institute of Standards and Technology. The Biological Evidence Preserva	ation	
Handbook: Best Practices for Evidence Handlers. 2013: 9-24. HT Szkuta B, et al. Potential degrading effect of sodium hypochlorite on exhibits		
containing DNA. Forensic Sci Int Genet Supp Ser 2015; 5: E52-E54.		
Forensic Science Regulator. 2015. The Control and Avoidance of Contamination i Laboratory Activities Involving DNA Evidence Recovery and Analysis. Issue	e 1	
(FSR-G-208): Sections 8.4 Control of Bench Environment, 8.5 Use of DNA Laboratories for Activities Other Than Casework, 8.6 Cleaning Processes, 8.7 Environment Magicalian 22.22 LIT		
8.7 Environmental Monitoring. 22-32. HT Gill P. The utility of 'substrate controls' in relation to 'contamination'. Forensic Sci I	Int	
1997; 85(2): 105-111. HT		
WSP CLD DNA summary for the potential cross-contamination between packaged samples (shipping contamination study). HT	d	
Sullivan K, et al. New developments and challenges in the use of the UK DNA database: addressing the issue of contaminated consumables. Forensic Sc Supp Ser 2004; 146S: S175-S176. HT Tech	ci Int	
Gill P, Kirkham A. Development of a simulation model to assess the impact of contamination in casework using STRs. J Forensic Sci 2004; 49(3): 485-49	1.	
Sundquist T, Bessetti J. Identifying and preventing DNA contamination in a DNA-ty laboratory. Profiles in DNA 2005; Sept: 11-13. HT Tech	yping	
Ladd C, et al. A systematic analysis of secondary DNA transfer. J Forensic Sci 19: 44(6): 1270-1272.		
Wickenheiser RA. Trace DNA: a review, discussion of theory, and application of the		
transfer of trace quantities of DNA through skin contact. J Forensic Sci 200 47(3): 442-450. HT)2;	
Fonneløp AE, et al. Secondary and subsequent DNA transfer during criminal investigation. Forensic Sci Int Genet 2015; 17: 155-162.		
Phipps M, Petricevic S. The tendency of individuals to transfer DNA to handled iter Forensic Sci Int 2007; 168(2-3): 162-168.	ms.	
Wiegand P, et al. Transfer of biological stains from different surfaces. Int J Legal N 2011; 125(5): 727-731.	Med	
Taylor D, et al. Helping to distinguish primary from secondary transfer events for tr DNA. Forensic Sci Int Genet 2017; 28: 155-177.	race	
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van Oorschot RAH, et al. DNA transfer in forensic science: a review. Forensic Sci Int	
Genet 2019; 38: 140-166.	
Toothman MH, et al. Characterization of human DNA in environmental samples.	
Forensic Sci Int 2008; 178(1): 7-15.	
Amick J, et al. Integrating DNA collection into the latent print section. J Forensic	
Identification 2004; 54(2): 170-177.	
California Criminalistics Institute. Summary of experiments investigating the impact of	
fingerprint processing and fingerprint reagents on PCR-based DNA typing	
profiles.	
WSP CLD Latent Prints Technical Manual: Items with Additional Requests for DNA or	
Biochemical Examination HT	
WSP CLD Biochemical Analysis Procedures: Introduction and General Exam	
Procedures HT read all, Tech read General Exam Procedures: General	
Instructions Regarding Relevance of Examinations only	
WSP CLD Biochemical Analysis Procedures: Whole Blood Processing	
WSP CLD CODIS Laboratory STR Analysis Procedures Manual: Processing of	
Convicted Offender Samples CODIS only	
WSP CLD DNA Analysis Quality Assurance Manual: Casework Evidence and Sample	
Control CODIS Tech	
WSP CLD DNA Casework STR Analysis Procedures: DNA Extract and Work Product	
Transfer/Return HT	
WSP CLD DNA Analysis Quality Assurance Manual: Convicted Offender Sample	
Control CODIS only	
WSP JusticeTrax W2 to LIMS-plus Interface User Guide CODIS only	
WSP CLD in-house study for Testing the Effectiveness of the Stratalinker UV	
Crosslinker in Eliminating Contaminating DNA from Laboratory Consumables	
CODIS only	
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4. Alternate Light Source

The purpose of this module is to familiarize the trainee with the proper use of the Alternate Light Source (ALS) for examining evidence for the presence of biological material.

4.1. Goals

At the completion of this module, the trainee should be able to operate the ALS safely to locate possible biological material.

4.2. Tasks

Instruction, demonstration and practical training will be provided.

- 4.2.1. Safety of operation of the ALS
- 4.2.2. Appropriate wavelengths and filters
- 4.2.3. Procedure for examination of evidence
- 4.2.4. Materials that fluoresce
- 4.2.5. Documentation of examination
- 4.2.6. Interpretation and conclusions

4.3. Assessment

Examine a variety of both known and unknown materials from biological, chemical, and physical sources to familiarize the trainee with a range of materials that may be encountered in casework. The substances should be examined on various substrates. No competency exam is provided for this module. The trainer will assess through discussion the trainee's knowledge of the subject areas as per the goals stated above and will document the training using the trainer's evaluation form.

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WSP DNA BIOCHEMISTRY AND STR TRAINING PROGRAM MODULE 4 – Alternate Light Source

Evaluation Form

		Cofety of an arction of the ALC
	Ш	Safety of operation of the ALS
		Appropriate wavelengths and filters
		Procedure for examination of evidence
		Materials that fluoresce
		Documentation
		Interpretation and conclusions
The t		as completed the above checked sections and is able to: te the ALS to locate possible biological material on items similar to what would be encountered in casework
Comr	ments:	
	Trainee	Printed Name + Initials Date Trainer Printed Name + Initials Date

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MODULE 4 ALTERNATE LIGHT SOURCE READING ASSIGNMENTS

Reference	INITIALS	DATE
Alternate Light Source User's Manual (lab specific)		
WSP CLD Biochemical Analysis Procedures: Alternate Light Source		

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Identification of Blood

The purpose of this module is to familiarize the trainee with accepted protocols for the presumptive and confirmatory testing for the presence and identification of blood.

5.1. Goals

At the completion of this module, the trainee should be able to:

- 5.1.1. Test stains using proper procedures for Kastle Meyer (phenolphthalein) and HemaTrace® tests, as applicable.
- 5.1.2. Interpret test results and draw appropriate conclusions.
- 5.1.3. Be familiar with other presumptive testing methods.
- 5.1.4. Know the components of blood and their functions

5.2. Tasks

Instruction, demonstration and practical training will be provided.

- 5.2.1. Physical and chemical characteristics of blood
 - 5.2.1.1. Components of blood and their function
 - 5.2.1.2. Visual appearance (overall)
 - 5.2.1.3. Stereomicroscopic appearance
 - 5.2.1.4. Effects of degradation and aging
- 5.2.2. Reagent preparation (Tech)
 - 5.2.2.1. Phenolphthalein
 - 5.2.2.2. Stock and working solutions
 - 5.2.2.3. Quality control testing of reagents and documentation
- 5.2.3. Presumptive testing
 - 5.2.3.1. Phenolphthalein
 - 5.2.3.1.1. Biochemical basis, procedure, and the value of a two-step test
 - 5.2.3.1.2. Control samples
 - 5.2.3.1.3. Potential false positives
 - 5.2.3.1.4. Documentation
 - 5.2.3.1.5. Interpretation and conclusions
 - 5.2.3.2. Other catalytic tests
 - 5.2.3.2.1. Leucocrystal violet
 - 5.2.3.2.2. Tetramethyl Benzidine (TMB, Hemastix®)
 - 5.2.3.2.3. Ortho-Tolidine
 - 5.2.3.3. Blood enhancement
 - 5.2.3.3.1. Leucocrystal violet
 - 5.2.3.3.2. Luminol and other luminescent reagents as available
- 5.2.4. Confirmatory testing
 - 5.2.4.1. Abacus OneStep HemaTrace® cards
 - 5.2.4.1.1. Biochemical basis and procedure
 - 5.2.4.1.2. High dose hook effect
 - 5.2.4.1.3. Specificity and sensitivity

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- 5.2.4.1.4. Documentation
- 5.2.4.1.5. Interpretation and conclusions
- 5.2.5. Effects of presumptive and confirmatory reagents on additional (e.g., STR) testing

5.3. Assessment

Test samples of known blood, rust, plant material, and other materials reported in the literature to give false positive presumptive tests. Create and test dilutions of blood up to 1:1000. Use various collection methods (e.g., moistened swab, filter paper, etc.). Test serum and whole blood, if possible. Prepare serial dilutions and laundered stains and test with phenolphthalein and all applicable confirmatory tests. Practice sets should include samples of varying strengths and mixtures of bodily fluids to ensure ability to determine differences between positive, negative, and inconclusive results, specifically when utilizing card-based testing methods.

COMPETENCY: A minimum of ten correctly characterized stains with the properly reported conclusions (five stains for HT trainees). This competency may be performed in concert with competencies for other body fluids.

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WSP DNA BIOCHEMISTRY AND STR TRAINING PROGRAM MODULE 5 – Identification of Blood

Evaluation Form

	Physical and chemical characteristics of blood
	Reagent preparation (Tech)
	Presumptive testing
	Confirmatory testing
	Effects of presumptive and confirmatory reagents on additional testing
	Documentation
	Interpretation and conclusions
	Competency
The t	trainee has completed the above checked sections and is able to: Test stains using proper procedures for phenolphthalein and HemaTrace®, as applicable Interpret test results and draw appropriate conclusions Be familiar with other presumptive testing methods Know the components of blood
Com	ments:
	Trainee Printed Name + Initials Date Trainer Printed Name + Initials Date

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MODULE 5 IDENTIFICATION OF BLOOD READING ASSIGNMENTS

REFERENCE	INITIALS	DATE
Gaensslen RE. 1983. Sourcebook in Forensic Serology, Immunology, and		
Biochemistry. Washington, D.C.: U.S. Department of Justice. Sections 4.2.4		
Hemochromogen crystal test: 85-87 and 6 Catalytic Tests: 101-116. HT		
Lee HC. 1982. Forensic Science Handbook. Identification and Grouping of Bloodstains.		
Englewood Cliffs (NJ): Prentice Hall. 272-279. HT		
Spalding RP, Cronin WF. 1984. Technical and Legal Aspects of Forensic Serology: A		
Laboratory Manual. Washington, D.C.: U.S. Department of Justice. Part I,		
Chapter I Blood – A General Facts Sketch, Sections I-III A; 1-3. HT		
Spalding RP, Cronin WF. 1984. Technical and Legal Aspects of Forensic Serology: A		
Laboratory Manual. Washington, D.C.: U.S. Department of Justice. Part I,		
Chapter II Chemistry of Tests used for the Identification of Blood, Section II B; 10.		
HT		
Spalding RP, Cronin WF. 1984. Technical and Legal Aspects of Forensic Serology: A		
Laboratory Manual. Washington, D.C.: U.S. Department of Justice. Part I,		
Chapter II Chemistry of Tests used for the Identification of Blood, Section III B;		
14-15. HT		
Spalding RP, Cronin WF. 1984. Technical and Legal Aspects of Forensic Serology: A Laboratory Manual. Washington, D.C.: U.S. Department of Justice. Part I,		
Chapter III Testing Methods for the Identification of Blood, Sections I and II B; 16,		
18-19. HT		
Spalding RP, Cronin WF. 1984. Technical and Legal Aspects of Forensic Serology: A		
Laboratory Manual. Washington, D.C.: U.S. Department of Justice. Part I,		
Chapter III Testing Methods for the Identification of Blood, Section IIIB; 22-24. HT		
Higaki RS, Philp WMS. A study of the sensitivity, stability and specificity of		
phenolphthalein as an indicator for blood. Can Soc Forensic Sci J. 1976; 9(3):		
97-102. HT		
Blake ET, Dillon DJ. Microorganisms and the presumptive tests for blood. J Police Sci		
and Admin. 1(4): 395-400. HT		
Cox M. A study of the sensitivity and specificity of four presumptive tests for blood. J		
For Sci. 1991; 36(5): 1503-1511. HT		
Cox M. Effect of fabric washing on the presumptive identification of bloodstains. J For		
Sci. 1990; 35(6): 1335-1341.		
WSP CLD Biochemical Analysis Procedures: Phenolphthalein Test (Modified Kastle-		
Meyer Test) Presumptive Test for Blood HT Tech		
Abacus HemaTrace® package insert HT Tech		
WSP CLD Biochemical Analysis Procedures: ABAcard HemaTrace® Human Blood Test		
Tech		
Hochmeister MN, et al. Validation studies of an immunochromatographic 1-step test for		
the forensic identification of human blood. J Forensic Sci. 1999; 44(3): 597-602.		
HT		
Rowley B. Commentary on Hochmeister MN, et al. Validation studies of an		
immunochromatographic 1-step test for the forensic identification of human		
blood. J Forensic Sci. 1999; 44(6): 1323-1324. HT		

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6. Bloodstain Pattern Interpretation

The purpose of this module is to familiarize the trainee with typical bloodstain patterns encountered in casework. This will not result in the trainee becoming proficient in bloodstain pattern analysis. However, the expectation is that they will be able to recognize when this analysis may be necessary.

6.1. Goals

6.1.1. Recognize when consultation with a qualified bloodstain pattern interpretation analyst would be beneficial.

6.2. Tasks

Instruction, demonstration and practical training will be provided.

- 6.2.1. Theory
- 6.2.2. Recognition of bloodstain patterns
- 6.2.3. Documentation of bloodstain patterns
- 6.2.4. Descriptive vocabulary

6.3. Assessment

Use liquid human blood (if possible) to create bloodstain patterns (e.g., dripping, contact transfer). Examine blood drops on various substrates, dropped from various angles, and in varying amounts. No competency exam is provided for this module. The trainer will assess through discussion the trainee's knowledge of the subject areas as per the goal stated above and will document the training using the trainer's evaluation form.

