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Overview of DNA Programs at NIST

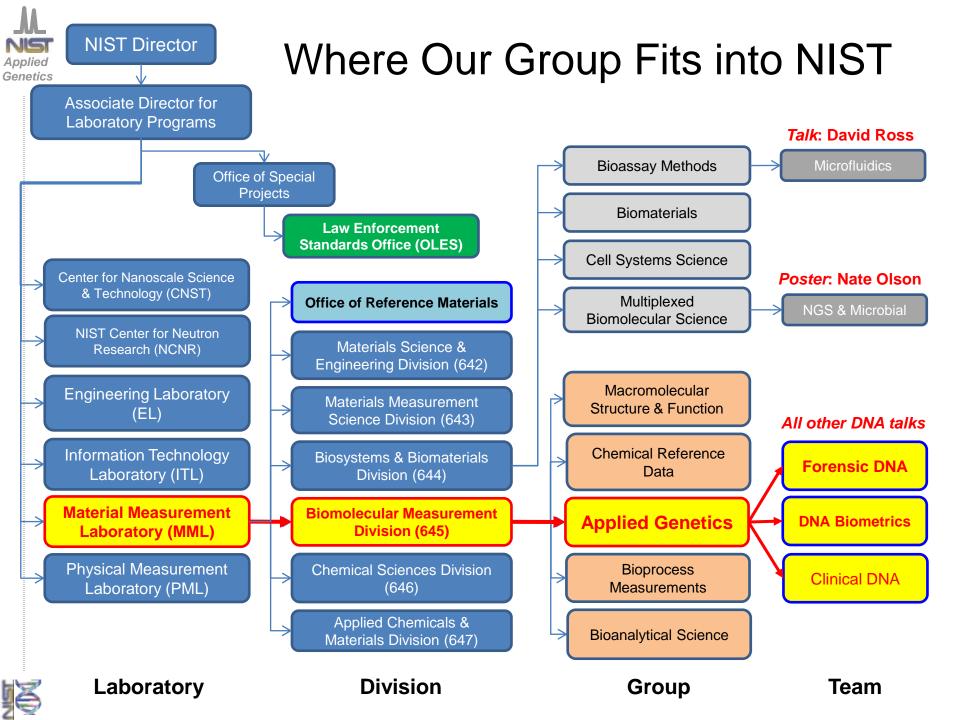
John M. Butler

NIST Fellow & Applied Genetics Group Leader

Forensics@NIST 2012 Meeting

Gaithersburg, MD November 28, 2012







NIST Human Identity Project Teams within the Applied Genetics Group

Forensic DNA Team



John

Butler



Mike

Coble





Kline

Support

Data Analysis

S

Dave Duewer



Pete Vallone Erica Butts



Funding from the **National Institute of Justice (NIJ)** through NIST Law Enforcement Standards Office (OLES)

Becky

Hill

Funding from the **FBI S&T Branch** through NIST Information Access Division

DNA Biometrics Team



Clinical DNA



Ross Haynes





http://www.nist.gov/mml/bmd/genetics/applied_genetics_pubs.cfm

Our Team's Consistent Funding and Productivity

Genetics

http://www.cstl.nist.gov/strbase/NISTpub.htm

	NIJ		http://www.csti.nist.gov/strbase/NISTpub.htm			
Year	Funding	<u>Staff</u>	Publications	Presentations	<u>Workshops</u>	
Pre-2000	\$100-200k	5	~3	~5	0	
FY2002	\$950k	6	6	33	0	
FY2003	\$900k	6	10	23	0	
FY2004	\$1.1M	7	15	24	1	
FY2005	\$1.2M	7	16	38	2	
FY2006	\$1.2M	6	14	42	7	
FY2007	\$1.1M	6	14	44	9	
FY2008	\$1.0M	7	11	29	6	
FY2009	\$1.1M +	7	7	50	11	
FY2010	\$1.2M <mark>+</mark>	9	11	51	2	
FY2011	\$1.2M <mark>+</mark>	9+1	18	72	10	
FY2012	\$1.2M +	8+1	22	58	9	
TOTALS	\$12.2 M		144	464	57	
Average ~\$1.1M 13.1 42.2 5.2 +FBI-funding (DNA biometrics)						



