

Association for Molecular Pathology November 17, 2010 – San Jose, CA

Dialogue with NIST: Standard Reference Materials for Molecular Diagnostics

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NIST Applied Genetics Group

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Presentation Outline

- NIST
 - location, role, organizational structure, funding
- Applied Genetics Group
 - members, expertise, equipment, funding
 - programs, projects, priorities
- Standard Reference Materials (SRMs)
 - Purpose, principles, production process, pricing
 - Differences from WHO standards (certification vs consensus)

Clinical DNA SRMs

- Pipeline, plans, and path forward
- Partnerships: needs/priorities from the community





Please ask

questions



NIST Background





Location of NIST





National Institute of Standards & Technology (NIST)

- Non-regulatory agency established in 1901 in the US Department of Commerce.
- Mission to promote US innovation and industrial competitiveness by advancing measurement science, standards & technology.
- NIST is at the top of the US standards pyramid for a wide variety of physical standards, test methods, and calibrations.







Organic Act of 1901; Updated in 2008

"It is therefore the unanimous opinion of your committee that no more essential aid could be given to

- manufacturing
- commerce
- the makers of scientific apparatus
- the scientific work of Government
- schools, colleges, and universities

than by the establishment of the institution proposed in this bill."

Functions and Activities include:

- Custody and dissemination of national standards
 - Calibrations, Certified Reference Materials, Reference Data, Reference Methods
- Determination of physical constants and properties of materials
- Comparison of US National standards with those of the world
- Solution of standards and measurement problems for industry and other agencies







Early Driver for U.S. Standards



1904

 Out-of-town fire companies arriving at a Baltimore fire cannot couple their hoses to the hydrants. 1526 buildings razed.

1905

 National Fire Protection Association adopted NBS-developed national hose coupling standard.





NIST

Since its inception NBS/NIST has focused its research and measurement services activities on contemporary societal needs.

2010



1901



Standards for train tracks, couplings, steel manufacturing

Standards for clinical analytes, medical imaging, cybersecurity





CHCH2CH2CH2CH2CH



NIST Today

Major Assets

- ~ 2,900 employees
- ~ 2600 associates and facilities users
- ~ 400 NIST staff on about 1,000 national and international standards committees
- 3 Nobel Prizes in past 15 years



Major Programs

- NIST Laboratories
- Baldridge National Quality Program
- Hollings Manufacturing Extension Partnership
- Technology Innovation Program

Joint NIST/University Institutes:

- JILA
- Joint Quantum Institute
- Institute for Bioscience & Biotechnology Research
- Hollings Marine Laboratory





Role of NIST

NIST is a <u>non-regulatory</u> federal agency within the Technology Administration of the U.S. Department of Commerce. NIST's primary mission is to <u>promote</u> <u>economic growth by working</u> <u>with industry to develop and</u> <u>apply technology,</u> <u>measurements, and standards.</u>

NIST carries out its mission through a portfolio of four programs:

Measurement and Standards Program

planned and conducted in cooperation with industry and focused on infrastructural technologies

Baldridge National Quality Program

an outreach program recognizing organizational performance excellence

Hollings Manufacturing Extension Partnership Program

a nationwide network of extension centers that provides hands-on technical assistance to smaller manufacturers

Technology Innovation Program

Planned to provide costshared awards to industry, universities and consortia for research on potentially revolutionary technologies that address critical national and societal needs.





The NIST Laboratories

New Structure for NIST Laboratory Programs



Traditionally focused research and measurement service activities on physical science and engineering disciplines

Bioscience and Health identified as a new area for significant emphasis for NIST labs



Applied Genetics NIST Biochemical Science Division







NIST Applied Genetic Group







Applied Genetics Group Mission Statement

Advancing technology and traceability through quality genetic measurements to aid work in

- forensic DNA testing
- clinical diagnostics
- cell line authentication
- agricultural biotechnology
- DNA biometrics







APPLIED GENETICS Group Major Programs Currently Underway

Forensic DNA

- New loci and assays (26plex)
- STR kit testing
- Ancestry SNP assays
- Low-template DNA studies
- Mixture interpretation
- STR nomenclature
- Variant allele cataloging and sequencing
- Expert systems review
- Training workshops to forensic DNA laboratories
- Validation information and software tools
- Textbook 3rd ed. (2 vol.)

Clinical Genetics

- Huntington's Disease SRM
- CMV SRM
- Exploring future needs

Ag Biotech

 – "universal" GMO detection/ quantitation (35S promoter)

DNA Biometrics

- Rapid PCR methods
- Efforts to standardize testing of future portable DNA systems
- Kinship analysis

Cell Line Authentication





NIST Applied Genetics Group

Group Leader





Group Expertise and Funding Sources

Group Expertise

- Reference Material Characterization
- Standard Information Resource Development
- Rapid Multiplex PCR Assay Construction
- Short Tandem Repeat (STR) Genotyping
- Single Nucleotide Polymorphism (SNP) Genotyping
- DNA Sequencing
- Training Materials and Workshops (validation info)

Current Funding Sources

- National Institute of Justice (Forensic DNA)
- FBI Science & Technology Branch (DNA Biometrics)
- **NIST SRM Program** (SRM development and production)
- Base funding from Congress (clinical DNA)





Applied Genetics Group Instrumentation

- ABI 3130xI for Sanger sequencing, SNP analysis, and STR genotyping
- ABI 7500, Qiagen Roto-Gene Q, Roche LightCycler, for qPCR (DNA quantitation)
- ABI 9700 and Veriti thermal cyclers for PCR
- Fluidigm BioMark for digital PCR (copy number determination)





NIST Digital PCR Instrument









NIST Standard Reference Materials (SRMs)







SRM Information

- Purpose & use
- Decision to create & design of components
- Pricing: how determined?
- Characterization
 - Stability
 - Homogeneity
 - Value assignment
- Certified value: how determined?
- Difference in approach from WHO (NIBSC)



Applied Senetics What are Standards and Metrology?

- Standards and Metrology are the things needed to supply good data that can be used to confidently support decision making.
 - in a good world, good decisions are informed with good data which are the results of excellent measurements!
- Metrology and standards comprise the 'formal' system that tells us how well we trust those data
- Establishing confidence in data







Examples of Standards

- 1. Documentary Standard: Guideline documentation that reflects agreements in practices by governmental bodies or nationally or internationally recognized industrial, professional, trade associations
- 2. <u>Certified or Standard Reference</u> <u>Material</u>: A physical entity to serve as a reference in measuring quantities or qualities, establishing practices or procedures, or evaluating results



Standard Reference Material® 1643d

Trace Elements in Water

This Standard Reference Material (SRM) is intended primarily for use in evaluating methods used in the determination of trace elements in fresh water. SRM 16434 consists of approximately 230 nn. of fittered and asidified water in a polyetlyme hot Netr, which is studied in an atuminized platic laye to mariant ashibitir. SRM 16434 atimatas the elemental composition of fresh water. Nitric acid is present at a concentration of 0.5 mol/L to stabilize the trace elements.