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WSP DNA BIOCHEMISTRY AND STR TRAINING PROGRAM MODULE 6 – Bloodstain Pattern Interpretation Evaluation Form

	Theory
	Recognition of bloodstain patterns
	Documentation of bloodstain patterns
	Descriptive vocabulary
The tra	ainee has completed the above checked sections and is able to: Recognize typical bloodstain patterns and determine when further analysis is necessary
Comm	ents:
_	
Т	rainee Printed Name + Initials Date Trainer Printed Name + Initials Date

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MODULE 6 BLOODSTAIN PATTERN INTERPRETATION READING ASSIGNMENTS

Reference	INITIALS	DATE
Bevel T, Gardner RM. 2008. Bloodstain Pattern Analysis. CRC Press. Chapter 1		
Bloodstain Pattern Analysis: Its Function and a Historical Perspective: 1-15.		
Bevel T, Gardner RM. 2008. Bloodstain Pattern Analysis. CRC Press. Chapter 2		
Bloodstain Pattern Terminology: 17-36.		
Bevel T, Gardner RM. 2008. Bloodstain Pattern Analysis. CRC Press. Chapter 3		
Bloodstain Classification: 37-87.		
Bevel T, Gardner RM. 2008. Bloodstain Pattern Analysis. CRC Press. Chapter 5 The		
Medium of Blood: 111-133.		
Bevel T, Gardner RM. 2008. Bloodstain Pattern Analysis. CRC Press. Chapter 10		
Understanding and Applying Characteristic Patterns of Blood: 231-259.		
Bevel T, Gardner RM. 2008. Bloodstain Pattern Analysis. CRC Press. Chapter 11		
Bloodstained Clothing Issues: 261-274.		
Bevel T, Gardner RM. 2008. Bloodstain Pattern Analysis. CRC Press. Chapter 13		
Documenting Bloodstains: 297-317.		

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7. Identification of Saliva

The purpose of this module is to familiarize the trainee with accepted protocols used to determine the presence of amylase, an enzyme found in elevated levels in saliva.

7.1. Goals

At the completion of this module, the trainee should be able to:

- 7.1.1. Test stains using proper procedures for Rapid Stain Identification (RSID)™-Saliva cards and Phadebas® paper.
- 7.1.2. Interpret test results and draw appropriate conclusions.

7.2. Tasks

Instruction, demonstration, and practical training will be provided.

- 7.2.1. Relevance of examination in casework
- 7.2.2. Examination approach
 - 7.2.2.1. ALS
 - 7.2.2.2. Amylase mapping
 - 7.2.2.3. Swabbing
 - 7.2.2.4. Sampling
- 7.2.3. Phadebas® paper (Tech)
 - 7.2.3.1. Biochemical basis and procedure
 - 7.2.3.2. Potential false positives
 - 7.2.3.3. Documentation
 - 7.2.3.4. Interpretation and conclusions
- 7.2.4. RSID™-Saliva cards (Tech)
 - 7.2.4.1. Biochemical basis and procedure
 - 7.2.4.2. Potential false positives
 - 7.2.4.3. Documentation
 - 7.2.4.4. Interpretation and conclusions

7.3. Assessment

Use ALS amylase mapping to identify potential saliva stains. Using RSID™-Saliva cards and Phadebas® paper, create and test dilutions of saliva up to 1:1000 and test at least one potential false positive. Practice sets must include samples of varying strengths and mixtures of bodily fluids to ensure ability to determine differences between positive, negative, and inconclusive results, specifically when utilizing card-based testing methods.

COMPETENCY: A minimum of five correctly characterized stains with the properly reported conclusions. This competency may be performed in concert with competencies for other body fluids.

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	Relevance of examination in casework
	Examination approach
	RSID™-Saliva cards (Tech)
	Phadebas® paper (Tech)
	Documentation
	Interpretation and conclusions
	Competency
The tr	ainee has completed the above checked sections and is able to: Test stains using proper procedures for Rapid Stain Identification (RSID)™-Saliva cards and Phadebas [®] paper
Comm	nents:
-	Trainee Printed Name + Initials Date Trainer Printed Name + Initials Date

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MODULE 7 IDENTIFICATION OF SALIVA READING ASSIGNMENTS

TRAINEE:	

Reference	INITIALS	DATE
Nelson DF, Kirk PL. The identification of saliva. J Forensic Med. 1963; 10(1): 14-20.		
Gaensslen RE. 1983. Sourcebook in Forensic Serology, Immunology, and Biochemistry. Washington, D.C.: U.S. Department of Justice. Sections 11.3 Amylase: 184-187 and 11.4 Immunological Methods: 187-189.		
Stiefel DJ, Keller PJ. Preparation and some properties of human pancreatic amylase including a comparison with human parotid amylase. Biochem Biophys Acta. 1973; 302(2): 345-361.		
Auvdel MJ. Amylase levels in semen and saliva stains. J Forensic Sci 1986; 31(2): 426-431.		
Keating SM, Higgs DF. The detection of amylase on swabs from sexual assault cases. J Forensic Sci Society 1994; 34(2): 89-93.		
Phadebas® product insert Tech		
WSP CLD Biochemical Analysis Procedures: Phadebas® Paper Amylase Diffusion Tech		
Willott GM, Griffiths M. A new method for locating saliva stains – spotty paper for spotting spit. Forensic Sci Int 1980; 15(1): 79-83.		
Meyers JR, Adkins WK. Comparison of modern screening techniques for saliva screening. J Forensic Sci 2008: 53(4): 862-867.		
Rapid Stain IDentification™ (saliva) product insert Tech		
WSP CLD Biochemical Analysis Procedures: Rapid Stain Identification of Human Saliva (RSID™) Cards Tech		
Ricci U, et al. False-positive results with amylase testing of citrus fruits. J Forensic Sci 2014; 59(5): 1410-1412.		

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8. Identification of Semen

The purpose of this module is to familiarize the trainee with the accepted protocols for the presumptive and confirmatory identification of semen.

8.1. Goals

At the completion of this module, the trainee should be able to:

- 8.1.1. Describe the physical and chemical characteristics of semen and the morphology of spermatozoa.
- 8.1.2. Test evidence items either directly or with a mapping technique to determine the location of possible semen stains by detecting acid phosphatase.
- 8.1.3. Produce a cell pellet, prepare a slide, stain the slide, and positively identify spermatozoa under a microscope. Characterize other material that may be present on the slide.
- 8.1.4. Test swabs and other material for the possible presence of semen using a p30 card.
- 8.1.5. Interpret results and draw appropriate conclusions.

8.2. Tasks

Instruction, demonstration, and practical training will be provided

- 8.2.1. Physical and chemical characteristics of semen
 - 8.2.1.1. Components of semen
 - 8.2.1.2. Spermatozoa morphology
 - 8.2.1.3. Typical volume of ejaculate
 - 8.2.1.4. Typical number of spermatozoa per volume
 - 8.2.1.5. Azoospermia
 - 8.2.1.6. Persistence of semen
- 8.2.2. Acid Phosphatase (Tech)
 - 8.2.2.1. Reagent preparation
 - 8.2.2.2. Quality control testing of reagents and documentation
 - 8.2.2.3. Mapping
 - 8.2.2.4. Sample swabbing and/or evidence swab testing
 - 8.2.2.5. Controls
 - 8.2.2.6. Biochemistry of reaction and time to color development
 - 8.2.2.7. Interpretation and conclusions
 - 8.2.2.8. False positives
- 8.2.3. Identification of spermatozoa and sample extraction (Tech)
 - 8.2.3.1. Cell pellet preparation
 - 8.2.3.2. Slide preparation
 - 8.2.3.3. Christmas tree staining, reagent preparation (if applicable)
 - 8.2.3.4. Sperm search, tails vs. no tails
 - 8.2.3.5. Sperm identification and epithelial cell identification, familiarity with commonly encountered organisms and substances (e.g., bacteria, yeast, lubricant, etc.)
 - 8.2.3.6. Interpretation and conclusions
- 8.2.4. p30 protein (Tech)

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- 8.2.4.1. Abacus OneStep p30 cards, biochemistry of reaction
 - 8.2.4.1.1. Sample preparation
 - 8.2.4.1.2. Controls
 - 8.2.4.1.3. Specificity and sensitivity
 - 8.2.4.1.4. High-dose hook effect
 - 8.2.4.1.5. False positives
 - 8.2.4.1.6. Interpretation and conclusions
- 8.2.5. Documentation
 - 8.2.5.1. Slide disposition
 - 8.2.5.2. Sperm search observations (including p30 and AP)
 - 8.2.5.3. Documentation of controls

8.3. Assessment

Create and test dilutions of semen up to 1:1000. Use a variety of substrates (clothing/fabric and swabs) with a variety of stains (e.g., semen, urine, vaginal secretions, etc.) using a combination of ALS, acid phosphatase reagent (spot test and mapping), microscopic examination for sperm, and Abacus OneStep p30 cards, as appropriate. Practice sets must include samples of varying strengths and mixtures of bodily fluids to ensure ability to determine differences between positive, negative, and inconclusive results, specifically when utilizing card-based testing methods.

A minimum of five satisfactory Christmas tree-stained slides must be prepared from mixed body fluids (e.g., semen/vaginal secretions, semen/saliva, etc. at various dilutions). Examine slides from various species to compare and contrast spermatozoa morphology.

COMPETENCY: A minimum of ten correctly characterized samples with the properly reported conclusions (five stains for HT trainees). This competency may be performed in concert with competencies for other body fluids.

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MODULE 8 – Identification of Semen Evaluation Form

	Physical and chemical characteristics of semen
	Acid phosphatase (Tech)
	Identification of spermatozoa and sample extraction (Tech)
	p30 protein (Tech)
	Documentation
	Interpretation and conclusions
	Competency
The tra	ainee has completed the above checked sections and is able to: Describe the physical and chemical characteristics of semen and the morphology of spermatozoa Test evidence items either directly or with a mapping technique to determine the location of possible semen stains by detecting acid phosphatase Produce a cell pellet, prepare a slide, stain the slide and positively identify spermatozoa under a microscope. Characterize other material that may be present on the slide Identify possible semen with the use of a p30 card
Comm	nents:
_	
٦	Trainee Printed Name + Initials Date Trainer Printed Name + Initials Date

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MODULE 8 IDENTIFICATION OF SEMEN READING ASSIGNMENTS

Reference	INITIALS	DATE
Baechtel F. 1988. Forensic Science Handbook. The Identification and Individualization of		
Semen Stains. Englewood Cliffs (NJ): Prentice Hall. 347-368. HT		
Gaensslen RE. 1983. Sourcebook in Forensic Serology, Immunology, and Biochemistry.		
Washington, D.C.: U.S. Department of Justice. Section 10.3 Seminal (Prostatic)		
Acid Phosphatase and Vaginal Acid Phosphatase: 155-169.		
Serological Research Institute. 2002. Serological Research Institute Methods Manual. Doc. MM I-B, rev.2. The Brentamine Reaction. 1-3.		
WSP CLD Biochemical Analysis Procedures: Acid Phosphatase Test Tech		
Enos WF, Beyer JC. Spermatozoa in the anal canal and rectum and in the oral cavity of		
female rape victims. J Forensic Sci 1978; 23(1): 231-233. HT		
Davies A, Wilson E. The persistence of seminal constituents in the human vagina. Forensic Sci 1974; 3(1): 45-55. HT		
Willott GM, Allard JE. Spermatozoa – their persistence after sexual intercourse. Forensic Sci Int 1982; 19(2): 135-154. HT		
Allard JE. The collection of data from findings in cases of sexual assault and the		
significance of spermatozoa on vaginal, anal and oral swabs. Sci Justice 1997; 37(2): 99-108. HT		
Joshi UN, et al. Effect of water immersion on seminal stains on cotton cloth. Forensic Sci		
Int 1981; 17(1): 9-11. HT		
Kafarowski E, et al. The retention and transfer of spermatozoa in clothing by machine		
washing. Canadian Society of Forensic Sci J 1996; 29(1): 7-11. HT		
Oppitz E. A new staining method for the detection of sperm in sexual offenses. Arkiv Fur Krimin 1969; 144: 145-148. HT		
Serological Research Institute. 2002. Serological Research Institute Methods Manual. Doc. MM II-C, rev.2. A Gram Modified Christmas Tree Stain. 1-2. HT		
WSP CLD Biochemical Analysis Procedures: Sperm Search/Christmas Tree Stain HT		
Tech		
Sensabaugh GF. Isolation and characterization of a semen-specific protein from human		
seminal plasma: A potential new marker for semen identification. J Forensic Sci 1978; 23(1): 106-115. HT		
Denison SJ, et al. Positive prostate-specific antigen (PSA) results in semen-free		
samples. Canadian Society of Forensic Sci J 2004; 37(4): 197-206. HT		
Seratec®. PSA in Body Fluids. 1-19. HT		
Sippel H, Lunetta P. Positive prostate-specific antigen (PSA) reaction in rectal samples		
from deceased males. Promega International Symposium on Human		
Identification, October 2004. HT		
Lunetta P, Sippel H. Positive prostate-specific antigen (PSA) reaction in post-mortem		
rectal swabs: A cautionary note. J Forensic Leg Med 2009; 16(7): 397-399. HT		
Hochmeister MN, et al. Evaluation of prostate-specific antigen (PSA) membrane test		
assays for the forensic identification of seminal fluid. J Forensic Sci 1999; 44(5): 1057-1060. HT		
Abacus Diagnostics p30 card product insert HT Tech		
WSP CLD Biochemical Analysis Procedures: Abacard® p30 Semen Test HT Tech		

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9. Evaluation of Apparent Hairs

The purpose of this module is to teach the trainee how to evaluate root material on apparent hairs for the suitability of DNA analysis.

9.1. Goals

At the completion of this module, the trainee should be:

- 9.1.1. Familiar with differences between a fiber, an animal hair, or a human hair.
- 9.1.2. Able to determine if an apparent human hair is suitable for DNA analysis.
- 9.1.3. Able to determine when to consult a Materials Analysis hair analyst.

9.2. Tasks

Instruction, demonstration, and practical training will be provided by a qualified Materials Analysis hair analyst.