Summary of NIJ-Funded Research



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Required under FBI Quality Assurance Standard 9.5.5 for labs connected to the national DNA database or receiving federal \$





DNA quantity measurement calibration

Autosomal and Y-chromosome short tandem repeat (STR) measurement calibration





- Standard Reference Materials
 - SRM 2391c (autosomal & Y-STRs), 2395 (Y-STRs), 2392-I (mtDNA)
 - SRM 2372 (DNA quantitation) being re-certified
- Technology Evaluation and Development
 - The presentations you will hear today represent a portion of our work over the past few years

Training Materials

- Workshops & STRBase website
- Textbooks (Forensic DNA Typing)



Applied Genetics

Plan for Presentations This Morning

DNA typing process



DNA Stability, Extractions and Quantitation

9:00-9:05	Overview of DNA Programs at NIST – John Butler
9:05-9:25	Stability Studies – Margaret Kline
9:25-9:40	DNA Extraction – Erica Butts
9:40-10:00	DRAGEN (NIST Forensics Grand Challenge Project) – David Ross
10:00-10:20	Digital PCR & DNA Quantitation – Ross Haynes

Report

10:20-10:50Break and Poster Viewing/Exhibitor Displays

STRs, mtDNA, Rapid DNA

- 10:50-11:10 STR Kits & New Loci Becky Hill
- 11:10-11:30 STRBase and Information Resources John Butler
- 11:30-11:50 Mixture Interpretation & True Allele Mike Coble
- 11:50-12:10 mtDNA base composition Kevin Kiesler
- 12:10-12:30 Rapid DNA Pete Vallone

12:30-1:45 Lunch and Poster Viewing/Exhibitor Displays





DNA Posters (Lecture Room B)

- 1. DNA Sequencing Error Estimation and SNP Validation for Microbial Forensics Applications
- 2. Characteristics of 24 Commonly Used Autosomal STR Loci
- 3. Population Statistics on the Proposed Expanded U.S. Core Loci
- 4. Concordance Testing Comparing STR Multiplex Kits with a Standard Data Set
- 5. SE33 Variant Alleles: Sequences and Implications
- 6. Evaluation of Additional Y-STR Loci to Resolve Common Haplotypes
- 7. An Evaluation of Additional Y-STR Loci in the PowerPlex Y-23 Kit
- 8. Direct PCR Amplification of STR Loci: Protocols and Performance
- 9. Rapid Amplification of Commercial STR Typing Kits
- 10. Validation of PowerPlex 16 HS in Comparison to Identifiler Plus on the ABI 3500 Genetic Analyzer
- 11. Setting Interpretation Thresholds and Results with Low-Level DNA Analysis
- 12. The New SRM 2391c: PCR-based DNA Profiling Standard
- 13. Using SRM 2372 Human Quantitation Standard: Are there differences between qPCR assays?
- 14. Forensic Performance of Insertion-Deletion Marker Systems
- 15. Candidate Reference Family Data: A Tool for Validating Kinship Analysis Software
- 16. Rapid DNA Testing Approaches for Reference Samples





Disclaimer

NIST Disclaimer: Certain commercial equipment, instruments and materials are identified in order to specify experimental procedures as completely as possible. In no case does such identification imply a recommendation or it imply that any of the materials, instruments or equipment identified are necessarily the best available for the purpose.

Points of view are those of the presenters and do not necessarily represent the official position of the National Institute of Standards and Technology or the U.S. Department of Justice.





For More Information on Forensic DNA Efforts at NIST...

http://www.cstl.nist.gov/strbase 301-975-4049; john.butler@nist.gov



Our team publications and presentations are available at: http://www.cstl.nist.gov/strbase/NISTpub.htm