The certified values for 26 elements in SRM 1643d are listed in Table 1. The information values for an additional four elements are provided in Table 2. The analytical methods used for the characterization of this SRM are given in Table 3. All values are reported as mass concentrations [1].

NOTICE AND WARNINGS TO USERS

Expiration of Certification: This certification of SRM 1643d is valid, within the measurement uncertainty(ies) specified, until 31 July 2003, provided the SRM is handled in accordance with instructions given in this certificate. This certification is multified if the SRM is damaged, constraintated, err modified.

Maintenance of SRM Certification: NIST will monitor this SRM over the period of its certification. If substantive technical changes occur that affect the certification before the expiration of this certification, NIST will notify the purchaser. Return of the attuched registration card will facilitate notification.

Presentions: The SRM should be shaken before use because of possible water condensation. To prevent possible contamination of the SRM do not inner pipets into the bottle. Samples should be decated at a noom temperature of 27 C = 5° C. After use, the bottle should be recargeded ightly and returned to the administed platch leag. Which should be folded and sealed with scaling gape. This safeguard will protect the SRM from possible environmental contamination and long-term expectation.

The accuracy of trace element determinations, especially at the µg/L level, is limited by contamination. Apparatus should be scrupulously cleaned and only high purity reagents employed. Sampling and manipulations, such as evaporations, should be done in a clean environment, such as a Class-100 clean bodd.

Coordination of the NIST technical measurements was under the direction of J.R. Moody of the NIST Analytical Chemistry Division.

The technical and support aspects involved in the certification and issuance of this SRM were coordinated through the Standard Reference Materials Program by J.S. Kane and B.S. MacDonald.

> Willie E. May, Chie Analytical Chemistry Division

> > Thomas E. Gills, Chief

Statistical analysis of the experi SRM 1643d

Gaithersburg, MD 20899

Revised Certificate Issue Dat





Importance of Reference Materials

Dr. Karen Mann, President of the Association of Molecular Pathology, in her testimony before Congress (Feb 24, 2010):

"Reference materials are important to ensure the necessary sensitivity, specificity and level of reproducibility of intra- and inter-laboratory test results. The best approach to achieve consistent and comparable quantitative data amongst laboratories is by the use of internationally established reference reagents."







Comparing Results

Results are only useful when compared

- to other results
 - e.g., to observe a trend
- to limits
 - e.g., a threshold for action
- different results in different places or measured at different times...
 - "comparability over spaceand-time"







Comparability of results

- Results linked to a common reference can be compared
- Scope of reference defines scope of comparability
 - global network
 - SI







Measurement Uncertainty

- Are these results the same?
- How well do you know the result?
 - essential part of being able to compare!
- Are these results good enough?
 – fit-for-purpose









"Quality" In Measurements

Method Validation

am I measuring what I set out to measure?

Uncertainty

how well do I know the result of what I've measured?

Traceability of Result (Reference Materials)

> can I compare this result with other results?







Helps meet ISO 17025 needs for traceability to a national metrology institute



How Do We Know an SRM is Needed?

- May be mandated or recommended by a scientific society or even Congress
 - SRMs are useful when there is a need for between laboratory comparability of measurements; especially if the labs use different measure equipment and/or methods
- Other agencies may come to NIST requesting an SRM
 - it is valuable to document who is requesting the reference material and why
 - The National Institute of Justice (NIJ) requested we develop SRMs for the Forensic DNA Testing Community when DNA technology started entering the courtroom
 - For NIST in general other agencies include: CDC, EPA, NIH...
 - The FDA when inspecting looks at the calibration of critical instruments and may stipulate the use of a higher order standard when available







NIST and other National Metrology Institutes (NMIs) worldwide:

- provide and maintain primary standards
- provide linkages for traceability to the international system of measurement
- disseminate these realizations in a manner and of a quality that is consistent with the needs of the measurement community

The typical role of an NMI is to establish and maintain:

Scientifically-Sound, Metrologically-Based Competencies and Measurement Capabilities that are Internationally Vetted and Recognized

To provide calibration and measurement services disseminated to Customers via mechanisms such as:

- Validated Reference Methods
- Certified Reference Materials
- Reference Data
- Value-assignment of customer-provided samples or materials
- Value-assignment of Proficiency Testing samples
- Measurement Services for other Government Agencies
- Etc.

Although physical quantities such as length, mass, temperature, time, etc. are the first to come to mind when one thinks about metrology and measurement standards, chemical measurement research and standards have been a major activity at NIST since its inception in 1901.







X

Applied Genetics



CIPM Mutual Recognition Arrangement

Objectives:



- Establish the degree of equivalence of national measurement standards maintained by NMIs
- Provide for the mutual recognition of both calibration and CRM certificates issued by NMIs
- Provide a secure technical foundation for wider agreements related to international trade, commerce and regulatory affairs

The CIPM Mutual Recognition Arrangement (MRA) was signed in October, 1999 by the directors of the NMIs of 47 member states of the *Metre Convention*, and representatives of two international organizations.

CIPM Mutual Recognition Arrangement



National Metrology Laboratories Must:

- Declare measurement capabilities that underpin services delivered to customers
- Participate in relevant International Key comparisons to validate claims
- Provide evidence of competence and <u>Quality</u> <u>systems</u> that underpin delivery of measurement services

Outcome:

 Statements of the measurement capabilities of each NMI in a database publicly available on the Web



Requirements for a NIST SRM

Material must be fit for purpose:

- Homogeneity
 - All tubes are the same
- Test random samples

- Stability
 - Will withstand shipping and normal storage and is periodically tested over the life time of the SRM
- Recoverability

Appropriate storage containers

- · What went in the tubes comes out
- Traceability
 - Values assigned are traceable to the designated certification method
- Commutability

Inter-laboratory study

• Is the SRM what the intended user needs?



Steps Involved in SRM Production

Attend conferences, read the literature, talk to potential customers

Sequence & Copy Number









Receive input on priorities for projects and potential SRMs

Research potential properties and samples to be characterized and measurement method to be used

Obtain candidate components/make measurements

Decide on number of SRM units to produce (impacts price/unit), sample packaging, concentration, etc.

Bottle components and conduct homogeneity and stability studies; finalize uncertainty

Write Report of Analysis and Certificate of Analysis

Certificate Reviewed and Approved by NIST Measurement Services Division

SRM Made Available for Purchase http://www.nist.gov/srm




Your

help

needed

Pipeline for SRM Production

- I. Request for SRM and internal decision to go forward
- 2. Understanding needs and scope of SRM (research)
- 3. SRM design

what formats and amounts are best?