- 9.2.1. Relevance of examination in casework
 - 9.2.1.1. Transference theory
 - 9.2.1.2. Persistence theory
- 9.2.2. Examination methods
 - 9.2.2.1. Handling loose hairs
 - 9.2.2.2. Handling hairs on sticky notes
 - 9.2.2.3. Stereomicroscope
 - 9.2.2.4. Illumination source and direction
 - 9.2.2.5. Background color and material
 - 9.2.2.6. Oblique light test
 - 9.2.2.7. Phase contrast microscope
 - 9.2.2.8. Glass slide preparation
- 9.2.3. Morphology of hairs and fibers
 - 9.2.3.1. Hair anatomy
 - 9.2.3.2. Human hair growth stages
 - 9.2.3.3. Tissue on human hairs
 - 9.2.3.4. Visual and microscopic features of strands
 - 9.2.3.4.1. Features used for categorization
 - 9.2.3.4.1.1 Human hair
 - 9.2.3.4.1.2 Animal hair
 - 9.2.3.4.1.3 Hair fragment
 - 9.2.3.4.1.4 Fiber
 - 9.2.3.4.1.5 Other
 - 9.2.3.4.2 How to identify these features
 - 9.2.3.4.2.1 Side view length, width, general form, surface, color
 - 9.2.3.4.2.2. End of view cross sectional shape
 - 9.2.3.4.2.3. Interior features delustrant, pigmentation, layers
 - 9.2.3.5. Features for recommendation of DNA analysis
- 9.2.4. When to consult a Material Analysis hair analyst

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recovery
Tape lifts
Vacuumings
Large items
Large quantity of hairs
tigative information – case scenario
Postmortem roots (necrotic, putrid)
Dyed/bleached hairs
Hair diseases
Somatic origin
Forcibly removed hairs
Acquired damage (crushed, burned, insect, fungal)
ssment difficulty
Damage to features
Atypical features
graphy
of known animal hairs

9.3. Assessment

Examine visually and with a stereomicroscope a set of known samples (loose and adhered to sticky notes). Samples must include human hairs, dog hairs, cat hairs, manufactured clothing fibers, manufactured carpet fibers, cotton fibers, and wool fibers. Perform an oblique light test on a dark, heavily delustered fiber and a dark, heavily pigmented human hair. Prepare a temporarily mounted human hair from a sticky note and observe with phase contrast microscopy. Examine known human hairs mounted in Permount slides with phase contrast microscopy and evaluate their roots. Practice identification of strands on a set of unknowns provided by your trainer using any of the methods taught.

COMPETENCY: The trainee will examine a sample set consisting of at least 10 hairs and fibers. For each sample, the trainee will identify whether the sample is suitable for STR DNA analysis.

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WSP DNA BIOCHEMISTRY AND STR TRAINING PROGRAM MODULE 9 – Evaluation of Apparent Hairs

Evaluation Form

Т	rainee Printed Name + Initials Date Trainer Printed Name + Initials Date
-	
Comm	ents:
The tra	ninee has completed the above checked sections and is: Familiar with differences between a fiber, an animal hair, or a human hair. Able to determine if an apparent human hair is suitable for DNA analysis. Able to determine when to consult a Materials Analysis hair analyst.
	Competency
	Interpretation and conclusions
	Documentation
	Practical examination of tape lifts from clothing and/or car seats
	Examination of known animal hairs
	Examination of known human head, pubic, and body hairs
	Examination of known fibers
	Illumination methods and using the stereomicroscope
	Theory and relevance of examination in casework

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MODULE 9 EVALUATION OF APPARENT HAIRS READING ASSIGNMENTS

TRAINEE:	

REFERENCE	INITIALS	DATE
Bisbing RE. 2002. Forensic Science Handbook. Volume 1, second edition. Upper Saddle River (NJ): Prentice Hall. The forensic identification and association of human hair. 392-393.		
Ogle RR, Fox MJ. 1999. Atlas of Human Hair – Microscopic Characteristics. CRC Press. Visual resource: Examine Chapter 5 Human hair microscopic characteristics: photographs and drawings of variate archetypes and examples: 37-48 and plates.		
Petraco N, Kubic T. 2003. Color Atlas and Manual of Microscopy for Criminalists, Chemists, and Conservators. New York: CRC Press. Visual resource: Examine images in Identification and Comparison of Human Hair: 57-67.		
Petraco N, Kubic T. 2003. Color Atlas and Manual of Microscopy for Criminalists, Chemists, and Conservators. New York: CRC Press. Visual resource: Examine images in Animal Hair Identification: 69-76.		
Petraco N, Kubic T. 2003. Color Atlas and Manual of Microscopy for Criminalists, Chemists, and Conservators. New York: CRC Press. Visual resource: Examine images in Appendix A: Human Hair Atlas: 217-237.		
Petraco N, Kubic T. 2003. Color Atlas and Manual of Microscopy for Criminalists, Chemists, and Conservators. New York: CRC Press. Visual resource: Examine images in Appendix B: Animal Hair Atlas: 239-255.		
Moore TD, et al. 1974. Identification of the Dorsal Guard Hair of Some Mammals of Wyoming. Cheyenne: Wyoming Game and Fish Department. Glossary: 3-17.		
Linch CA, et al. Evaluation of the human hair root for DNA typing subsequent to microscopic comparison. J Forensic Sci 1998; 43(2): 305-314.		
Pettenati MJ, Rao PN. Commentary on "Linch CA, Smith SL, Prahlow JA. Evaluation of the human hair root for DNA typing subsequent to microscopic comparison. J Forensic Sci. 1998; 43(2): 305-314". J Forensic Sci 1999; 44(6): 1329-1330.		
Dachs J, et al. The persistence of human scalp hair on clothing fabrics. Forensic Sci Int 2003; 138(1): 27-36.		
Chewning DD, et al. Persistence of fibers on ski masks during transit and processing. Forensic Sci Comm 2008; 10(3).		
Exline D. Frequency of pubic hair transfer during sexual intercourse. J Forensic Sci 1998; 43(3): 505-508.		
Gaudette BD, Tessarolo AA. Secondary transfer of human scalp hair. J Forensic Sci 1987; 32(5): 1241-1253.		
Siegel JA, Mirakovits K. 2010. Forensic Science: The Basics. Boca Raton (FL): CRC Press. Textile Fibers, Typical Fibers, and Fiber Morphology: 410-412.		
WSP CLD Biochemistry Analysis Procedures: Hair		

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10. Cellular DNA Collection Techniques

The purpose of this module is to teach the knowledge and techniques required for successful collection of cellular DNA from various items.

10.1. Goals

At the completion of this module, the trainee should be able to:

- 10.1.1. Identify the locations on a given item that will yield the most successful DNA results.
- 10.1.2. Demonstrate proper documentation of the collection process and subsequent sample.

10.2. Tasks

Instruction, demonstration and practical training will be provided

- 10.2.1. Observation of qualified analysts collecting cellular material (including the M-Vac, if applicable)
 - 10.2.1.1. Documentation
 - 10.2.1.2. Collection considerations
 - 10.2.1.3. Proper labeling
 - 10.2.1.4. Storage/preservation of collected samples prior to DNA analysis
- 10.2.2. Review of 3-5 completed case files with cellular collection (including the M-Vac, if applicable)
 - 10.2.2.1. Documentation
 - 10.2.2.2. Sampling techniques
 - 10.2.2.3. Sampling areas
- 10.2.3. Hands-on training in DNA collection from various sources and observation of the trainee's techniques (including the M-Vac, if applicable)

10.3. Assessment

COMPETENCY: The trainee will use collection techniques on five mock items (including use of the M-Vac, if applicable), which will be subsequently analyzed by a qualified DNA analyst for evaluation. The trainer will document the successful use of cellular collection methods using the trainer's evaluation form.

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WSP DNA BIOCHEMISTRY AND STR TRAINING PROGRAM MODULE 10 – Cellular DNA Collection Techniques Evaluation Form

	Single and double-swab techniques	
	Safety of operation of the M-Vac	
	Procedure for examination of evidence	
	Documentation	
	Interpretation and conclusions	
	Competency	
The t	rainee has completed the above checked s Successfully use DNA collection techniq Properly operate the M-Vac	ections and is able to: ues to attain DNA profiles of item seen in casework
Comr	ments:	
		Trainer Printed Name + Initials Date
	Trained Traine + Illitials Date	

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MODULE 10 CELLULAR DNA COLLECTION TECHNIQUES READING ASSIGNMENTS

REFERENCE	INITIALS	DATE
Sweet D, et al. An improved method to recover saliva from human skin: the double swab		
technique. J Forensic Sci 1997; 42(2): 320-322. HT		
Butler JM. 2012. Advanced Topics in Forensic DNA Typing: Methodology. San Diego:		
Elsevier Academic Press. Chapter 1 Sample Collection, Storage, and		
Characterization: 1-19. HT		
Butler JM. 2012. Advanced Topics in Forensic DNA Typing: Methodology. San Diego:		
Elsevier Academic Press. Chapter 1 Sample Collection, Storage, and		
Characterization: 1-10. CODIS		
MSI M-Vac Systems® System. SEC Series 100 and 150 User Guide (if applicable)		
WSP CLD DNA internal validation for the M-Vac System® (if applicable)		
WSP CLD Biochemistry Analysis Procedures: MSI M-Vac System®		
WSP CLD DNA Analysis Casework STR Procedures Manual: Recovering Slide-Mounted		
Hairs or Semen Smears		

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11. Fundamental Scientific DNA Knowledge

The purpose of this module is to ensure the trainee has the formal education and understanding of the fundamental scientific basis of forensic DNA analysis as required by national standards (see FBI QAS Standards).

11.1. Goals

At the end of this session the trainee should be able to:

- 11.1.1. Document college level course work covering the fundamental principles of genetics, biochemistry, population genetics/statistics, and molecular biology.
- 11.1.2. Understand fundamental scientific knowledge as it applies to forensic DNA analysis.
- 11.1.3. Pass a written exam as part of Module 25 on in-depth knowledge appropriate to their duties.
- 11.1.4. Discuss forensic DNA topics in depth, appropriate to their duties.

11.2. Tasks (Tech)

- 11.2.1. All trainees must produce a curriculum vitae stating their education, work experience, and professional activities.
- 11.2.2. All trainees must also provide a copy of their college transcripts.

11.3. Assessment

College level coursework must have been successfully completed by the Casework DNA Analyst and CODIS DNA Analyst trainees in genetics, biochemistry, population genetics/statistics and molecular biology. The trainer will document the approval of the trainee's education by the DNA Technical Leader using the trainer's evaluation form.

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MODULE 11 – Fundamental Scientific Knowledge Evaluation Form

	Produce curriculum vitae stating	education, wo	rk experience, and professional activi	ities (Tech)
	Provide a copy of college transcr	ipts (Tech)		
The tra	foundation for understand	ng the fundame genetics/statis ding forensic D	ental principles of genetics, stics, and molecular biology which pro	
Comm	nents:			
<u>-</u>	Frainee Printed Name + Initials	 Date		Date

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MODULE 11 FUNDAMENTAL SCIENTIFIC DNA KNOWLEDGE READING ASSIGNMENTS

TRAINEE:	

REFERENCE	INITIALS	DATE
Butler JM. 2005. Forensic DNA Typing: Biology, Technology, and Genetics of STR		
Markers. New York: Elsevier Academic Press. Chapter 1 Overview and History of DNA Typing: 1-13.		
Butler JM. 2005. Forensic DNA Typing: Biology, Technology, and Genetics of STR Markers. New York: Elsevier Academic Press. Chapter 2 DNA Biology Review: 17-30.		
National Research Council Committee on DNA Forensic Science. 1992. DNA Technology in Forensic Science. Washington, D.C.: National Academies Press. Summary: 1-26.		
National Research Council Committee on DNA Forensic Science. 1996. The Evaluation of Forensic DNA Evidence. Washington, D.C.: National Academies Press. Executive Summary: 1-8 (compare to 1992 Summary).		
Shutler GG. 2005. Forensic Botany: Principles and Applications to Criminal Casework. Boca Raton: CRC Press. Chapter 8 An Overview of Historical Developments in Forensic DNA Analysis: 117-135.		
Gill P, et al. Forensic application of DNA 'fingerprints'. Nature 1985; 318(6046): 577-579.		

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Applied Scientific DNA Knowledge and Lab Work

The purpose of this module is to provide practical instruction to the trainee on the analytical procedures used in the laboratory. This module builds on the foundation of the fundamental scientific knowledge relating to the study of forensic DNA analysis and includes lab work using this applied knowledge. The Casework DNA Analyst trainee will perform analysis on biological samples that would be normally encountered in forensic casework. The CODIS DNA Analyst trainee will perform analysis on reference samples normally encountered in convicted offender submissions. The methods detailed in the WSP CLD Casework STR Analysis Procedures manual or the WSP CLD CODIS Laboratory STR Analysis Procedures manual, as appropriate, will be employed.

12.1. Goals

At the end of this session the trainee should be able to:

- 12.1.1. Discuss forensic DNA topics in depth, appropriate to their duties.
- 12.1.2. Competently perform DNA STR analysis on biological samples similar to what would be encountered in forensic DNA casework or convicted offender samples.
- 12.1.3. Demonstrate good laboratory technique for DNA STR analysis.

12.2. Tasks

There will be instruction and demonstration of the procedures that relate to the trainee's work place duties.

