

# NIST On-Going Projects to Aid the Human Identity **Testing Community**

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Visit to Applied Biosystems - July 19, 2005



# Team Impact on Forensic Community

- 27 publications since June 2004 (61 since 2000)
- 31 presentations to the community since June 2004
- All NIST publications and presentations available on STRBase: http://www.cstl.nist.gov/biotech/strbase/NISTpub.htm
- · Training materials: 2 workshops conducted with Bruce McCord NEAFS (Sept 29-30, 2004) - Albany DNA Academy (June 13-14, 2005)
- Forensic DNA Typing: Biology, Technology, and Genetics of STR Markers, 2<sup>nd</sup> Edition (John Butler)

#### National Institute of Justice of the U.S. Department of Justice

#### Current Areas of NIST Research Effort

- Resources for "Challenging Samples" (miniSTRs)
- Information on New Loci (SNPs, Y-Chromosome, new STRs)
- Standard Information Resources (STRBase website, training materials/review articles, validation standardization)
- Allele Sequencing and Interlaboratory Studies (Real-time qPCR, mixture interpretation)

# Mike Coble

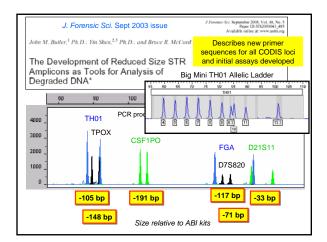
# miniSTRs for Degraded DNA

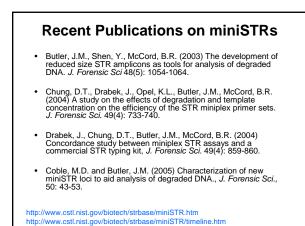
John Butler

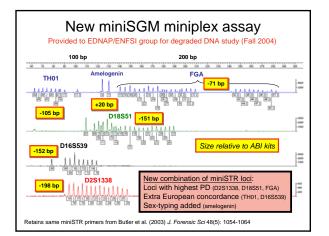
- Original miniSTR paper with CODIS loci, D2, D19, Penta D, Penta E Butler et al. (2003) J. Forensic Sci. 48: 1054-1064
- Many CODIS loci are too big and make poor miniSTRs.
- New miniSTRs and assays: NC01, NC02 - Coble, M.D. and Butler, J.M. (2005) J. Forensic Sci. 50:43-53
- · New miniSGM miniplex: AMEL, TH01, FGA, D18, D16, D2
- EDNAP/ENFSI degraded DNA study coordinated by Peter Gill
- · Creation of miniSTR information on STRBase

http://www.cstl.nist.gov/biotech/strbase/miniSTR.htm

A miniSTR is a reduced size STR amplicon that enables higher recovery of information from degraded DNA samples miniSTR Conventional PCR primer primer STR repeat region -miniSTR Conventional PCR primer Testing must be performed to show allele concordance between primer sets prime 260 Conventional STR test 4000 (COfiler™ kit) 9 11 246.27 254.37 70 75 80 85 90 95 100 105 110 115 120 12. 141 MiniSTR assay (using ~150 bp smaller Butler et al. 2003 primers) (03.50) 102.07 er, J.M. (2005) Forensic DNA Typ







#### Many CODIS Loci Make Poor miniSTRs

- Large allele range (e.g., FGA)
- Large alleles (e.g., D21S11 and FGA)
- Poor flanking regions prohibiting reliable primer annealing immediately adjacent to the repeat region (e.g., D7S820)

# Why go beyond CODIS loci

"STRs have proven to be highly successful [for mass disasters] in the past e.g. Waco disaster and various air disasters. However, even if the DNA is high quality there are occasions when there are insufficient family members available to achieve a high level of confidence with an association."

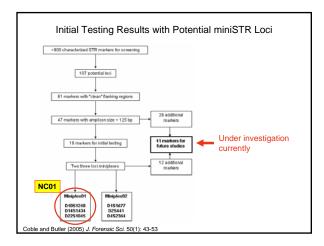
Gill, P., Werrett, D.J., Budowle, B. and Guerrieri, R. (2004) An assessment of whether SNPs will replace STRs in national DNA databases-Joint considerations of the DNA working group of the European Network of Forensis Science hsitutuse (ISHS) and the Scientific Working Group on DNA Analysis Methods (SWGDAM). Science&Justice, 44(1): 51-53. Why go beyond CODIS loci
 "To achieve this purpose, either <u>new STRs</u> <u>could be developed</u>, or alternatively, existing STRs could be supplemented with a SNP panel."
 "There are also efforts for modifying existing STR panels by decreasing the size amplicons by designing new primers."