- 4. Gather materials for SRM components
- 5. Characterization of components

Measure amounts with uncertainties, homogeneity, stability

- Complete paperwork
 - Report of Analysis (ROA) and Certificate of Analysis
- 7. Paperwork reviewed and SRM approved
- 8. SRM released (http://www.nist.gov/srm)
- 9. Customers use SRM and provide feedback

SD (standards development)

WCF (working capital funds)

Association for Molecular Patholog

SRM office

Applied Senetics How is NIST SRM Cost Determined?

 SRM price is based on cost recovery of working capital funds requested in order to produce the SRM components





Estimated Cost of an SRM

\$50K - 500 units produced\$50K - 100 units producedCalculate SRM Unit PriceCalculate SRM Unit Price

Laboratory	CSTL	Laboratory
SRM Category	105	SRM Category
Production Costs	\$50,000.00	Production Costs
Units	500	Units
NIST Unit Production Cost	\$100.00	NIST Unit Production
MSD SRM Operating Cost (Surcharge)	\$187.00	MSD SRM Operating
Cold	Storage Fee	
AD VALOREM Surcharge	\$50.00	AD VALOREM Surc
49% Service Development: \$49.00		49% Service Devel
1% Obsolescence: \$1.00		1% Obsolescence
Dept. of Commerce surcharge	\$4.32	Dept. of Commerce su
Lab Operations, if applicable	\$48.00	Lab Operations, if app
Total Price	\$389.32	Total Price
Rounded to Nearest Dollar *	\$389.00	Rounded to Nearest D

Laboratory SRM Category	CSTL 105
Production Costs	\$50,000.00
Jinis	100
NIST Unit Production Cost	\$500.00
MSD SRM Operating Cost (Surcharge) \$187.00
Cold	l Storage Fee
AD VALOREM Surcharge	\$250.00
49% Service Development: \$245.00)
1% Obsolescence: \$5.00)
Dept. of Commerce surcharge	\$4.32
Lab Operations, if applicable	\$48.00
lotal Price	\$989.32
Rounded to Nearest Dollar *	\$989.00



How does NIST Determine the Number of Units to Produce?

We (the staff producing the SRM) are responsible for determining the appropriate number of units to produce considering items such as:

- the perceived demand for each SRM based on historical sales rates of the SRM or of similar SRMs,
- the technical difficulty of future renewals,
- the marginal cost of producing additional units,
- the cost of storage space,
- material stability

The sales price is set to collect all production costs (no profit) and repay the WCF in **no more than five years**.





If we build it, will they come?

Will clinical DNA labs or commercial calibrant suppliers consider the cost of the NIST material too expensive or not important to their work?





Lessons Learned from Forensic DNA Experience

Mandate use of SRMs

- This mandates NIST participation & SRM production
- Consistent, significant funding
 - Enables research as well as SRM production
 - National Institute of Justice partnership
- Experienced staff and appropriate equipment
- Standard techniques and targets
 - Short tandem repeat (STR) typing and core loci
- Information sharing
 - Creation of STRBase website to help community
- Inter-laboratory studies
 - Lessons learned can help with SRM needs and design



Congress Passed the DNA Identification Act of 1994 (Public Law 103 322)

Formalized the FBI's authority to establish a national DNA index for law enforcement purposes.

FBI's DNA Advisory Board Quality Assurance Standards for Forensic DNA Testing Laboratories

(October 1, 1998)

STANDARD 9.5

The laboratory shall check its DNA procedures annually or whenever substantial changes are made to the protocol(s) against an appropriate and available NIST standard reference material or standard traceable to a NIST standard.





1995: SRM 2391 PCR-based DNA Profiling Standard





2003: NIST SRM 2391b

Driven primarily by commercial kit loci...

National Institute of Standards & Technology

Certificate of Analysis

Standard Reference Material® 2391b

PCR-based DNA Profiling Standard

This Standard Reference Material (SRM) is intended primarily for use in the standardization of forensic and paternily quality assurance procedures for Polymersue Chain Reaction (PCR)-based genetic testing and for instructional law enforcement or non-clinical research purposes. This SRM can also be used for quality assurance when assigning values to in-house control materials. It is not intended for any human or animal clinical diagnostic use. Note that SRM 2301b is alightly modified from SRM 2301, in that there is more emphasis on Short Tandem Repetits (STR4) and less emphasis on D1880 [1,2] reflecting the growing interest and utility of STRs [3 to 14]. Additional information on each STR locus can be found at a NIST-approximated database on the interest. <u>http://www.cstl.aisit.gov/bitech/utilose[14]</u>.

This SRM is composed of well-characterized human deoxyribonucleic acid (DNA) in two forms: genomic DNA and DNA to be extracted from cells aported onto filter paper. A unit of the SRM is composed of 12 fixeen components packaged in one box. See the section in this certificate entitled *Description of Components* for a complete listing of the components.

Certified Values: The SRM is certified for genetic loci of forensic interest that were commercially available at the time of production. Genetic types for these loci can be found in Tables 1, 2, and 3. The tables are organized as follows: Table 1 lists the genetic types for the Federal Bureau of Investigation's (FBF's) CODIS (<u>COmbined DNA</u> Index System) core STR loci; Table 2 lists additional STR loci of interest; and Table 3 lists the genetic types for D1S80, AmpliType^{III} PM + HLADQA1, and Amelogenin.

Expiration of Certification: The certification of this SRM is valid until 31 December 2008, provided the SRM is handled and stored in accordance with the instructions given in this certificate. However, the certification is invalid if the SRM is contaminated or otherwise modified.

Maintenance of SRM Certification: NIST will monitor this SRM over the period of its certification. If substantive technical changes occur that affect the certification before the expiration of certification, NIST will notify the purchaser. Return of the attached registration card will facilitate notification.

Storage: Store frozen at a temperature of -20 °C. DO NOT use a self-defrosting freezer because periodic cycling of temperatures may cause shortened shelf life of this SRM.

The overall direction and econdination of the technical activities leading to certification were under the chairmanship of J.M. Butler of the NIST Biotechnology Division.

Analytical determination and technical measurements leading to the certification of this SRM were performed by M.C. Kline and J.W. Redman of the NIST Biotechnology Division.

The support aspects involved in the preparation, certification, and issuance of this SRM were coordinated through the NEST Standard Reference Materials Group by C.S. Davis.