- 12.2.1. Trainers will discuss with trainees subject matter and published references (Tech)
 - 12.2.1.1. Casework Direct
 - 12.2.1.2. DNA extraction and purification (organic and EZ1)
 - 12.2.1.3. DNA quantification
 - 12.2.1.4. Polymerase chain reaction (PCR) based DNA typing methodology.
 - 12.2.1.5. Direct amplification (DA)
 - 12.2.1.6. Short tandem repeat polymorphisms
- 12.2.2. Casework DNA Analyst trainees will be assigned a number of samples sufficient to demonstrate their ability to competently conduct the laboratory's analytical procedures and produce reliable and accurate results. The following is a typical assignment: at least 25 single source samples followed by ten single source competency samples, at least seven samples for differential extraction and analysis, five contact/touch DNA samples (e.g., for wearer DNA), ten hair samples, and three non-probative cases. These three non-probative cases may serve as the competency/mock cases for Module 23. These samples will reflect the variability, range, type, and complexity of casework analysis.
 - 12.2.2.1. Samples will be processed using organic extraction, Qiagen EZ1 robotic protocols, STARlet and/or AutoLys robotic protocols. Assignment of samples for use with direct amplification of reference samples is optional. If direct amplification of reference samples is used for some of the single source samples, at least ten will be done using this procedure
- 12.2.3. CODIS DNA Analyst trainees will be assigned

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- 12.2.3.1. A practice set of five samples to be processed manually via extraction in the laboratory under direct observation of the trainer
- 12.2.3.2. A GeneMapper® ID-X data set containing different types of contamination
- 12.2.3.3. At least two training sets of ten samples (eight buccal and two blood) to process manually via extraction
- 12.2.3.4. At least two training sets of ten samples to process manually via direct amplification
- 12.2.3.5. At least two 96-well plate training sets of about 40 samples each to process with the BSD600 Duet/Ascent Puncher and direct amplification
- 12.2.3.6. CODIS DNA Analyst trainees will also be assigned five manual competency samples (extraction), five manual competency samples (direct amplification), and a set of 30 samples to process using the BSD600 Duet/Ascent puncher and direct amplification
- 12.2.4. The following materials are available for further study should the trainer or trainee deem additional practice is necessary
 - 12.2.4.1. GeneMapper® ID-X data sets for data analysis practice
 - 12.2.4.2. PowerQuant runs for standard curve and/or quantification value evaluation
 - 12.2.4.3. Example case files for worksheets and workflow practice
- 12.2.5. Laboratory analysis is to be performed for:
 - 12.2.5.1. DNA extraction (lysis) and purification
 - 12.2.5.2. DNA quantification
 - 12.2.5.3. Polymerase chain reaction (PCR) based DNA typing methodology
 - 12.2.5.4. Short tandem repeat DNA typing profiles
- 12.2.6. The trainer will discuss with the trainee the Sample Switch Detection Procedure

12.3. Assessment

All trainees must be able to generate reliable genotype data in a proficient manner. The trainer will document the achievement of the trainee's lab work using the trainer's evaluation form.

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Riochomi	cal and STP	Training Program	Manual

WSP DNA BIOCHEMISTRY AND STR TRAINING PROGRAM MODULE 12— Applied Scientific DNA Knowledge Evaluation Form

	blank for HT) (7 samples and a reagent blank done using automation, if applicable)
	DNA extraction and purification
	DNA quantification
	Polymerase chain reaction
	Direct amplification
	Short tandem repeat polymorphisms
Casa	work DNA Analyst
	Single source stains (≥25 – if using DA, at least 10 will be done using DA) (20 for HT) (5 using the EZ1 and 5 DA for Tech)
	Differential extraction and analysis (≥7) (20 for HT) (2 for Tech)
	Contact/touch (3)
	Hair (10)
	Non-probative cases (3) (HT)
CODI	S DNA Analyst
	Manual extraction of practice samples (5)
	Contamination data set
	Manual extraction of training samples (2 sets of 10)
	Direct amplification of training samples (2 sets of 10)
	96-well plate processing of training samples (2 sets of ~40)

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The trainee has completed the above checked sections and is able to:

Discuss and display an in-depth knowledge appropriate to their duties (Tech)

Prepare for laboratory analysis work assignments

Competently perform PCR STR analysis on biological samples similar to what would be encountered in forensic DNA casework or forensic databases

Demonstrate good laboratory technique for PCR STR analysis

Operate the following instruments: general laboratory equipment and instruments associated with the procedures used in STR analysis such as autoclaves, heat blocks, pipettes, vortex mixers, centrifuges, etc.; real-time PCR instruments; thermal cyclers; genetic analyzers; applicable robots (EZ1, QIAgility, STARlet, AutoLys, BSD600 Duet/Ascent Puncher, etc.) (Tech)

Comments:			
Trainee Printed Name + Initials	Date	Trainer Printed Name + Initials	 Date

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MODULE 12 APPLIED SCIENTIFIC DNA KNOWLEDGE AND LAB WORK READING

ASSIGNMENTS TRAINEE:	

REFERENCE	INITIALS	DATE
WSP CLD DNA Casework STR Analysis Procedures Manual: Introduction HT Tech		
WSP CLD CODIS Laboratory STR Analysis Procedure Manual Introduction, Fusion 6C Kit Loci, and CODIS Case Approach CODIS only		
Y-screening		
Graham EK, et al. Developmental validation of the casework direct kit, custom: a method for the rapid processing of casework samples. 2018. HT		
Promega. 2019. Casework Direct Kit application note. AN300. HT		
WSP CLD DNA internal validation summary for the Promega Casework Direct Kit HT		
WSP CLD DNA lab-specific internal validation summary for the Promega Casework Direct Kit HT		
WSP CLD DNA Casework STR Analysis Procedures Manual: Y-Screening for Sexual Assault Evidence HT Tech		
WSP CLD DNA Internal Validation Summary for the Hamilton Microlab AutoLys STAR for Y-Screening and Quantification Set-up (if applicable)		
WSP CLD DNA Casework STR Analysis Procedures Manual: Automated Sample Processing Using the Hamilton AutoLys (if applicable)		
Extraction		
Butler JM. 2012. Advanced Topics in Forensic DNA Typing: Methodology. San Diego:		
Elsevier Academic Press. Chapter 2 DNA Extraction Methods: 29-40. HT		
WSP CLD DNA Casework STR Analysis Procedures Manual: Non-Differential Lysis		
Using WSP Buffer HT Tech WSP CLD DNA Casework STR Analysis Procedures Manual: Differential Lysis		
Procedure for Semen Stains HT Tech		
WSP CLD DNA Casework STR Analysis Procedures Manual: Lysis: Hair Samples Using		
WSP Buffer		
WSP CLD DNA Casework STR Analysis Procedures Manual: Qiagen EZ1 Pretreatment Protocols Using G2 Buffer HT Tech		
Purification	<u>_</u>	
Shutler GG, et al. Removal of a PCR inhibitor and resolution of DNA STR types in mixed		
human-canine stains from a five year old case. J Forensic Sci 1999; 44(3): 623-626. HT		
Montpetit SA, et al. A simple automated instrument for DNA extraction in forensic		
casework. J Forensic Sci 2005; 50(3): 555-563. HT		
Anslinger K, et al. Application of the BioRobot® EZ1 in a forensic laboratory. Leg Med 2005; 7(3): 164-168. HT		
Kishore R, et al. Optimization of DNA extraction from low-yield and degraded samples		
using the BioRobot® EZ1 and BioRobot® M48. J Forensic Sci 2006; 51(5): 1055-1061. HT		
Qiagen. EZ1 DNA Investigator® handbook CODIS HT Tech		
Promega. 2012. DNA IQ™ System technical manual: small sample casework protocol. TB296. HT		
Promega. 2012. DNA IQ™ System technical manual: database sample protocol. TB297. HT		
WSP CLD DNA internal validation summary for the EZ1 HT		
WSP CLD CODIS lab validation summary for the EZ1 CODIS only		

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WSP CLD CODIS lab supplemental internal validation for the EZ1 Advanced XL CODIS	
only	
WSP CLD DNA Casework STR Analysis Procedures Manual: Qiagen BioRobot® EZ1 and EZ2 Connect Workstations HT Tech	
WSP CLD DNA Casework STR Analysis Procedures Manual: Purification: EZ1 Trace Method HT Tech	
WSP CLD DNA Casework STR Analysis Procedures Manual: Purification: EZ2 Connect Trace Method HT Tech	
WSP CLD DNA Casework STR Analysis Procedures Manual: Purification: EZ1 Large Volume Method HT Tech	
WSP CLD DNA Casework STR Analysis Procedures Manual: Purification: EZ2 Connect Large Volume Method HT Tech	
WSP CLD DNA Casework STR Analysis Procedures Manual: Purification: EZ1 Tip Dance Method HT Tech	
WSP CLD DNA Casework STR Analysis Procedures Manual: Purification: EZ2 Connect Tip Dance Method HT Tech	
BSD600 DUET automated punch instrument user manual CODIS only	
WSP CLD CODIS Laboratory STR Analysis Procedures Manual: BDS600 Duet Puncher Protocol CODIS only	
BSD600 Ascent automated punch instrument user manual CODIS only	
WSP CLD CODIS Laboratory STR Analysis Procedures Manual: BSD600 Ascent Puncher Protocol CODIS only	
WSP CLD CODIS Laboratory STR Analysis Procedures Manual: EZ1 DNA Investigator® Kit Extraction CODIS only	
Quantification	
Butler JM. 2012. Advanced Topics in Forensic DNA Typing: Methodology. San Diego: Elsevier Academic Press. Chapter 3 DNA Quantitation: 49-64. HT	
Ewing MM, et al. Human DNA quantification and sample quality assessment:	
developmental validation of the PowerQuant® system. Forensic Sci Int 2016; 23: 166-177. HT	
Promega. 2020. PowerQuant® System technical manual. TMD047. HT Tech	
Bode and WSP supplemental validation summary reports and lab binders for the PowerQuant® system HT	
WSP CLD DNA Casework STR Analysis Procedures Manual: DNA Quantification: Standards Preparation PowerQuant® System HT Tech	
WSP CLD DNA Casework STR Analysis Procedures Manual: DNA Quantification: Sample Set-Up PowerQuant® System = HT Tech	
WSP CLD DNA Casework STR Analysis Procedures Manual: DNA Quantification: PowerQuant® System HT Tech	
WSP CLD DNA Casework STR Analysis Procedures Manual: DNA Quantification: PowerQuant® System Data Analysis HT Tech	
WSP CLD DNA Casework STR Analysis Procedures Manual: DNA Quantification: PowerQuant® System Data Interpretation HT Tech	
WSP CLD CODIS lab performance verification reports for the AB 7500 Sequence Detection system CODIS only	
Green RL, et al. Developmental validation of the Quantifiler™ real-time PCR kits for the quantification of human nuclear DNA samples. J Forensic Sci 2005; 50(4): 809-825. CODIS only	
Applied Biosystems. Quantifiler™ user manual CODIS only	
WSP CLD CODIS Laboratory STR Analysis Procedures Manual: Quantifiler™ Template	
Setup CODIS only	

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WSP CLD CODIS Laboratory STR Analysis Procedures Manual: Quantifiler™ Data Analysis and Interpretation CODIS only	
Concentration	
Millipore. 2005. Concentrating and desalting DNA or RNA with Microcon or Centricon centrifugal filters. HT	
WSP CLD DNA Casework STR Analysis Procedures Manual: Microcon Concentration of DNA HT	
WSP CLD DNA Casework STR Analysis Procedures Manual: Vacufuge Procedure – Concentration, Preservation, and Recovery of DNA Extracts/Work Product HT Tech	
WSP CLD CODIS Laboratory STR Analysis Procedures Manual: Microcon® Concentration of DNA CODIS only	
Amplification	
Saiki RK, et al. Primer-directed enzymatic amplification of DNA with a thermostable DNA polymerase. Science 1988; 239(4839): 487-491. HT	
Butler JM. 2012. Advanced Topics in Forensic DNA Typing: Methodology. San Diego: Elsevier Academic Press. Chapter 4 PCR Amplification: 69-91. HT	
Butler JM. 2012. Advanced Topics in Forensic DNA Typing: Methodology. San Diego:	
Elsevier Academic Press. Chapter 5 Short Tandem Repeat (STR) Loci and Kits: 99-132. HT	
Butler JM. 2005. Forensic DNA Typing: Biology, Technology, and Genetics of STR	
Markers. New York: Elsevier Academic Press. Chapter 7 Forensic Issues: 152-154 only. HT	
WSP CLD DNA summary for the evaluation of expanded loci amplification kits HT	
Ensenberger MG, et al. Developmental validation of the PowerPlex® Fusion 6C system. Forensic Sci Int Genet 2016; 21: 134-144. HT Tech	
Promega. PowerPlex® Fusion 6C System technical manual. TMD0045. HT	
WSP CLD DNA internal validation summary for the PowerPlex® Fusion 6C system HT	
WSP CLD DNA lab-specific internal validation summary for the PowerPlex® Fusion 6C system HT	
WSP CLD DNA Casework STR Analysis Procedures Manual: Amplification: Sample Set- Up PowerPlex® Fusion 6C	
WSP CLD DNA Casework STR Analysis Procedures Manual: Amplification: PowerPlex® Fusion 6C	
WSP CLD DNA Casework STR Analysis Procedures Manual: Direct Amplification Set-Up: PowerPlex® Fusion 6C	
WSP CLD DNA Casework STR Analysis Procedures Manual: Direct Amplification: PowerPlex® Fusion 6C HT Tech	
Promega. 2016. SwabSolution™ Kit technical manual. TMD037. HT Tech	
WSP CLD DNA validation of buccal cotton swab direct amplification using the	
PowerPlex® Fusion 6C system and 3500 genetic analyzer HT	
WSP CLD DNA Marysville lab PowerPlex® Fusion 6C direct amplification of DNA reference samples supplemental validation HT	
Federal Bureau of Investigation. Quality assurance standards for forensic DNA testing laboratories. Standard 9.4 including discussion. Current version. HT	
Promega. PunchSolution™ Kit technical manual. TMD038. CODIS only	
WSP CLD CODIS Laboratory STR Analysis Procedures Manual: Fusion 6C Direct Amplification CODIS only	
WSP CLD CODIS Laboratory STR Analysis Procedures Manual: Fusion 6C Extract	
Amplification CODIS only Capillary Electrophoresis/Data Collection	
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Washington State Patrol – Crime Laboratory Division Biochemical and STR Training Program Manual Butler JM. 2012. Advanced Topics in Forensic DNA Typing: Methodology. San Diego: Elsevier Academic Press. Chapter 6 Capillary Electrophoresis: 141-162. HT Butler JM. 2015. Advanced Topics in Forensic DNA Typing: Interpretation. San Diego: Elsevier Academic Press. Chapter 2 Data, Models, and Thresholds: 25-44. HT	
Butler JM. 2012. Advanced Topics in Forensic DNA Typing: Methodology. San Diego: Elsevier Academic Press. Chapter 6 Capillary Electrophoresis: 141-162. HT Butler JM. 2015. Advanced Topics in Forensic DNA Typing: Interpretation. San Diego:	
Elsevier Academic Press. Chapter 6 Capillary Electrophoresis: 141-162. HT Butler JM. 2015. Advanced Topics in Forensic DNA Typing: Interpretation. San Diego:	
Butler JM. 2015. Advanced Topics in Forensic DNA Typing: Interpretation. San Diego:	
Elsevier Academic Press. Chapter 2 Data, Models, and Thresholds: 25-44. HT	
Butler JM. 2015. Advanced Topics in Forensic DNA Typing: Interpretation. San Diego:	
Elsevier Academic Press. Chapter 3 STR Alleles and Amplification Artifacts: 47- 68 only.	
HT	
Butler JM. 2015. Advanced Topics in Forensic DNA Typing: Interpretation. San Diego:	
Elsevier Academic Press. Chapter 8 Troubleshooting Data Collection: 183-207.	
HT	
Applied Biosystems. 3500/3500xL user manual, product enclosures and/or manual (as	
applicable) HT Tech	
Applied Biosystems. 2012. Technical Note. Considerations for evaluating carryover on	
Applied Biosystems capillary electrophoresis platforms in a HID laboratory. HT	
VSP CLD DNA validation summary for direct amplification using the PowerPlex® Fusion	
6C system and 3500 genetic analyzer HT VSP CLD DNA Casework STR Analysis Procedures Manual: Amplification Product	
Preparation: Fusion 6C HT Tech	
WSP CLD CODIS lab internal validation reports for the 3500xL CODIS only	
Bode validation of direct amplification using the PowerPlex® Fusion 6C system, 3500xL	
genetic analyzer, and BSD600 DUET automated punch instrument. 2016. CODIS	
only	
VSP CLD CODIS Laboratory STR Analysis Procedures Manual: Amplification Product	
Preparation for the 3500XL CODIS only	
VSP CLD CODIS Laboratory STR Analysis Procedures Manual: Running Plates on the	
3500XL Genetic Analyzer CODIS only	
DNA Analysis of Various Sample Types	
Vegel JG, Jr, Herrin G, Jr. Deduction of the order of sexual assaults by DNA analysis of	
two condoms. J Forensic Sci 1994; 39(3): 844-846. HT	
Viegand P, Kleiber M. DNA typing of epithelial cells after strangulation. Int J Legal Med	
1997; 110(4): 181-183. HT	
orente M, et al. Dandruff as a potential source of DNA in forensic casework. J Forensic	
Sci 1998; 43(4): 901-902.	
Sweet D, Shutler GG. Analysis of salivary DNA evidence from a bite mark on a body	
submerged in water. J Forensic Sci 1999; 44(5): 1069-1072. HT	
Abaz J, et al. Comparison of the variables affecting the recovery of DNA from common	
drinking containers. Forensic Sci Int 2002; 126(3): 233-240.	
Primorac D. The role of DNA technology in identification of skeletal remains discovered	
in mass graves. Forensic Sci Int Supp Ser 2004. 146S: S163-S164.	