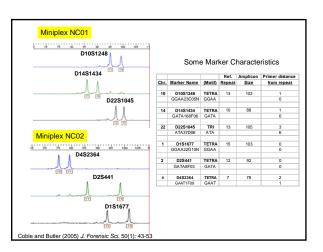
# Why go beyond CODIS loci

- Desirable to have markers unlinked from CODIS loci (different chromosomes) for some applications
- Small size ranges to aid amplification from degraded DNA samples
- New miniSTR loci will benefit missing persons investigations and paternity testing (and perhaps national databases in the future)

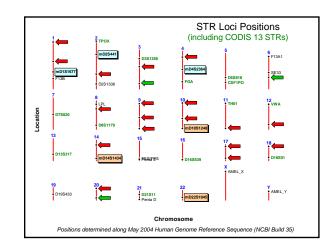
#### Characterization of New miniSTR Loci

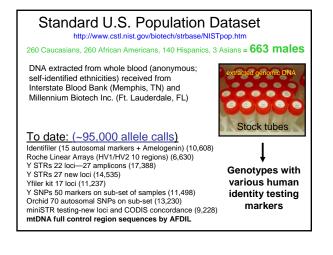
- Candidate STR marker selection
- Chromosomal locations and marker characteristics
- PCR primer design
- Initial testing results
- Population testing
- Allelic ladder construction
- Miniplex assay performance

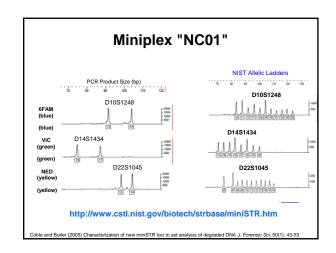


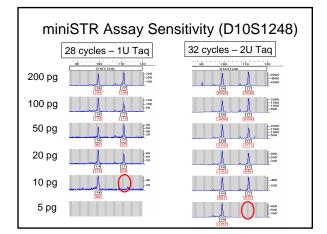


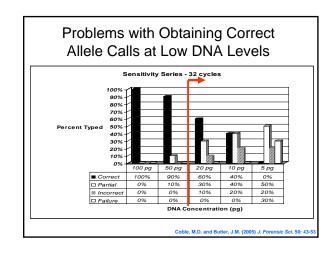


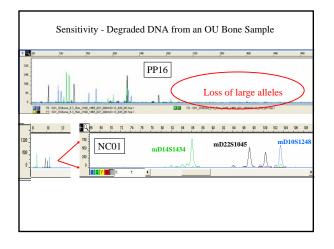






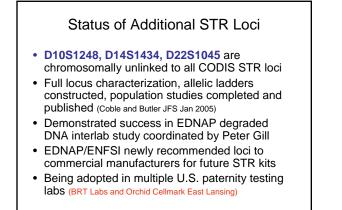


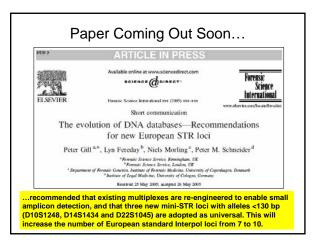








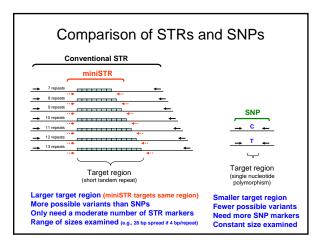






- Construction of T2plex autosofial SNF assay
- Creation of Forensic SNP Information website on STRBase
   see Gill et al. Science&Justice 44(1): 51-53

http://www.cstl.nist.gov/biotech/strbase/SNP.htm





# Work with Y-STRs

John Margaret Pete Amy Butler Kline Vallone Decker

- Beta-testing of all commercial Y-STR kits
- Population data supplied to Yfiler haplotype database
- · 49 Y-STR loci evaluated with ~650 U.S. samples
- New Y-chromosome information on STRBase linking to all available haplotype databases
- Nomenclature defined for new loci
- Human Y-Chromosome DNA Profiling Standard Reference Material (SRM 2395) – updates with DYS635 for Yfiler
- Separation of two brothers with 47 Y-STRs
  http://www.cstl.nist.gov/biotech/strbase/y\_strs.htm