> Vincent Vilker, Acting Chief Biotechnology Division John Ramble, Jr., Chief

Measurement Services Division

Gaithenburg, MD 20899 Certificate Issue Date: 06 December 2002

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2. Certified Values for Additional STR Loci

F13B	FES/FPS	LPL	Penta D	Penta E	D2S1338	D19S433
10,10	12,12	10,11	10,15	7,12	17,23	13,16.2
8,10	10,11	00 -				16
9,10	11,12	22 8	iutos	oma	51 K	S 4
6,9	10,13	chai	racter	ized	acros	S 3
8,9	11,13	12	2 DNA	A sam	nples	,14
9,10	11,11	10,12	9,12	12,14	25,25	12,14
6,8	11,11*	11,12	3.2,11	12,16	17,22	13,15.2
6,8	10,11	9,11	8,9	5,10	22,22	12.2,15
8,10	10,12	11,12	12,12	12,13	19,23	14,15
8,8	11,11	10,12	8,12	11,11	23,23	13,14
8,10	10,12	11,12	12,12	12,13	19,23	14,15
8,8	11,11	10,12	8,12	11,11	23,23	13,14



NIST Standard Reference Material (SRM) for Forensic DNA Testing

SRM 2391b (2003-2011)

- 48 autosomal STR loci with certified values
- **10 liquid genomic DNA** components + **2 punches** (cells on 903 paper)
- All single source samples
- 4 males + 6 females
- 9947A & 9948 included

SRM 2391c (2011-future)

- 23 autosomal STR loci and 17 Y-STRs certified
- 4 liquid genomic DNA components + 2 punches (cells on FTA & 903 paper)
- 5 single source + 1 mixture
- 3 males + 2 females (unique)
- All new samples
 - no 9947A or 9948





How to use an SRM...

- The SRM material has been assigned a value through vigorous analysis methods.
- The appropriate SRM is used to value assign/validate materials and/or methods or calibrate.
- Specific SRM usage instructions are included in the Certificate of Analysis supplied with each SRM material
- The format of the SRM determines the analysis step for it's use. (carried through the complete analysis, or only at a later stage of analysis)





How to Create NIST Traceable Materials?

- Make a "lot" of material that you want to use as your control material.
- Test your control material for homogeneity and stability
- Analyze your control material in parallel with the SRM material, using the SRM material to assign the value to the control material.
- This "lot" of control material is now traceable to the NIST SRM.
- When this "lot" of control material is consumed or yields a different result it can no longer be used
- New "lots" of control materials must be analyzed in parallel with the SRM; not to one another





What is Documented When Creating a NIST SRM?







What is Documented When Creating a NIST SRM?

- Homogeneity, Stability, Fit for Purpose, Purity
- Well characterized material
 - 3 levels of certification: certified, reference, or informational values
- Well documented:
 - An established paper trail that would allow a future NIST scientist to reproduce the material if needed
 - ROA: Report of Analysis
 - An internal NIST document that is very detailed
 - COA: Certificate of Analysis
 - What is released with the reference material that describes the contents of the SRM along with the certified values





SRM 2395 Documentation

Store all data in a central, secure location

National Institute of Standards & Technology Certificate of Analysis

R

Standard Reference Material* 2395

Human Y-Chromosome DNA Profiling Standard (in Cooperation with the Netional Internate of Justice - U.S. Department of Justice)

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Viscan L. Vilker, Chief Joins Rambie, Jr., Chief

Page 1 of 3

Desiderations, MD 20099 Constructions Status 2003

1012 1022





SRM 2395 file drawer contents









Paper trail leads to a Certificate

National Institute of Standards & Technology

HUNE CORPORATION

Certificate of Analysis

Standard Reference Material[®] 2395

Human Y-Chromosome DNA Profiling Standard (In Cooperation with the National Institute of Justice – U.S. Department of Justice)

This Standard Reference Material is intended primarily for use in the standardization of forensic and paternity quality assurance procedures for Polymerase Chain Reaction (PCR)-based genetic testing and for instructional law enforcement or non-clinical research purposes that involve the human Y-chromosome. This SRM can also be used for quality assurance when assigning values to in-house control materials. It is not intended for any human or animal clinical diagnostic use. Additional information on each Y-chromosome marker can be found at a NIST-sponsored database on the Internet: <u>http://www.cstl.nist.gov/biotech/strbase</u>.

This SRM is composed of well-characterized human genomic deoxyribonucleic acid (DNA) in liquid form. A unit of the SRM is composed of 6 frozen components packaged in one box. There are five male samples and one female sample in this SRM. See the section in this certificate entitled *Description of Components* for a complete listing of the components.

This certificate will be provided with the reference material and describes what is certified





Report of Analysis (ROA)

US DEPARTMENT OF COMMERCE MATTOMAL INSTITUTE OF STANDARDS AND TECHNOLOOPY CHEMICALSCIENCE AND TECHNOLOOPY LABORATORY BIOCHEMICAL SCIENCE DIVISION GATTHER SBURG, MARYLAND 30300

REPORT OF ANALYSIS

25-Jime-2007

- Submitted to: Laurie E. Locascio, Chief Biochemical Science Division
- Authors: Margaret C. Kline Anny E. Decker David L. Duewer Peter M. Vallone
- Title:
 Preparation and Homogeneity Testing of Materials for SRM

 2372 Human DNA Quantitation Standard

Constituents:

Human Genomic DNA

The ROA is kept on file at the SRM Office (and the Division) and describes in great detail experimental data supporting the specified certified values

This ROA is 54 pages





Some of the details of SRM 2372

<u>Material Qualification</u>

Preliminary absorption spectra of the bulk materials over the wavelength range 220 nm to 345 nm were obtained using the BioCary 100 spectrophotometer located in 227/B261.

The molecular mass of the DNA in the bulk materials was assessed with a FlashGel (Cambrex Bio Science Rockland, Inc., Rockland, ME). The gel was loaded with 1 uL of each component and the proprietary FlashGel DNA Marker. The gel was electrophoresed for 7 minute at 275 Volts. Results of the analysis were imaged with the system's proprietary gel-stain and recorded as a digital photograph on the FlashGel Dock with viewing light.

Selection of Vials for Analysis

Vials of each component were pulled for four different sets of analysis: UV/vis certification measurements, UV/vis homogeneity assessment, qPCR homogeneity assessment, and interlaboratory validation. The certification measurements required the total volume of 51 vials; three vials were taken from three of the corner cells (coded red in the diagram to the right) of each of the 17 boxes. The UV/vis homogeneity assessment required one vial from each box; this was taken from the "near center" cells (coded

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teal green in the diagram). The qPCR homogeneity assessment also required one vial from each box; this was taken from one of the edge cells (along one edge between the red cells). The interlaboratory validation study consumed 34 vials of the A, B_{oB} and C materials; these were selected from the "near center" and the remaining corner.

UV/vis Verification and Homogeneity

Absorbance homogeneity for all components was assessed using the BioCary 100 spectrophotometer located in 227/B261. Seventeen vials, one from each of the storage boxes, were assayed for each component using 80 μ L cuvettes. The temperature of the cuvettes was 22 $\% \pm 1 \%$, the same as used in the HAS II certification measurements.