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13. DNA Interpretation and STRmix

The purpose of this module is to provide practical instruction on how to interpret and report analytical results as designated by laboratory policy. The Casework DNA Analyst trainee will receive instruction on how to use GeneMapper™ ID-X, STR interpretation guidelines, the interpretation of mixtures, the use of the STRmix™ program and evaluation of results for intuitiveness, STRlite, basic report writing, and wording of conclusions. A brief introduction to the laboratory's CODIS program will also be provided. There will be a set of mixture data (~ 20 mixtures to demonstrate competency) representative of casework provided to the Casework DNA analyst trainee on which to conduct interpretation according to laboratory policy. The Casework DNA analyst trainee will document appropriate manual evaluation of a subset of the provided mixtures and demonstrate correlation with STRmix results.

13.1. Goals

At the end of this session the trainee should be able to:

13.1.1. Utilize GeneMapper™ ID-X to correctly interpret casework STR data, deconvolute DNA profiles using STRmix™ and correlate these results with manual interpretation of the data, use STRlite to summarize STRmix™ data, and write interpretation results in accordance with laboratory policy

13.2. Tasks

Instruction, demonstration, and practical training will be provided

- 13.2.1. GeneMapper™ ID-X set-up and use (CODIS) (Tech)
- 13.2.2. STR interpretation guidelines (Tech)
- 13.2.3. Use of the STRmix[™] program for deconvolution
- 13.2.4. Use of the STRlite workbook
- 13.2.5. Deducing profiles for CODIS, MME, and a brief overview of CODIS eligibility
- 13.2.6. Wording of interpretation conclusions in reporting of results
- 13.2.7. Work assigned to complete
 - 13.2.7.1. Interpretation of ~20 sets of mixture data representative of casework will be assigned
 - 13.2.7.2. The Casework DNA Analyst trainee is to provide a written interpretation according to laboratory policy
- 13.2.8. The trainee must complete paperwork to be approved for CODIS access.
- 13.2.9. CODIS DNA Analyst trainees will be assigned
 - 13.2.9.1. A GeneMapper™ ID-X training data set
 - 13.2.9.2. A GeneMapper™ ID-X training data set composed of anomalies

13.3. Assessment

Interpretation results and associated paperwork of the ~20 mixtures will be evaluated by experienced Casework DNA STR analysts. The trainer will document the completion of the module using the trainer's evaluation form. The practical portions of modules 16 and 18 should be worked on in tandem with the mixtures to be completed for this module.

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WSP DNA BIOCHEMISTRY AND STR TRAINING PROGRAM MODULE 13 – DNA Interpretation and STRmix Evaluation Form

Ш	GeneMapper™ ID-X (CODIS) (Tech)
	STR interpretation guidelines (Tech)
	Use of STRmix™ and deconvolution
	Interpretation of mixture data sets (~20)
	Trainees fill out paperwork for CODIS access
COD	DIS DNA Analyst
	GeneMapper® ID-X data training set
	GeneMapper® ID-X advanced data training set (anomalies)
The t	trainee has completed the above checked sections and is able to: Correctly interpret casework STR data compatible with laboratory policy Use STRmix™ and STRlite to assist in deconvolution and determining CODIS eligibility
Comi	ments:
	Trainee Printed Name + Initials Date Trainer Printed Name + Initials Date

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MODULE 13 DNA INTERPRETATION AND STRMIX READING ASSIGNMENTS

TRAINEE:	

REFERENCE	INITIALS	DATE
WSP CLD DNA Casework STR Analysis Procedures Manual: GeneMapper™ ID-X Setup		
Tech		
WSP CLD DNA Analysis Quality Assurance Manual: Appendix I – Extraneous DNA		
Guidelines		
WSP CLD DNA Analysis Quality Assurance Manual: Appendix II – Extraneous DNA		
Guidelines – CODIS CODIS only		
Meldgaard M, Morling N. Detection and quantitative characterization of artificial extra		
peaks following polymerase chain reaction amplification of 14 short tandem repeat		
systems used in forensic investigations. Electrophoresis 1997; 18(11): 1928-1935.		
Walsh PS, et al. Sequence analysis and characterization of stutter products at the		
tetranucleotide repeat locus vWA. Nucleic Acids Research 1996; 24(14): 2807-		
2812.		
Butler JM. 2015. Advanced Topics in Forensic DNA Typing: Interpretation. San Diego:		
Elsevier Academic Press. Chapter 3 STR Alleles and Amplification Artifacts: 68-		
80 only.		
Rolf B, et al. Somatic mutations at STR loci – a reason for three-allele pattern and		
mosaicism. Forensic Sci Int 2002; 126(3): 200-202.		
Hendrickson BC, et al. Accurate STR allele designations at the FGA and vWA loci		
despite primer site polymorphisms. J Forensic Sci 2004; 49(2); 250-254.		
Jiang W, et al. Identification of dual false indirect exclusions on the D5S818 and FGA		
loci. Leg Med 2011; 13(1): 30-34.		
Butler JM. 2015. Advanced Topics in Forensic DNA Typing: Interpretation. San Diego:		
Elsevier Academic Press. Chapter 1 Data Interpretation Overview: 3-22.		
Butler JM. 2015. Advanced Topics in Forensic DNA Typing: Interpretation. San Diego:		
Elsevier Academic Press. Chapter 4 STR Genotypes: 87-105.		
Butler JM. 2015. Advanced Topics in Forensic DNA Typing: Interpretation. San Diego:		
Elsevier Academic Press. Chapter 5 STR Profiles: 109-123.		
Butler JM. 2012. Advanced Topics in Forensic DNA Typing: Methodology. San Diego:		
Elsevier Academic Press. Chapter 10 Degraded DNA: 293-304.		
Applied Biosystems. 2009. Investigations to assist in the interpretation of DNA profiles.		
Butler JM. 2012. Advanced Topics in Forensic DNA Typing: Methodology. San Diego:		
Elsevier Academic Press. Chapter 11 Low-Level DNA Testing: 311-341.		
Butler JM. 2015. Advanced Topics in Forensic DNA Typing: Interpretation. San Diego:		
Elsevier Academic Press. Chapter 6 DNA Mixtures: 129-154.		
Butler JM. 2015. Advanced Topics in Forensic DNA Typing: Interpretation. San Diego: Elsevier Academic Press. Chapter 7 Low-Level DNA and Complex Mixtures: 159-		
· ·		
177. Butler JM. 2015. Advanced Topics in Forensic DNA Typing: Interpretation. San Diego:		
Elsevier Academic Press. Appendix 4 Worked Mixture Example: 537-548 only.		
Butler JM, et al. NIST interlaboratory studies involving DNA mixtures (MIX05 and		
MIX13): Variation observed and lessons learned. Forensic Sci Int Genet 2018; 37:		
81-94.		
WSP CLD DNA Casework STR Analysis Procedures Manual: Guidelines for Evaluating		
DNA Profile Data Tech		

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WSP CLD DNA Casework STR Analysis Procedures Manual: Interpretation of STR Profiles	
WSP CLD DNA Casework STR Analysis Procedures Manual: Kinship Analysis	
Perlin MW, Szabady B. Linear mixture analysis: a mathematical approach to resolving mixed DNA samples. J Forensic Sci 2001; 46(6): 1372-1378.	
Bright J, et al. Developmental validation of STRmix expert software for the interpretation of forensic DNA profiles. Forensic Sci Int Genet 2016; 23: 226-239.	
STRmix™ support solutions. 2018. Why do I have unintuitive genotypes at vWA (involving allele 14)?	
WSP CLD DNA estimation of STRmix™ parameters for Fusion 6C within STRmix™ v2.5	
WSP CLD DNA internal validation summary for STRmix™. Presentations at the DNA Functional Area Meeting. Seattle. 2018.	
WSP CLD DNA internal validation summary for STRmix™ v2.8 probabilistic genotyping software for PowerPlex® Fusion 6C and the 3500 genetic analyzer	
WSP Estimation of STRmix v2.8 Parameters for the Washington State Patrol Crime Laboratory Division	
WSP CLD DNA Casework STR Procedures Manual: Probabilistic Genotyping Using STRmix™	
Wagner M. Simplifying the STRmix™ workflow with a custom software solution. Poster presentation at the International Symposium on Human Identification, Phoenix. 2018.	
WSP CLD DNA STRlite performance check plan and performance check summary	
WSP CLD DNA STRlite PowerPoint and demo. Presentation at the DNA Functional Area Meeting. Seattle. 2019.	
WSP CLD DNA Casework STR Analysis Procedures Manual: STRlite for STRmix 2.8	
WSP CLD CODIS Laboratory STR Analysis Procedures Manual: Evaluating Fusion 6C profiles with GeneMapper ID-X v1.6.2® as an Analysis ToolCODIS only	
WSP CLD CODIS Laboratory STR Analysis Procedures Manual: Evaluating Fusion 6C profiles with GeneMapper® ID-X v1.6.2 as an Expert System6C CODIS only	

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14. Population Genetics and Statistics

The purpose of this module is to give detailed background information on the different statistics used in forensic DNA testing (focusing on likelihood ratios), the considerations that need to be taken into account when calculating each statistic, the use of STRmix[™] to calculate likelihood ratios, and statistics dealing with kinship and paternity.

14.1. Goals

At the end of this session the trainee should be able to:

- 14.1.1. Understand the principles and mathematics behind population statistics, including the use of theta.
- 14.1.2. Perform likelihood ratio calculations in STRmix™.
- 14.1.3. Conduct statistics for kinship and paternity using CODIS Popstats.

14.2. Tasks

Instruction, demonstration, and practical training will be provided

- 14.2.1. Population genetics and statistics pertaining to forensic DNA analysis (CODIS)
- 14.2.2. Use of the STRmix™ program for statistical calculations
- 14.2.3. Use of CODIS Popstats for statistical calculations involving kinship

14.3. Assessment

Trainees will incorporate STRmix[™] and Popstats statistics in their work with the ~20 mixtures assigned in module 13. The trainer will document the completion of the module using the trainer's evaluation form.