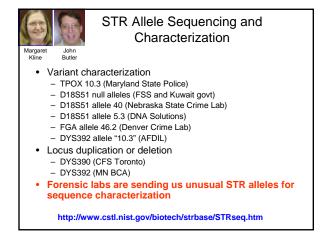


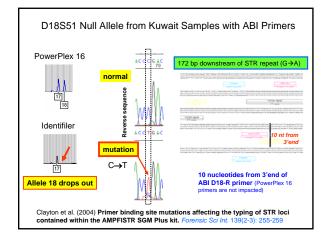
#### Evaluation of qPCR Assays

Evaluation of published assays on same samples

- Characterization of Quantifiler lot-to-lot performance
- · Additional studies under way utilizing qPCR:
  - Examining the challenge of multiplexing qPCR assays
  - Studies to track DNA recovery from various types of tubes
    Characterizing potential SRM 2372 components (Human DNA
  - Quantitation Standard)

http://www.cstl.nist.gov/biotech/strbase/DNAquant.htm







Validation Standardization

- Survey initiated at June 2004 NIJ meeting and conducted last summer resulted in 53 responses
- Talk at Promega meeting Oct 2004
- Validation summary sheets .
- Validation website on STRBase .
- · We invite submission of your internal validation studies for inclusion in the NIST validation website

http://www.cstl.nist.gov/biotech/strbase/validation.htm





#### STRBase Updates

Jan Redman

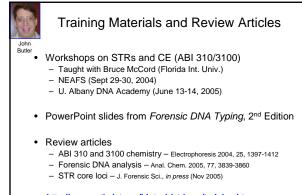
#### Primary updates performed monthly

- · Summary of variant alleles and tri-allelic patterns
- List of STR references (Reference Manager database)
- NIST publications and presentations
- · New content is being added regularly to aid training and to support forensic DNA laboratories

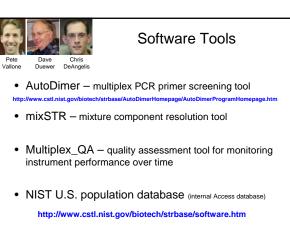
http://www.cstl.nist.gov/biotech/strbase/ http://www.cstl.nist.gov/biotech/strbase/NISTpub.htm http://www.cstl.nist.gov/biotech/strbase/var\_tab.htm

Content of STRBase Website http://www.cstl.nist.gov/biotech/strbase STR Fact Sheets on Core Loci .../str\_fact.htm .../multiplx.htm Multiplex STR Kit Information .../y\_strs.htm Y-Chromosome Information Variant Alleles Reported .../var\_tab.htm .../mutation.htm Mutation Rates for Common STRs Reference List with ~2,300 Papers .../str\_ref.htm Downloadable PowerPoints for Training • .../training.htm .../validation.htm Validation Information ٠ .../miniSTR.htm miniSTR Information ٠ .../address.htm Addresses for Scientists • .../NISTpub.htm Publications & Presentations from NIST •

# Presentation to Applied Biosystems



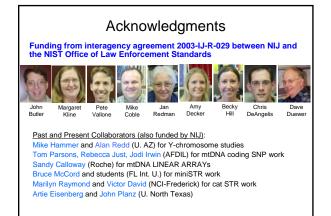
http://www.cstl.nist.gov/biotech/strbase/training.htm http://www.cstl.nist.gov/biotech/strbase/NISTpub.htm





- 8 DNA samples supplied
- 84 laboratories signed up (80 labs returned results)
- 287 data sets using 19 different methods
- 60 data sets with real-time qPCR (37 Quantifiler data sets)
- Publication in May 2005: J. Forensic Sci. 50(3): 571-578
- Mixture Interpretation Study (MIX05)
  - 91 labs signed up (64 labs returned data)
  - Interpretation requested of provided e-grams for 4 mock sexual
  - assault cases
  - Data analysis is still on-going...

http://www.cstl.nist.gov/biotech/strbase/interlab.htm



#### **Disclaimers and Collaborations**

Funding: Interagency Agreement 2003-IJ-R-029 between the National Institute of Justice and NIST Office of Law Enforcement Standards

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