The detailed UV/V is homogeneity protocol is given in Appendix B.



<u>qPCR Commutability Studies.</u>

While the interlaboratory study supplied some information about systematic differences in response to the A, B_{old} and C materials among the available qPCR methods, studies were conducted at NIST to provide greater detail for specific assays [6,7,8] and to evaluate Component B_{mw} . Four commercially available DNA quantitation standards were used for these commutability studies. These studies are documented in MCK Notebook QPCR 05 pages 38 – 43, SRM2372\2372Certification\qPCR\qPCR_Usage.

One 96-well plate was used for each of the three qPCR methods evaluated. The design of the studies is shown schematically in Figure 4.

	1	2	З	4	5	6	7	8	9	10	11	12
A		A_1	B_1	0_1	SP_1A	SP_1A	SP_1C	SP_1C	A_1	B_1	0_1	Blank
Ð		A_2	B_2	C_2	SP_2A	SP_2A	SP_2C	SP_2C	A_2	B_2	C_2	
С		A_3	В_З	C_3	SP_3A	SP_3A	SP_3C	SP_3C	A_3	В_З	C_3	
D		A_4	B_4	C_4	SP_4A	SP_4A	SP_4C	SP_4C	A_4	B_4	C_4	
Е		A_5	B_5	C_5	SP_1B	SP_1B	SP_1D	SP_1D	A_5	B_5	C_5	
F		A_6	B_6	C_6	SP_2B	SP_2B	SP_2D	SP_2D	A_6	B_6	C_6	
G		A_7	B_7	C_7	SP_3B	SP_3B	SP_3D	SP_3D	A_7.	B_7	C_7	
Н		A_8	B_8	C_8	SP_4B	SP_4B	SP_4D	SP_4D	A_8	B_8	C_8	

Figure 4. qPCR Commutability Design

Note: "X_1" (where "X" represents the component A, B or C) is the 1:10 dibted material, X_2 is a 1:5 dibtion of X_1, X_3 is a 1:2 dibtion of X_2, X_5 is a 1:2 dibtion of X_4, X_6 is a 1:2 dibtion of X_5, X_7 is a 1:2 dibtion of X_6, and X_8 is a 1:2 dibtion of X_7. "SP_1A" to "SP_1D" are 1:10 dibtions of four commercially available DNA calibration materials, "SP_2x" (where "x" represents one of the four commercial materials) is a 1:5 dibtion of SP_1x, SP_3x is a 1:2 dibtion of SP_2x, and SP_4x is a 1:2 dibtion of SP_3x. Blank" denotes a buffer-only negative control. SP_A Applied Biosystems (Foster City, CA) Quantifiler Human DNA Standard Patt # 4343893 Lot #0602018, SP_C Applied Biosystems Quantifiler Human DNA Standard Patt # 4343893 Lot #0604020, SP_D Promega Corp (Madison, WI) Human Genomic DNA Male, Pat# G147A Lot #13636102. Applied Biosystems Quantifiler Human DNA Standard Patt # 2343893 Lot #0604020, SP_D Promega Human Genomic DNA Male reported [DNA] 262 ng/ µL.

Note book details

The SRM must be useful to the community. 2372 was checked against the most frequently used qPCR methods for the forensic Human Id community. Additional commercial stds were also tested.



1



Table 1. 2 µL Pipette Calibration

	Beatro	nic Pipet Reagent Lot	E00007 Code 54	98E	2 uL			
	Vet 1	Neagent Dr.	Mai 2	1601		Mal 2	VAL	
Tipe Box1	1	1 025	1	1 064		1	1 071	
rips box i		1.855		1.004			1.055	
	4	1.904		1.902		4	1.900	
	3	1.924	3	1.990		3	1.921	
	4	1.963	4	1.949		4	1.946	
	5	1.913	5	1.96		5	1.942	
	6	1.925	6	1.963		6	1.959	
	7	1.922	7	1.951		7	1.95	
	8	1.997	8	2.008		8	1.953	
	9	1.998	9	1.952		9	1.958	
	10	1.99	10	1.955		10	1.932	
	11	1.954	11	1.948		11	1.969	
	12	1.992	12	1.943		12	1.955	
	13	1.939	13	1.942		13	1.971	
	14	1.917	14	1.944		14	1.973	
	15	1.962	15	1.935		15	1.969	
	16	1.933	16	1.993		16	1.992	
	17	1.938	17	2.016		17	1.947	
	18	1.982	18	1.932		18	1.933	
	19	1.974	19	1.97		19	1.881	
	20	194	20	1.931		20	1.961	
Tips Box2	21	1 948	21	1.95		21	1 972	
1105 0002	22	2 196	22	1 932		22	1 963	
		2.100		1.302			1.000	
	Awa	1.963		1.960			1.953	Г

SĎ

0.059

Grand Mean SD C∨ 1.959 0.039 2.0 %

Setpoint

-18 %

0.023

Pipettes are Checked

When equipment is used for CRITICAL

Measurements

The calibration of the equipment must be checked!

In this case with qPCR the pipettes used to aliquot the samples where verified to be functioning properly

Table 2. 10 µL Pipette Calibration

0.025

Electro	nic Pipet	G010134E	10 uL
	Vol		
1	9.98		
2	9.946		
з	9.986		
4	9.94		
5	9.99		
6	10.019		
7	9.985		
8	9.95		
9	9.975		
		Setpoint	
Avg	9.975	-0.25 %	
SD	0.025		
CV	0.253	6	

The 2 µL is within -1.8 % of the setpoint

The 10 µL is within -0.25 % of the setpoint





Definition of RM vs CRM (SRM at NIST)

reference material RM

material, sufficiently homogeneous and stable with reference to specified properties, which has been established to be fit for its intended use in **measurement** or in examination of **nominal properties** (JCGM 200: 2008)

certified reference material CRM

reference material, accompanied by documentation issued by an <u>*authoritative body*</u> and providing one or more specified property values with associated uncertainties and traceabilities, using valid procedures. (JCGM 200: 2008)

Standard Reference Material® (SRM®): A CRM issued by NIST that also meets additional NIST certification criteria. (NIST SP 260-136: 2000)







Definition of SRM Materials

Certified Reference Materials from NIST that are:

- Well-characterized
- Using state-of-the-art measurement methods and/or technologies
- For the determination of chemical composition and/or physical properties

NIST Special Publication 260-136 "Definitions of Terms and Modes Used at NIST for Value-Assignment of Reference Materials for Chemical Measurements

Three Possible Data Quality Descriptors

- NIST Certified Value
 - Highest confidence in its accuracy
 - All known sources of bias have been investigated
- NIST Reference Value
 - Best estimate of the true value
 - All known sources of bias have not been fully investigated
- NIST Information Value
 - Value will be of interest and use
 - Insufficient information is available to assess uncertainty