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WSP DNA BIOCHEMISTRY AND STR TRAINING PROGRAM MODULE 14 – Population Genetics and Statistics Evaluation Form

	Population genetics and statistics in fo	orensic DN/	A analysis (CODIS)	
	Statistical calculation			
	Paternity/kinship			
The tra	rainee has completed the above checke Understand the principles behind vari Perform likelihood ratios using STRm with laboratory policy	ious statistic		ce
Comm	nents:			
-				_
I	Trainee Printed Name + Initials	Date	Trainer Printed Name + Initials Date)

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MODULE 14 POPULATION GENETICS AND STATISTICS READING ASSIGNMENTS

HT

REFERENCE	INITIALS	DATE
Butler JM. 2015. Advanced Topics in Forensic DNA Typing: Interpretation. San Diego:		
Elsevier Academic Press. Chapter 9 Statistical Interpretation Overview: 213-236.		
Butler JM. 2015. Advanced Topics in Forensic DNA Typing: Interpretation. San Diego:		
Elsevier Academic Press. Chapter 10 STR Population Data Analysis: 239-270.		
Butler JM. 2015. Advanced Topics in Forensic DNA Typing: Interpretation. San Diego:		
Elsevier Academic Press. Chapter 11 DNA Profile Frequency Estimates and		
Match Probabilities: 281-305.		
Butler JM. 2015. Advanced Topics in Forensic DNA Typing: Interpretation. San Diego: Elsevier Academic Press. Chapter 12 DNA Mixture Statistics: 309-329.		
Lander ES, Budowle B. DNA fingerprinting dispute laid to rest. Nature 1994; 371: 735-		
738.		
National Research Council Committee on DNA Forensic Science. 1992. DNA		
Technology in Forensic Science. Washington, D.C.: National Academies Press.		
Chapter 3 DNA Typing: Statistical Basis for Interpretation: 74-96.		
National Research Council Committee on DNA Forensic Science. 1996. The Evaluation		
of Forensic DNA Evidence. Washington, D.C.: National Academies Press.		
Chapters 4 Population Genetics and 5 Statistical Issues: 89-165.		
Evett IW, Weir BS. 1998. Interpreting DNA Evidence: Statistical Genetics for Forensic		
Scientists. Sunderland (MA): Sinauer Associates. Chapter 4 Population Genetics:		
79-131.		
Evett IW, Weir BS. 1998. Interpreting DNA Evidence: Statistical Genetics for Forensic		
Scientists. Sunderland (MA): Sinauer Associates. Chapter 5 Statistical Genetics: 132-162.		
Evett IW, Weir BS. 1998. Interpreting DNA Evidence: Statistical Genetics for Forensic		
Scientists. Sunderland (MA): Sinauer Associates. Chapter 8 Calculating Match		
Probabilities: 206-216.		
Evett IW, Weir BS. 1998. Interpreting DNA Evidence: Statistical Genetics for Forensic		
Scientists. Sunderland (MA): Sinauer Associates. Chapter 9 Presenting Evidence:		
217-246.		
Adams C. 2010. Essential Mathematics and Statistics for Forensic Science. Hoboken		
(NJ): John Wiley & Sons. Chapter 11 Statistics and the significance of evidence.		
279-311.		
Budowle B, et al. CODIS STR loci data from 41 sample populations. J Forensic Sci		
2001; 46(3): 453-489.		
Moretti T, et al. Population data on the expanded CODIS core STR loci for eleven		
populations of significance for forensic DNA analyses in the United States. Forensic Sci Int Genet 2016; 25: 175-181.		
Gettings KB, et al. STRSeq: a catalog of sequence diversity at human identification short	+	
tandem repeat loci. Forensic Sci Int Genet 2017; 31: 111-117.		
Bright J, et al. A guide to forensic DNA interpretation and linkage. Promega Corporation		
website. 2014. http://www.promega.com/resources/profiles-in-dna/2014/a-guide-		
to-forensic-dna-interpretation-and-linkage/		
Taylor D, et al. Testing likelihood ratios produced from complex DNA profiles. Forensic		
Sci Int Genet 2015; 16: 165-171.		

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Slooten K. Likelihood ratio distributions and the (ir)relevance of error rates. Forensic Sci Int 2019. https://doi.org/10.1016/j.fsigen.2019.102173	
WSP CLD DNA Casework STR Analysis Procedures Manual: STRmix™ V2.8	
Butler JM. 2015. Advanced Topics in Forensic DNA Typing: Interpretation. San Diego: Elsevier Academic Press. Chapter 14 Relationship Testing: 349-387.	
Chakraborty R, Stivers DN. Paternity exclusion by DNA markers: Effects of paternal mutations. J Forensic Sci 1996; 41(4): 671-677.	
Gunn PR, et al. DNA analysis in disputed parentage: The occurrence of two apparently false exclusions of paternity, both at short tandem repeat (STR) loci, in the one child. Electrophoresis 1997; 18(9): 1650-1652.	
Panke ES, et al. DNA paternity tests: Technology is outpacing the law. Family Law Newsletter 2001.	
WSP CLD DNA Casework STR Analysis Procedures Manual: Kinship Analysis	

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15. Combined DNA Index System

The purpose of this module is to provide instruction on the Combined DNA Index System (CODIS) uploads, searches, further CODIS eligibility guidelines, and report wording of CODIS-related conclusions.

15.1. Goals

At the end of this session the trainee should be able to:

- 15.1.1. Explain the laboratory's CODIS program including eligibility guidelines and how samples are searched and/or uploaded.
- 15.1.2. Use CODIS to determine the MME/MRE of a forensic specimen.
- 15.1.3. Understand the relationship of CODIS between the local, state, and national levels.

15.2. Tasks

Instruction and demonstration will be provided

- 15.2.1. Observe the calculation of the MME/MRE of a sample
- 15.2.2. Observe the entry of a CODIS profile into LDIS
- 15.2.3. Complete the assigned training modules in the CODIS software

15.3. Assessment

No practical exam or competency is provided for this training section. The trainer will assess through discussion of the trainee's knowledge of the subject matter and will document the training using the trainer's evaluation form.

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WSP DNA BIOCHEMISTRY AND STR TRAINING PROGRAM MODULE 15 – CODIS

Evaluation Form

	CODIS sample eligibility, sample entry, search,	and upload
	Use of CODIS Popstats Moderate Match Estima	ate and Match Rarity Estimate (MME and MRE)
The tra	ainee has completed the above checked sections Explain the laboratory's CODIS program includ searched and/or uploaded Use CODIS to enter a DNA profile and determin	ng eligibility guidelines and how samples are
Comm	ents:	
- T	rainee Printed Name + Initials Date	Trainer Printed Name + Initials Date

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MODULE 15 CODIS READING ASSIGNMENTS

REFERENCE	INITIALS	DATE
Butler JM. 2012. Advanced Topics in Forensic DNA Typing: Methodology. San Diego:		
Elsevier Academic Press. Chapter 8 DNA Databases: 213-264.		
Hares D. Selection and implementation of expanded CODIS core loci in the United		
States. Forensic Sci Int Genet 2015; 17: 33-34.		
WSP CLD CODIS Standard Operating Procedures Manual		
Federal Bureau of Investigation. A Guide to Determining What is Allowable in the		
Forensic Index at NDIS.		
WSP CLD CODIS Training PowerPoints: CODIS New Analyst Training		
WSP CLD CODIS Training PowerPoints: CODIS Eligibility		
WSP CLD CODIS Training PowerPoints: CODIS Forensic Mixture Eval		
CODIS software modules. CODIS CJIS-SEN LMS.		
WSP CLD DNA Casework STR Analysis Procedures Manual: CODIS & CODIS Match		
Prediction		
WSP CLD DNA Casework STR Analysis Procedures Manual: CODIS Export		
Butler JM. 2012. Advanced Topics in Forensic DNA Typing: Methodology. San Diego:		
Elsevier Academic Press. Appendix 2 Familial DNA Searches: 603-609.		
Butler JM. 2005. Forensic DNA Typing: Biology, Technology, and Genetics of STR		
Markers. New York: Elsevier Academic Press. D.N.A. Box 18.1 The business		
case for using forensic DNA technology: 436-437.		
Steinberger E, Sims G. Finding criminals through the DNA of their relatives – familial		
searching of the California offender DNA database. Prosecutor's Brief 2008; 31(1-2): 28-32.		
Myers S, et al. Searching for first-degree familial relationships in California's offender		
DNA database: validation of a likelihood ratio-based approach. Forensic Sci Int		
Genet 2011; 5(5): 493-500.		
WSP CLD CODIS Laboratory STR Analysis Procedures Manual: Reporting Profiles &	1	
CODIS Database CODIS only		
WSP CLD CODIS Laboratory STR Analysis Procedures Manual: Appendix A:		
Administrative Procedures for Processing Offender Samples CODIS only		

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16. Case Files, Report Writing, and Review

The Casework DNA Analyst trainee will receive instruction on requirements for report writing, wording of conclusions, and organization of the case file. Trainees will also become familiar with the technical and administrative review process.

16.1. Goals

At the end of this session the trainee should be able to:

- 16.1.1. Correctly report conclusions of serological and DNA analysis.
- 16.1.2. Understand the required components of a report.
- 16.1.3. Organize a complete case file.
- 16.1.4. Explain the review process through both technical and administrative reviews, including LIMS milestone and activities.

16.2. Tasks

16.2.1. Review at least five completed case files representing the scope of expected casework, focusing on their organization and contents

16.3. Assessment

Trainees will demonstrate report writing and case file preparation in their work involving the ~20 mixtures assigned in module 13. The trainer will document the completion of the module using the trainer's evaluation form.

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WSP DNA BIOCHEMISTRY AND STR TRAINING PROGRAM MODULE 16 – Case Files, Report Writing, and Review Evaluation Form

	Organization and contents of case	files		
	Report writing and LIMS			
The tra	ninee has completed the above checonomic Correctly write reports compatible was Explain the steps taken in LIMS to	with laborator	ry policy	
Comm	ents:			
Ţ	rainee Printed Name + Initials	Date	Trainer Printed Name + Initials	Date

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MODULE 16 CASE FILES, REPORT WRITING, AND REVIEW READING ASSIGNMENTS

	17	
-	- 1	

TRAINEE:	

REFERENCE	INITIALS	DATE
WSP CLD Quality Operations Manual: Case Management, Sections 1-9		
WSP CLD DNA Casework STR Analysis Procedures Manual: STR Case File Content		
WSP CLD DNA Analysis Quality Assurance Manual: Reports		
WSP CLD DNA Analysis Quality Assurance Manual: Review		
WSP CLD DNA Casework STR Analysis Procedures Manual: Laboratory Reports		
WSP CLD DNA Casework STR Analysis Procedures Manual: Guidelines for		
Paternity/Kinship: Guidelines for Report Writing – Paternity, Parentage, and		
Kinship		
WSP CLD DNA Casework STR Analysis Procedures Manual: Guidelines for		
Paternity/Kinship: Guidelines for Conclusion Statements – Paternity, Parentage,		
and Kinship		
WSP CLD DNA Casework STR Analysis Procedures Manual: Reports Involving CODIS		
WSP CLD CODIS Laboratory STR Analysis Procedures Manual: CODIS Case File		
Content CODIS only		
WSP CLD DNA Casework STR Analysis Procedures Manual: STR Case File Review		
WSP CLD CODIS Laboratory STR Analysis Procedures Manual: CODIS Case File		
Technical Review with GeneMapper® ID-X v1.6.2 as an Analysis ToolCODIS		
only		
WSP CLD CODIS Laboratory STR Analysis Procedures Manual: CODIS Case File		
Technical Review with GeneMapper® ID-X v1.6.2 as an Expert System" CODIS		
only		

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17. Additional DNA Sources, Automation, and Trends

This purpose of this module is to introduce trainees to other types of DNA used for forensic purposes, along with their advantages and drawbacks. Also, trainees will learn about automation in the forensic laboratory and other trends regarding forensic DNA testing. This is to educate the trainee in areas of DNA testing that may not be part of their specific workflow, but exist in the field of forensic DNA testing.

17.1. Goals

At the end of this training session, the trainee should be able to:

- 17.1.1. Explain the difference between SNP, Y-STR, and mitochondrial DNA, and how they differ from autosomal STRs (Tech understand Y-STR typing).
- 17.1.2. Feel familiar with the use of robotics in the forensic laboratory.
- 17.1.3. Become familiar with the following topics:
 - 17.1.3.1. Expert system software (CODIS)
 - 17.1.3.2. Rapid DNA systems (CODIS)
 - 17.1.3.3. Massively parallel sequencing (CODIS)

17.2. Tasks

Trainers will familiarize trainees with automation specific to their job duties (CODIS)

17.3. Assessment

No practical examination or competency is provided for this training section. The trainer will assess through discussion of the trainee's knowledge of the subject matter and will document the training using the trainer's evaluation form.

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WSP DNA BIOCHEMISTRY AND STR TRAINING PROGRAM MODULE 17 – Additional DNA Sources, Automation, and Trends Evaluation Form

	Single nucleotide polymorphisms	
	Y chromosome DNA typing (Tech)	
	Mitochondrial DNA	
	Automation in the forensic DNA laboratory (CODIS)	
	Expert systems software (CODIS) (Tech)	
	Rapid DNA systems (CODIS)	
	Massively parallel sequencing (CODIS)	
The t	trainee has completed the above checked sections and is able to: Explain the types of DNA used in forensic analysis other than autosomal STRs Understand the trends in forensic DNA analysis, including the use of automation (CO)	DIS)
Comr	nments:	
	Trainee Printed Name + Initials Date Trainer Printed Name + Initials	Date

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MODULE 17 ADDITIONAL DNA SOURCES, AUTOMATION, AND TRENDS READING ASSIGNMENTS

REFERENCE	INITIALS	DATE
Additional DNA Sources		
Butler JM. 2012. Advanced Topics in Forensic DNA Typing: Methodology. San Diego: Elsevier Academic Press. Chapter 12 Single Nucleotide Polymorphisms and Applications: 347-362. HT		
Butler JM. 2012. Advanced Topics in Forensic DNA Typing: Methodology. San Diego: Elsevier Academic Press. Chapter 13 Y-Chromosome DNA Testing: 371-396. HT		
Butler JM. 2015. Advanced Topics in Forensic DNA Typing: Interpretation. San Diego: Elsevier Academic Press. Chapter 15 Lineage Marker Statistics: 403-425. HT		
Gill P, et al. DNA Commission of the International Society of Forensic Genetics: recommendations on forensic analysis using Y-chromosome STRs. Int J Legal Med 2001; 114(6): 305-309. (also Forensic Sci Int 2001; 124(1): 5-10) HT		
Jobling MA, Tyler-Smith C. The human Y chromosome: an evolutionary marker comes of age. Nature Reviews Genetics 2003; 4: 598-612. HT		
Butler JM. Recent developments in Y-short tandem repeat and Y-single nucleotide polymorphism analysis. Forensic Sci Review 2003; 15(2): 92-100. HT		
Gusmão L, et al. DNA Commission of the International Society of Forensic Genetics (ISFG): an update of the recommendations on the use of Y-STRs in forensic analysis. Int J Legal Med 2006; 120(4): 191-200. (also Forensic Sci Int 2006; 157: 187-197) HT		
Isenberg AR. Forensic mitochondrial DNA analysis: a different crime-solving tool. FBI Law Enforcement Bulletin 2002; 71(8): 16-22. HT		
Butler JM. 2012. Advanced Topics in Forensic DNA Typing: Methodology. San Diego: Elsevier Academic Press. Chapter 14 Mitochondrial DNA Analysis: 405-444. HT		
Automation		
WSP CLD DNA internal validation summary for the Qiagen QIAgility™ (if applicable)		
WSP CLD DNA Casework STR Analysis Procedures Manual: Qiagen QIAgility™ Instructions (if applicable)		
WSP CLD DNA internal validation summary for the Hamilton Microlab STARlet (if applicable)		
WSP CLD DNA Casework STR Analysis Procedures Manual: Hamilton Microlab STARlet (if applicable)		
De Jong, B et al. Hamilton AutoLys developmental validation: Successful validation of a fully automated sample lysis workstation. Forensic Sci Int Genet 2013; 4: e93-e94.		
Hamilton AutoLys and/or STARlet Operator's manuals.		
Baron, L et al. Breakthrough in forensic workflow automation, eliminating the sample preparation and lysis bottlenecks with the AutoLys STAR: Technology and validation study. Netherlands Forensic Institute Poster. EAFS 2012.		
Butler JM. 2012. Advanced Topics in Forensic DNA Typing: Methodology. San Diego: Elsevier Academic Press. Chapter 17 New Technologies and Automation: 497-509. HT		
WSP CLD CODIS lab internal validation summary for GMID-X as an NDIS-approved expert system CODIS only		

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Budowle B. Typing megaplex human identity marker panels on multiple CGS platforms. Presentation at the International Symposium on Human Identification, Atlanta. 2013. HT	
Tan E, et al. Fully integrated, fully automated generation of short tandem repeat profiles. Investigative Genetics 2013; 4(1): 16. HT	

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18. Y-STR DNA Typing for Casework

The analysis of STRs on the Y chromosome utilizes the same technology and principles as autosomal STRs. The Y-STR trainee must be currently or previously qualified in autosomal STR analysis before undergoing the Y-STR training module. This module will provide the in-depth scientific knowledge relating to the application of Y-STRs to forensic DNA analysis. This module will provide practical instruction to the trainee on the analytical protocols used in the laboratory for Y-STR amplification and analysis. This module will also provide instruction on how to interpret and report Y-STR analytical results with established laboratory policy.