NIST Special Publication 260-136 "Definitions of Terms and Modes Used at NIST for Value-Assignment of Reference Materials for Chemical Measurements





Modes Used at NIST for Value-Assignment of Reference Materials for Chemical Measurements to Obtain a NIST Certified Value

- Certification <u>at NIST</u> Using a Single Primary Method with Confirmation by Other Method(s)
- Certification <u>at NIST</u> Using Two Independent Critically-Evaluated Methods
- Certification/Value-Assignment Using One Method <u>at NIST</u> and Different Methods by Outside Collaborating Laboratories

NIST Special Publication 260-136 "Definitions of Terms and Modes Used at NIST for Value-Assignment of Reference Materials for Chemical Measurements, Table 1 (p.2)





Steps Involved in Production

- Tubes need to be purchased
- Boxes need to be put together
- Labels need to be ordered from the SRM Office for each component and the box (6 wk lead time)
- Samples need to be dispensed into separate tubes as uniformly as possible to maintain homogeneity
- Homogeneity studies need to be performed
- Data collected will be evaluated by a statistician from the NIST Statistical Engineering Division





Bottling SRM 2372 Materials



Teflon container holding ≈ 250 mL of *Candidate* SRM 2372. *It's not an SRM until it passes all testing.*

With a multi-channel pipettor 8 tubes can be filled at a time. That's \approx 214 reps to fill 1700 tubes per component.



The assembly line closing the recently filled tubes

Genetics



Protecting the SRM Product from the Staff : Lab Coats, Masks and Hair nets or full face shields **P**ersonal **P**rotective **E**quipment (PPE) or **P**roduct **P**rotective **E**quipment.





Safety Considerations: The Blister Brigade



Closing the 1,700 component A tubes (SRM 2372) caused some blisters even while wearing gloves. **Safety resolution:** Band-aids applied prior to closing SRM component tubes the next session helped reduce the number of blisters formed!

Applied Genetics

World Health Organization (WHO)



http://www.who.int

National Institute for Biological Standards and Control (NIBSC)

A centre of the Health	BSC Protection Agency	Health Protection Agency	http:	//www.	nibsc.a	Site sea	arc
About Us	Products	Services	Science	Spotlight	Partners	Contact	

Home Page > Products > Product Catalogues > Biological Reference Materials

Biological Reference Materials	Biological Reference Materials
ECBS established products 2009 A-Z Listing Search Catalogue	You can browse products by subject area by selecting the relevant link below. Products can then be ordered by type of standard or product code.
Return to previous page	Endocrinological Bacteriological Virological Cytokine, Growth Factor, Allergens, Sera, Miscellaneous Blood Products
	Haemostasis, Thrombosis, Transfusion Science, Immunohaematology Retroviral

http://www.nibsc.ac.uk/products/biological_reference_materials.aspx







National Institute for Biological Standards and Control

- The World Health Organization established a program in biological reference materials with the intended use for vaccine production, immunological and biological assays.
 NIBSC provides the International Standards (IS)
- An important motivation for this work was the safety of the blood supply
- They were initially designed to be qualitative standards
- The first WHO International Standard, for HCV, was released in 1997
- IS include: HCV, HBV DNA, HIV1 RNA, HAV, HPV type 16 and 18, Parvovirus B19, CMV





NIST and NIBSC have different models for reference materials production

NIBSC

- Clinical material or whole cultured virus
- Consensus evaluation,
 assay dependent value
- Quantity expressed as
 International Units
 unique to that material
- No uncertainty

NIST

- Pure viral DNA
- Independent RM
- Units are genome copy number per volume, traceable to the International system of units or SI
- Reported uncertainty





WHO International Standards - IU/mL

- An International Standard is a collection of ampoules containing as far as is possible the same amount of the analyte in as stable a form as possible (e.g lyophilised, low moisture, low oxygen, stored at -20°C)
- The material has a stable quantifiable biological activity. It is similar to the real life analyte.
- Stability studies are carried out before it is established and real time stability is monitored
- The lifetime is unspecified and depends on data or need. Replacement where necessary is an issue.
- The unit is arbitrarily defined as a fraction of the contents of an ampoule; when the collection of ampoules is exhausted the replacement is chosen to be as close to the old one as possible but there is uncertainty and strictly speaking the unit changes.
- Information on this and the next few slides provided by Phil Minor, Head of Virology, NIBSC



Process for establishing WHO International Standards

- Identify need for standard, including requirements for source material and formulation (in consultation with experts within the field)
- Proposal for development of standard presented to WHO ECBS for adoption into standardization program
- Source and evaluate candidate materials, process development (trial fills)
- Prepare candidate standard (definitive fills)
- Launch worldwide collaborative study to evaluate candidate standard, samples are sent to a number of high quality laboratories for their analysis
- Data return and statistical analysis
- Prepare report for approval by study participants
- Present report to WHO ECBS for establishment of standard (candidate standard assigned concentration in International Units (IU), usually based on the mean titer determined in the collaborative study)
- NIBSC acts as the custodian and worldwide distributor of established WHO International Standards



Summary of NIST & NIBSC/WHO

- Two standards developing organizations are working on primary reference materials for molecular diagnostics for infectious diseases with a current focus on viral infections of immune-compromised patients
- The organizations use two different models in the development of these reference materials
- NIST provides SRMs and NIBSC/WHO provide International Standards
- In the case of CMV, the NIST material is pure viral DNA (Towne strain) in buffer and the NIBSC material will be lyophilized intact virus (Merlin strain)
- The SRM will be certified for sequence and quantity traceable to the SI and with calculated uncertainties
- The IS will have an assigned unitage that is derived from a collaborative study consensus value
- The proposed standards are to be used as primary standards to calibrate secondary standards that will be subsequently used as calibrants





Clinical DNA SRMs: Current & Future






GO

GO

Clearly we need to improve our SRM offerings AND search engine! Material Measurement Laboratory Standard Reference Materials

Search Results

November 12, 2010

SRM/RM Number:		GO
Keywords:	molecular genetics	GO

D - Detail T - Table C - Certificate

Archived Certificates

Full Certificate Archive >>

	SRM	Description
С	SRM 705	Polystyrene (Narrow Molecular Weight Distribution), February 1, 1963
С	SRM 705	Polystyrene (Narrow Molecular Weight Distribution), November 6, 1969
С	SRM 706	Polystyrene (Broad Molecular Weight Distribution), April 17, 1995





Current NIST DNA Reference Materials

Forensic Applications

Date of release or certificate revision (r)

- STR PCR DNA Profiling (SRM 2391b) 1995, r2008
- Mitochondrial DNA Sequencing (SRM 2392-I, 2392) 1999, 2003
- Human Y-Chromosome DNA Profiling (SRM 2395) 2003, r2008
- RFLP DNA Profiling (SRM 2390) 1992, r2001, now obsolete

Clinical Applications

- Fragile X Human DNA Triplet Repeat (SRM 2399) 2004, r 2007
- Huntington's Disease CAG Repeats (SRM 2393)
- Cytomegalovirus (CMV) Copy Number & Sequence (SRM 2366)

Platform Testing

- Human DNA Quantitation (SRM 2372) 2007
- Heteroplasmic mtDNA Mutation Detection (SRM 2394) 2004
- DNA Sequence Library for External RNA Controls (SRM 2374)

Several in development...

S





SRM 2374 – DNA Sequence Library for External RNA Controls

Work by another group at NIST





External RNA Control Consortium (ERCC)

- NIST SRM to contain 96 unique control sequences inserted in common plasmid DNA
 - engineered to be readily *in vitro* transcribed to make RNA controls
 - RNA controls intended to mimic mammalian mRNA
- Developed *sequence library* from submission by ERCC members, as well as synthesis
 - evaluated performance of RNA controls on variety of platforms
 - selected 96 well-performing sequences in collaborative study
 - from library of 176
- Array manufacturers modified products to include SRM sequences
- Prepared 400 units of SRM
 - 96 tubes in each





SRM 2393 HD Alleles

- Alleles included in the SRM are:
- 15, 17, 29, 35, 36, 39, 40, 45, 50, & 75







NIST SRM projects for molecular diagnostics of infectious disease

Cytomegalovirus (CMV) – under development

Epstein-Barr virus and BK virus – in the planning stages





Who are our customers for infectious disease standards?

- Reagent and calibrant manufacturers
- Clinical laboratories that prepare their own calibrants
- NIST standards should be primary standards traceable to the SI
- NIST standards should be used to establish traceability for secondary standards/calibrants
- A central reference point to link the standards in use in clinical laboratories

Improved measurements through better calibration should lead to reduced variability among clinical labs







CMV Standard Reference Material

Type of material - pure viral DNA in buffer

- CMV DNA, Towne strain, cloned into a bacterial artificial chromosome (Towne $\Delta 147$)
 - Developed by Dr. Hua Zhu, New Jersey Medical School
- The BAC/CMV DNA is essentially a very large plasmid 240,000 bp, which is propagated and purified by a process that is similar to plasmids
- Advantages Consistent genome size and ease of propagation





∆ 147 Towne BAC

- BAC DNA is considered stable and can accommodate up to 300,000 base pairs
- DNA has been removed (20,000 base pairs) from the CMV viral genome to accommodate the BAC related DNA
- DNA that has been removed is <u>not</u> in the regions used for amplification





Properties Being Certified

- DNA sequence of regions of the CMV genome that are used as targets of PCR amplification
 - The following regions have been Sanger sequenced UL34, UL54, UL55-56, UL80, UL83, UL97, UL122-126, UL132, & US17
 - Sequence to date matches GenBank Towne strain sequence
- Copy number by direct measurement using digital PCR





CMV DNA Sequence Characterization



Regions sequenced

http://www.nist.gov/mml/biochemical/genetics/cmv_structure.cfm

Schematic map of the CMV genome. The CMV genome is organized as two regions of unique sequences, unique long (UL) and unique short (US), flanked by two sets of inverted repeats (light shaded boxes). *Kotenko et al. (2000) PNAS 97(4): 1695-1700*

Sequence alignments from available GenBank CMV strains being compared against published PCR primers and probes – this information is being included on the NIST Clinical DNA Information Resource website

http://www.nist.gov/mml/biochemical/genetics/clinical_dna.cfm



Sanger sequence result obtained at NIST from UL54 (section shown is the reverse primer region for the CP1 assay used in digital PCR and matches the Towne strain and published primer)





Certification of the CMV DNA genome copy counting

Digital PCR – Quantify the amount of DNA (copies/volume) by counting amplification from single molecules

- Nano scale reactions (6 nL)
- DNA concentration where some reaction chambers are negative
- $_{\odot}\,$ Based on Poisson statistics, number of copies is determined
- Thousands of replications/assay repeated with multiple assays targeting regions on the CMV genome
- Traceable to the SI via the Mole
- New tool, with active research at other National Metrology Institutes to validate this approach





NIST Digital PCR Instrument









765 individual chambers / panel 12 panels/chip



S



Preparation of the SRM

- Three levels of DNA concentration in buffer
- Packaging in Teflon tubes
- Certification for copy number/volume will be done on each of the concentrations
- Monitoring for homogeneity and stability for the life of the SRM (5 years from time of issue)
 - We currently have a stock that we have been monitoring for 18 months at three temperatures



Variability in Digital PCR Measurements Across Different CMV Targets (all three candidate components for SRM 2366)





Variability in Digital PCR Measurements Across Different CMV Targets (single candidate component of SRM 2366)



TaqMan Assay

ANOVA of the data showed no significant differences between groups

Average across all groups						
Average SD	2118 150	copies/μL copies/μL				
Uk=2	300	copies/µL				





QCMD CMV EQA 2010 Inter-laboratory Study

- NIST collaborated with Quality Control for Molecular Diagnostics (QCMD) for an inter-laboratory study conducted as part of the QCMD CMV 2010 EQA program
- NIST provided aliquots of CMV DNA, component B of the candidate CMV SRM, to QCMD
- A vial of DNA was sent out to each participating laboratory along with the 10 QCMD samples (lyophilized virus in plasma or VTM)
- While the QCMD samples required extraction, the NIST DNA was to be added directly to the assay. Participants were asked to run the assay in triplicate and report results in copies / mL.
 - 181 data sets were submitted





QCMD Data Analyzed by Assay Type

	# Data	<mark>Log₁₀ (copies/m</mark>		
QCMD CMV EQA - Participants and assays	sets	Median	MADe	
Total Datasets	181	5.900	0.486	
Conventional Commercial	5	5.854	0.872	
Real-Time Laboratory developed - Total	78	6.002	0.650	
Real-Time Commercial - Total	96	5.826	0.451	
Argene CMV HHV6,7,8 R-gene	6	5.864	0.150	
Argene CMV R-gene	15	6.205	0.332	
Nanogen Q-CMV Real time Complete Kit	21	5.733	0.794	
QIAGEN artus CMV PCR Kit (RG, LC,TM)	28	5.821	0.326	
Roche LightCycler CMV Quant Kit	12	5.776	0.298	







Information Resources for the Clinical Genetics/Healthcare Community

http://www.nist.gov/mml/biochemical/genetics/clinical_dna.cfm

http://www.nist.gov/mml/biochemical/genetics/index.cfm



Clinical DNA Informational Resource (CDIR)

http://www.nist.gov/mml/biochemical/genetics/clinical_dna.cfm

- In 2009, we started an internet based information resource – the goal is to provide information on DNA sequence alignments and primer and probes for clinical assays as a start
- We plan to load this resource with accessible and useful information for the clinical diagnostics community
- We welcome input from you on additions and changes to this information resource