18.1. Goals

At the end of this training session, the trainee should be able to:

- 18.1.1. Pass a test (oral or written) on the basic concepts of the Y-chromosome and forensic Y-STR analysis.
- 18.1.2. Competently perform Y-STR analysis on biological samples that would normally be encountered in forensic casework and issue properly reported conclusions.
- 18.1.3. Demonstrate the understanding and use of a haplotype database and statistical interpretation.

18.2. Tasks

Trainers will discuss with trainees subject matter and published references

- 18.2.1. Evolution, molecular biology, and properties of the Y-chromosome
- 18.2.2. Forensic applications of Y-STR analysis
- 18.2.3. Amplification with the currently validated Y-STR amplification kit (Tech)
- 18.2.4. Typing of Y-STR amp product on a genetic analyzer (Tech)
- 18.2.5. Interpretation and reporting of Y-STR results (Tech)
- 18.2.6. Population databases and Y-STR statistics
- 18.2.7. Testimony, practice, and observation
- 18.2.8. Analysis of three single source male DNA extracts (Tech)
- 18.2.9. Trainee will be provided with 6 sets of Y-STR data. The data sets will include one of each of the following types of samples: single source, partial profile, mixture with a major component, mixture with a deducible minor component, mixture with a known contributor, and an indistinguishable mixture.

18.3. Assessment

Completion of a competency exam is required to complete this module of training regardless of prior Y-STR analysis experience. The exam will consist of 50% written questions and 50% oral questions. The trainee will complete a competency test consisting of one non-probative case and will prepare full documentation of the analysis and interpretations in the format used for regular casework following the established WSP Y-STR, STR, and quality assurance casework procedures. (Tech)

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WSP DNA BIOCHEMISTRY AND STR TRAINING PROGRAM MODULE 18 – Y-STR DNA Typing for Casework

Evaluation Form

•	Trainee Printed Name + Initials Date Trainer Printed Name + Initials	Date	
Comn	nents:		
The tr	rainee has completed the above checked sections and is able to: Discuss and display an in-depth knowledge of forensic Y-STR analysis Competently perform Y-STR analysis on biological samples similar to what would be encountered in forensic DNA casework Correctly interpret Y-STR data and write reports compatible with laboratory policy		
	Written exam (passing grade of 80% or higher)		
	Y-STR non-probative competency test (Tech)		
	Interpretation of Y-STR data (6 sets)		
	Single source male extracts (3) (Tech)		
	Y-STR testimony practice and observation		
	Y-STR statistics and population databases		
	Interpretation of Y-STR data (Tech)		
	Y-STR amplification kit/typing on the genetic analyzer (Tech)		
	Forensic application of Y-STRs		
	Y chromosome: evolution and biology		

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MODULE 18 Y-STR DNA TYPING FOR CASEWORK READING ASSIGNMENTS

REFERENCE	INITIALS	DATE
WSP CLD DNA Casework STR Analysis Procedures Manual: Y-STR Casework Tech		
WSP CLD DNA Casework STR Analysis Procedures Manual: Y-STR Amplification Tech		
WSP CLD DNA Casework STR Analysis Procedures Manual: Y-STR Amplification		
Product Preparation Tech		
Genescan™ 600 LIZ® size standard product insert		
Thompson J, et al. Developmental validation of the PowerPlex® Y23 system: a single multiplex Y-STR analysis system for casework and database samples. Forensic Sci Int Genet 2013; 7(2): 240-250.		
Promega. 2021. PowerPlex® Y23 technical manual. TMD035. Tech		
WSP CLD DNA validation summary for the Promega PowerPlex® Y23 system using the AB® 3500		
WSP CLD DNA supplemental internal validation summary for the Promega PowerPlex® Y23 system		
WSP CLD DNA PowerPlex® Y23 training day presentations		
Moore D, et al. Description of artefacts in the PowerPlex® Y23 system associated with excessive quantities of background female DNA. Forensic Sci Int Genet 2016; 24: 44-50.		
Lee E, et al. Off-ladder alleles due to a single nucleotide polymorphism in the flanking region at DYS481 detected by the PowerPlex® Y23 system. Forensic Sci Int Genet 2016; 24: e7-e8.		
Butler JM, et al. Chromosomal duplications along the Y-chromosome and their potential impact on Y-STR interpretation. J Forensic Sci 2005; 50(4): 853-859.		
SWGDAM Interpretation Guidelines for Y-Chromosome STR Typing. 2014.		
WSP CLD DNA Casework STR Analysis Procedures Manual: Y-STR Guidelines for Evaluating GeneMapper® Data Tech		
WSP CLD DNA Casework STR Analysis Procedures Manual: Y-STR Mixture Deconvolution		
Budowle B, et al. Basic principles for estimating the rarity of Y-STR haplotypes derived from forensic evidence. Presentation at the 18 th Annual International Symposium on Human Identification. 2007.		
Willuweit S, Roewer L. The new Y chromosome haplotype reference database. Forensic Sci Int Genet 2015; 15: 43-48.		
WSP CLD DNA Casework STR Analysis Procedures Manual: Y-STR Statistical Calculations		
WSP CLD DNA Casework STR Analysis Procedures Manual: Y-STR Laboratory Reports		

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Validation and Quality Assurance

The purpose of this module is to inform trainees of the processes and requirements of validation and performance checks, and to give them a detailed explanation of the quality assurance program as related to DNA testing, instrumentation, and procedures.

19.1. Goals

At the end of this session the trainee should be:

- 19.1.1. Familiar with the requirements of WSP CLD's internal validation program as it applies to procedures and instrumentation
- 19.1.2. Able to recognize the importance of the calibration and maintenance of equipment
- 19.1.3. Understanding the proficiency testing program
- 19.1.4. Know what is required for accreditation
- 19.1.5. Aware of the audit process

19.2. Tasks

Trainers will familiarize trainees with quality assurance systems specific to their job duties.

- 19.2.1. Validation
- 19.2.2. Instrument calibration and maintenance
- 19.2.3. Proficiency testing
- 19.2.4. Accreditation and auditing

19.3. Assessment

No practical examination or competency is provided for this training section. The trainer will assess through discussion of the trainee's knowledge of the subject matter and will document the training using the trainer's evaluation form.

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WSP DNA BIOCHEMISTRY AND STR TRAINING PROGRAM MODULE 19– Validation and Quality Assurance Evaluation Form

	Validation, calibration, and main	tenance of in	strumentation	
	Accreditation and auditing			
The tra	ainee is able to: Understand the validation proce instruments Explain accreditation, auditing, a		niliar with the maintenance required for iency program	applicable
Comm	ents:			
- ר	Frainee Printed Name + Initials	 Date		 Date

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MODULE 19 VALIDATION AND QUALITY ASSURANCE READING ASSIGNMENTS

TRAINEE:	

REFERENCE	INITIALS	DATE
WSP CLD Quality Operations Manual: Traceability and Quality Control HT		
WSP CLD Quality Operations Manual: Method Validation HT		
WSP CLD DNA Analysis Quality Assurance Manual: Methods Validation CODIS HT Tech		
WSP CLD DNA Analysis Quality Assurance Manual: Equipment Calibration and		
Maintenance CODIS HT Tech		
WSP CLD DNA Analysis Quality Assurance Manual: Proficiency Testing CODIS HT Tech		
WSP CLD DNA Analysis Quality Assurance Manual: Non-Conforming Work CODIS HT		
Tech		
WSP CLD DNA Analysis Quality Assurance Manual: Audits and Quality Review CODIS		
HT Tech		
WSP CLD DNA Casework STR Analysis Procedures Manual: Sample Switch Detection		
HT		
WSP CLD DNA Casework STR Analysis Procedures Manual: Quality Assurance/Quality		
Control HT Tech		
WSP CLD CODIS Laboratory STR Analysis Procedures Manual: Quality		
Assurance/Quality Control CODIS		
WSP CLD DNA Casework STR Analysis Procedures Manual: Reagent Quality Control HT Tech		
WSP CLD DNA Casework STR Analysis Procedures Manual: Reagent Lot Numbers and		
Expiration Dates HT Tech		
WSP CLD DNA Casework STR Analysis Procedures Manual: Reagent Preparation HT		
Tech		
WSP CLD DNA Casework STR Analysis Procedures Manual: Calibration of Instruments		
HT Tech		
WSP CLD DNA Casework STR Analysis Procedures Manual: Tempgenius – Setup and		
Maintenance HT Tech		
WSP CLD DNA Casework STR Analysis Procedures Manual: Qiagen BioRobot® EZ1 and		
EZ2 Connect – Maintenance HT Tech		
WSP CLD DNA Casework STR Analysis Procedures Manual: AB 7500 Real Time PCR		
Systems Instrument – Setup HT Tech		
WSP CLD DNA Casework STR Analysis Procedures Manual: AB 7500 Real Time PCR		
Systems Instrument – Maintenance HT Tech		
WSP CLD DNA Casework STR Analysis Procedures Manual: Hamilton Microlab STARlet		
- Maintenance		
WSP CLD DNA Casework STR Analysis Procedures Manual: Qiagen Qiagility™ –		
Maintenance		
WSP CLD DNA Casework STR Analysis Procedures Manual: AB 3500 Genetic Analyzer		
- Setup and Maintenance HT Tech		
Autoclave User's Manual (lab specific) HT Tech		
DNA QC Forms. SharePoint: DNA Forms & Templates HT Tech		
WSP CLD CODIS Laboratory STR Analysis Procedures Manual: Performance Check		
and Calibration of Instruments CODIS only		
WSP CLD CODIS Laboratory STR Analysis Procedures Manual: Tempgenius Wireless		
Data Acquisition & Monitoring System CODIS only		

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WSP CLD CODIS Laboratory STR Analysis Procedures Manual: Tempgenius Wireless Data Acquisition & Monitoring System Maintenance CODIS only	
WSP CLD CODIS Laboratory STR Analysis Procedures Manual: Reagent Preparation CODIS only	
WSP CLD CODIS Laboratory STR Analysis Procedures Manual: UV Irradiator Operating Instructions CODIS only	
WSP CLD CODIS Laboratory STR Analysis Procedures Manual: BSD600 Duet Puncher Maintenance CODIS only	
WSP CLD CODIS Laboratory STR Analysis Procedures Manual: BSD600 Ascent Puncher Maintenance CODIS only	
WSP CLD CODIS Laboratory STR Analysis Procedures Manual: EZ1 Advanced XL Maintenance CODIS only	
WSP CLD CODIS Laboratory STR Analysis Procedures Manual: 7500 Real Time PCR System Maintenance CODIS only	
WSP CLD CODIS Laboratory STR Analysis Procedures Manual: 9700 Thermal Cycler Maintenance CODIS only	
WSP CLD CODIS Laboratory STR Analysis Procedures Manual: ProFlex PCR System Maintenance CODIS only	
WSP CLD CODIS Laboratory STR Analysis Procedures Manual: 3500XL Genetic Analyzer Maintenance CODIS only	
WSP CLD CODIS Laboratory STR Analysis Procedures Manual: GeneMapper® ID-X v.1.6.2 Server Maintenance CODIS only	
WSP CLD CODIS Laboratory STR Analysis Procedures Manual: Tuttnauer Tabletop Autoclave – Instrument Maintenance CODIS only	

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20. Testimony, Legal Issues, and Ethics

The purpose of this module is to provide instruction and prepare the Casework DNA Analyst trainee for court presentation in the State of Washington. Unless the Casework DNA Analyst trainee has previous DNA typing testimony experience, at least one moot court session must be conducted in preparation for giving testimony. The trainee should be encouraged to attend court and observe experienced forensic scientists testify.