Clinical DNA Informational Resource Website

http://www.nist.gov/mml/biochemical/genetics/clinical_dna.cfm



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Date created: November 3, 2009 | Last updated: April 7, 2010 Contact: CSTL Webmaster

Sequence Alignments for CMV Strains

🕖 SeqB 🙆 File Selectio	uilder - [AY315197 p Edit Features Enzym on: Top strand 7839	<mark>cr targets(cut).s</mark> mes Sequence (3 78464 length	bd] Cloning Format V 1 = 72	iew Net-Search V	Vindow Help					_□× _₽× 221 987 kh
	10	20	30	40	50	60	70	80	90	
					For Seq	Pol 2	·····			
0										
5'	ccggcctcgta	gtgaaaatta	atggtgttgaa	cagategeges	ccaatacgg	cgtcctgcagac	agtaacggcct	tacctgggcg	cggccc	
0	+++++++++++++++++++++++++++++++++++++++			***	* * * * * * * *	* * * * * * * * *	+ • • • • + • • • • • •		 ++++ 7	8480
5	ggccggagcat			gtetagegegt	ggttatgeeg	jeaggaegtetg	rcattgeegg:	atggacccgc	gccggg	
0	•	SMV CP1-r		CMV CP1-IM	a Rol 2		UP14			
			<		iq roi z					
0										
5	teggeattage	cacgaaacaa	cgcggggatgtc	cttgtaggaca	iggtcatccti	-gegttgeegea	liggtaaageteg	ggccatagtg	ttgagc Luurul 7	8570
3'	agccgtaatcg	atactttatt	reaccetacad	maacateetot	ccautauga:		ccatttcgag	contateac	aactco	0370
0		9-999		9						
TT See	nMan - [Alignment of	Contia 1]								
PP Fil	e Edit Sequence C	ontia Project SN	IP Net Search W	/indow Help						
ÐF	osition: 14193	4							2	222.066kb
			143,5	40 143,550	143,560	143,570	143580 14	3590 143	600 143	610
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皇	BK000394.seq			ccgttactgtet(lcaddacdccdi	tattggtgegega tattggtgegega	tetetteaaeae tetetteaaeae	cattaattttc: cattaattttc:	actacgagge actacgagge	cgggggccat
	X17403.seq	AD169	gtagg	ccgttactgtct	gcaggacgccgt	attggtgcgcga	tetgtteaacae	cattaattttc	actacgaggc	cggggccat
	AC146905.seq	Toledo	→ gtagg	ccgttactgtct	gcaggacgccgt	attggtgcgcga	tetgtteaacae	cattaattttc	actacgaggc	eggggecat
	NC_006273.seq	Merlin	🕂 🤆 gtagg	ccgttactgtct(gcaggacgccgt	attggtgcgcga	tetgtteaacae	cattaattttc	actacgaggc	cggggccat
	MB320634 apr	(clinical)		ccgttactgtct	gcaggacgccgt		tetgtteaacae	cattaattttc:	actacgaggc	cgggggccat
	AD329034.864	(cinneal)	/ gcagg	eegilaeigiei	geaggaegeegi	lattggtgegega	cergereaacae	callaallics	actacyayyc	uggggeeac
			-							-
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X



Sequences from 14 CMV Strains

Sequence alignments

The qPCR assay below (Tanaka 2000 Journal of Medical Virology 60:455–462) has a large number of mismatches that cause a false negative rate of 24%. A little more time in the library, probably would have lead this researcher to redesign the primer and probe binding sites.









- We are asking for your input on what we are currently doing and the relevance to the clinical community
- This is the first time that NIST has worked on standards for clinical diagnostics of <u>infectious</u> disease – we want to get it right
- Please come and talk to us during the meeting: we have three posters G09, ID19, and OTH05.



NIST Presentations at AMP 2010



Margaret Kline



Marcia Holden



Ross Haynes

- G09 (poster): Characterizing the Electrophoretic Mobility of Huntington Disease Alleles 13-77: Are the Deviations from the Theoretical Values Intrinsic or Internal Sizing Standard Artifacts?
- ID19 (poster): NIST Candidate Standard Reference Material (SRM): Cytomegalovirus DNA
- OTH05 (poster): Cautionary Considerations when Exploring Cell Lines as Potential Reference Materials





Contact Information



<u>Group Members Working on</u> <u>Clinical DNA Projects</u>

- Marcia Holden (x4162)
- Margaret Kline (x3134)
- Ross Haynes (x4469)
- John Butler (x4049) Group Leader

Email: first name . last name @ nist.gov Phone: 301-975-x

http://www.nist.gov/mml/biochemical/genetics/clinical_dna.cfm





Your

help

needed

Pipeline for SRM Production

- I. Request for SRM and internal decision to go forward
- 2. Understanding needs and scope of SRM (research)
- 3. SRM design

what formats and amounts are best?

- 4. Gather materials for SRM components
- 5. Characterization of components

Measure amounts with uncertainties, homogeneity, stability

- Complete paperwork
 - Report of Analysis (ROA) and Certificate of Analysis
- 7. Paperwork reviewed and SRM approved
- 8. SRM released (http://www.nist.gov/srm)
- 9. Customers use SRM and provide feedback

SD (standards development)

WCF (working capital funds)

Association for Molecular Patholog

SRM office



Reference Materials Requested

in Karen Mann's Congressional Testimony (Feb 24, 2010)

- Immediate
 - CMV
 - BCR/ABL
 - KRAS
 - EGFR
- Short term
 - BK virus
 - Epstein Barr virus
- Medium term
 - Certified Gene Sequence Databases
 - Infectious agents: adenovirus, enterovirus, Hepatitis B virus, Herpes simplex, JC virus, …

Source: http://www.amp.org







- Materials being characterized for sequence and amount
 - Material: Towne strain
 - Components: 3 vials of liquid DNA containing the same sequence but different concentrations









• Specific Needs:

- What is the measurement issue?

SRM Design:

- What format (e.g., genomic DNA, amount)?
- How many variants (e.g., different types)?
- How many components desired?

Source of Materials:

 Who can supply appropriate starting material of sufficient quantity for SRM production?





KRAS

• Specific Needs:

- What is the measurement issue?

• SRM Design:

- What format (e.g., genomic DNA, amount)?
- How many variants (e.g., different types)?
- How many components desired?

Source of Materials:

 Who can supply appropriate starting material of sufficient quantity for SRM production?







• Specific Needs:

- What is the measurement issue?

SRM Design:

- What format (e.g., genomic DNA, amount)?
- How many variants (e.g., different types)?
- How many components desired?

Source of Materials:

 Who can supply appropriate starting material of sufficient quantity for SRM production?