20.1. Goals

At the end of this session the trainee should be:

- 20.1.1. Familiar with the legal system for Washington State as it pertains to expert witnesses
- 20.1.2. Able to provide unbiased, clear, and easy to understand expert testimony on forensic DNA analysis
- 20.1.3. Aware of legal issues surrounding DNA testing and testimony
- 20.1.4. Confident in their ability to meet the ethical standards expected of an expert witness

20.2. Tasks

Instruction will be provided

- 20.2.1. Courtroom procedures and rules of evidence processes
 - 20.2.1.1. Court structure of both trial and appeals courts
 - 20.2.1.2. Format of hearing or trial
 - 20.2.1.3. Discovery and admissibility rules
 - 20.2.1.4. Courtroom demeanor and attire
- 20.2.2. DNA analyst qualifications
- 20.2.3. Technical testimony
 - 20.2.3.1. Testifying to testing results
 - 20.2.3.2. The importance of communicating the statistical weight of DNA associations
 - 20.2.3.3. Framework for evidence evaluation, the hierarchy of propositions, and presentation of expert opinion
 - 20.2.3.4. Communicating assumptions and limitations of test methods and results
- 20.2.4. Testimony practice of direct and cross examination
- 20.2.5. Ethical responsibility of expert witnesses
- 20.2.6. Evidence and exhibit presentation
 - 20.2.6.1. Handling of evidence
 - 20.2.6.2. Exhibit continuity
- 20.2.7. State and federal DNA database legal authority
 - 20.2.7.1. Permissible samples and profiles
 - 20.2.7.2. Confidentiality and disclosure of information
- 20.2.8. Review curriculum vitae of trainee (Tech)
- 20.2.9. Observation of witness testimony (Tech)
- 20.2.10. Legal issues (NAS, PCAST reports; successful legal challenges)

20.3. Assessment

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Participation in a minimum of one successful moot court is required to complete this module. The results of the moot court performance evaluation will be retained by the laboratory as part of the trainee's file. The trainer will document completion of this module using the trainer's evaluation form.

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WSP DNA BIOCHEMISTRY AND STR TRAINING PROGRAM

MODULE 20 – Testimony, Legal Issues, and Ethics Evaluation Form

- T	rainee Printed Name + Initials Date Trainer Printed Name + Initials Date
Comm	ents:
The tra	ainee has completed the above checked sections and is able to: Understand the legal system for Washington State as it pertains to expert witnesses Provide unbiased, clear, and easy to understand expert testimony on forensic DNA analysis
	Legal issues
	Moot court (Tech - oral board)
	Review curriculum vitae and observe expert witness testimony (Tech)
	State and federal DNA database legal authority
	Evidence/exhibit presentation
	Ethical responsibility of expert witnesses
	Testimony practice (direct and cross examination)
	Technical testimony
	DNA analyst qualifications
	Courtroom procedures and process of rules of evidence

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MODULE 20 TESTIMONY, LEGAL ISSUES, AND ETHICS READING ASSIGNMENTS

ΗТ

TRAINEE:	

REFERENCE	INITIALS	DATE
WSP CLD Quality Operations Manual: Case Management, Sections 10-12		
A Citizen's Guide to Washington Courts. http://www.courts.wa.gov. Organization of WA		
courts.		
WSP CLD DNA Analysis Quality Assurance Manual: Discovery and Public Disclosure CODIS Tech		
Coleman H, Swenson E. 1994. DNA in the Courtroom: A Trial Watcher's Guide. Seattle: Genelex Press. Chapter 5 DNA in the Courtroom: 75-92.		
Matson JV, et al. 2004. Effective Expert Witnessing: Practices for the 21st Century. Boca Raton (FL): CRC Press. Chapter 1 The Legal Environment: 1-11.		
Matson JV, et al. 2004. Effective Expert Witnessing: Practices for the 21st Century. Boca Raton (FL): CRC Press. Chapter 2 Key Cases and Precedents Affecting Expert Witnessing: 13-22.		
Matson JV, et al. 2004. Effective Expert Witnessing: Practices for the 21st Century. Boca Raton (FL): CRC Press. Chapter 4 The Pre-Trial Process: 35-57.		
Matson JV, et al. 2004. Effective Expert Witnessing: Practices for the 21st Century. Boca Raton (FL): CRC Press. Chapter 5 Preparing for Trial: 59-66.		
Matson JV, et al. 2004. Effective Expert Witnessing: Practices for the 21st Century. Boca Raton (FL): CRC Press. Chapter 6 The Courtroom Drama: 71-87.		
Matson JV, et al. 2004. Effective Expert Witnessing: Practices for the 21st Century. Boca Raton (FL): CRC Press. Chapter 7 The Art of Expert Witnessing: 89-101.		
Butler JM. 2012. Advanced Topics in Forensic DNA Typing: Methodology. San Diego: Elsevier Academic Press. Chapter 18 Legal Aspects of DNA Testing and the Scientific Expert in Court: 515-543.		
National Judicial College and Justice Speakers Institute. 2020. Science Bench Book for Judges. Section 6.2 Pre-Trial Discovery: 209-218.		
National Judicial College and Justice Speakers Institute. 2020. Science Bench Book for Judges. Section 7 Trial: 219-243.		
National Judicial College and Justice Speakers Institute. 2020. Science Bench Book for Judges. Section 9 The Expert Witness: 261-278.		
National Research Council Committee on DNA Forensic Science. 1992. DNA Technology in Forensic Science. Washington, D.C.: National Academies Press. Chapters 6 Use of DNA Information in the Legal System and 7 DNA Typing and Society: 131-164.		
National Research Council Committee on DNA Forensic Science. 1996. The Evaluation of Forensic DNA Evidence. Washington, D.C.: National Academies Press. Chapter 1 Introduction: 47-59 and Chapter 6 DNA Evidence in the Legal System: 166-211. Note significant changes between the two NRC reports.		
Holmgren J. DNA evidence and jury comprehension. Canadian Society of Forensic Sci J 2005; 38(3): 123-141.		
Eldridge H. Juror comprehension of forensic expert testimony: a literature review and gap analysis. Forensic Sci Int: Synergy 2019; 1: 24-34.		
Donnelly P, Friedman RD. DNA database searches and the legal consumption of scientific evidence. Michigan Law Review 1999; 97(4): 931-984.		
Robertson J. Integrity issues impacting on the provision of forensic services. Australian J Forensic Sci 1999: 31(2): 87-97.		

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National Research Council. 2009. Strengthening Forensic Science in the United States:	
A Path Forward. Washington, D.C.: The National Academies Press.	<u> </u>
President's Council of Advisors on Science and Technology. 2016. Report to the	
President Forensic Science in Criminal Courts: Ensuring Scientific Validity of	
Feature-Comparison Methods. Executive Summary: 1-20.	
President's Council of Advisors on Science and Technology. 2016. Report to the	
President Forensic Science in Criminal Courts: Ensuring Scientific Validity of	
Feature-Comparison Methods. Chapter 1 Introduction: 21-24.	
President's Council of Advisors on Science and Technology. 2016. Report to the	
President Forensic Science in Criminal Courts: Ensuring Scientific Validity of	
Feature-Comparison Methods. Chapter 2 Previous Work on Validity of Forensic-Science Methods: 25-39.	
President's Council of Advisors on Science and Technology. 2016. Report to the	
President Forensic Science in Criminal Courts: Ensuring Scientific Validity of	
Feature-Comparison Methods. Chapter 3 The Role of Scientific Validity in the	
Courts: 40-43.	
President's Council of Advisors on Science and Technology. 2016. Report to the	
President Forensic Science in Criminal Courts: Ensuring Scientific Validity of	
Feature-Comparison Methods. Chapter 4 Scientific Criteria for Validity and	
Reliability of Forensic Feature-Comparison Methods: 44-66.	
President's Council of Advisors on Science and Technology. 2016. Report to the	
President Forensic Science in Criminal Courts: Ensuring Scientific Validity of	
Feature-Comparison Methods. Chapter 5 Evaluation of Scientific Validity for	
Seven Feature-Comparison Methods: Introduction, Section 5.1 DNA Analysis of	
Single-source and Simple-mixture Samples, Section 5.2 DNA Analysis of	
Complex-mixture Samples, and Conclusion: 67-75, 122-123.	<u> </u>
President's Council of Advisors on Science and Technology. 2016. Report to the	
President Forensic Science in Criminal Courts: Ensuring Scientific Validity of	
Feature-Comparison Methods. Chapter 6 Recommendations to NIST and OSTP:	
124-130.	
President's Council of Advisors on Science and Technology. 2016. Report to the	
President Forensic Science in Criminal Courts: Ensuring Scientific Validity of	
Feature-Comparison Methods. Chapter 10 Scientific Findings: 146-150.	
President's Council of Advisors on Science and Technology. 2016. Report to the	
President Forensic Science in Criminal Courts: Ensuring Scientific Validity of	
Feature-Comparison Methods. Appendix A Statistical Issues: 151-154.	
ANAB. 2018. Guiding principles of professional responsibility for forensic service	
providers and forensic personnel. GD 1350: 1-3. WSP internal training presentation. 2024. Expert Testimony: General Considerations,	+
Evaluative Framework, and the Hierarchy of Propositions.	
Evaluative Framework, and the meratchy of Fropositions.	

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21. Cognitive Bias

This training will provide the DNA analyst with an introduction to cognitive bias and its role in forensic science.

21.1. Goals

At the end of this session, the trainee should be:

- 21.1.1. Familiar with the different types of bias that can affect forensic science.
- 21.1.2. Recognize and minimize bias during the testing process.

21.2. Tasks

Trainers will discuss with trainees subject matter on the following topics:

- 21.2.1. Cognitive, contextual, and confirmation bias
- 21.2.2. Steps to minimize cognitive bias
- 21.2.3. Analysts should participate in a cognitive bias discussion annually in conjunction with the ASCLD Guiding Principles review.

21.3. Assessment

No practical examination or competency is provided for this training section. The trainer will assess through discussion of the trainee's knowledge of the subject matter and will document the training using the trainer's evaluation form.

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WSP DNA BIOCHEMISTRY AND STR TRAINING PROGRAM MODULE 21 – Cognitive Bias Evaluation Form

	Types of bias			
	Ways to minimize cognitive bias			
The tr	ainee has completed the above ch Recognize the different types of			
Comm	nents:			
-	 Trainee Printed Name + Initials	 Date	Trainer Printed Name + Initials	 Date

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MODULE 21 COGNITIVE BIAS READING ASSIGNMENTS

CODIS HT

REFERENCE	INITIALS	DATE
CLD cognitive bias PowerPoint presentation		
Dror I, Hampikian G. Subjectivity and bias in forensic DNA mixture interpretation. Sci and Justice 2011; 51(4): 204-208.		
Biedermann A. Prediction in forensic science: a critical examination of common understandings. Frontiers in Psychology 2015; 6: 1-4.		

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22. Outsourcing

This module outlines the outsourcing program used by the CLD to send DNA work to laboratories outside the Washington State Patrol Crime Laboratory system. Training specific to the review of outsourced casework will be dependent on the outsourcing laboratory and will be done separately.

22.1. Goals

At the end of this session, the trainee should be:

22.1.1. Able to describe the outsourcing process, including in-house review

22.2. Tasks

Trainers will briefly discuss when outsourcing is used by the laboratory and how the process of outsourcing works

22.3. Assessment

No practical examination or competency is provided for this training section. The trainer will assess through discussion of the trainee's knowledge of the subject matter and will document the training using the trainer's evaluation form.

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WSP DNA BIOCHEMISTRY AND STR TRAINING PROGRAM MODULE 22 – Outsourcing

Evaluation Form

	Outsourcing process			
The tr	ainee is able to: Explain the process of outsourcing	g cases, inclu	ding in-house review	
Comm	nents:			
<u>.</u>	Trainee Printed Name + Initials	 Date	Trainer Printed Name + Initials	 Date

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MODULE 22 OUTSOURCING READING ASSIGNMENTS

REFERENCE	INITIALS	DATE
WSP CLD Quality Operations Manual: Subcontracting of Tests		
WSP CLD DNA Analysis Quality Assurance Manual: Outsourcing CODIS		
WSP CLD DNA Casework STR Analysis Procedures Manual: Outsourced Ownership		
Review		

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23. Final Evaluations

23.1. Goals

- 23.1.1. Pass a written exam on in-depth knowledge appropriate to their duties
- 23.1.2. Each Casework/CODIS DNA Analyst trainee (optional for experienced staff training in this area) will prepare and give a lecture presentation of ~20-30 minutes in length to WSP CLD scientific staff on a topic in which in-depth knowledge is required.
 - 23.1.2.1. This will be followed by a brief question and answer period
 - 23.1.2.2. A written dissertation of the presentation is also required
- 23.1.3. Successfully complete competency consisting of mock case(s).

23.2. Tasks

- 23.2.1. Take written exams for biochemistry and DNA (HT trainees will also take two written tests, one for Phase I and one for Phase II)
- 23.2.2. Present a lecture presentation and written dissertation
- 23.2.3. Complete competency consisting of mock case(s)

23.3. Assessment

A passing score (≥80%) on the written exam is required to complete training. Successful completion of the lecture presentation and written dissertation is required for all Casework/CODIS DNA Analyst trainees. Competency samples in the form of a mock case (or non-probative case) will be provided to the Casework DNA Analyst trainee (samples from Module 14 work may be used). The Casework DNA Analyst trainee will prepare full documentation of the analysis and interpretations in the format used for regular casework. The CODIS DNA Analyst trainee will be provided with competency samples representative of what will be encountered in performing regular work duties. The CODIS DNA Analyst trainee will prepare full documentation of the analysis as required for convicted offender database entry.

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WSP DNA BIOCHEMISTRY AND STR TRAINING PROGRAM

MODULE 23 – Final Evaluations

Evaluation Form

	Written exam (passing grade of 80% or higher)					
	Lecture presentation and	written dissertation*				
	Training reviewed by DN	Training reviewed by DNA technical leader				
	Name	 Initials	 Date			
Case	ework DNA Analyst					
	Competency test					
COD	IS DNA Analyst					
	Competency test (manua	al extraction and direct amp	olification)			
compre provide of one with DI	ng meets the following criteria: piehension of subject, able to answed to explain subject. Trainee ma or two of the preceding criteria nNA Technical Leader). rainee has completed the a	ver questions (from lecture) to the y pass on condition of successful ot being fully met (additional wo	ne satisfaction of the trainer, sul completion of further specifork assigned by trainer to mee	ufficient scientific detail ied work or in the instance		
	Show complete understa casework, shown	nding of the principles and through written examinati ensic DNA laboratory proc	d knowledge involved in ion and/or an oral prese	ntation		
Comr	ments:					
	Trainee Printed Name + Ini	itials Date Tr	ainer Printed Name + In	itials Date		

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