

A network diagram with a blue background and white lines connecting nodes. Several circular icons are scattered across the network, including a biohazard symbol, a padlock, a map of the United States, a building, a hand holding a globe, and a biohazard symbol with a cross.

# Rapid DNA

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DHS Science & Technology Directorate



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# Rapid DNA Overview

## Reduces Multi-million Dollar Laboratory Processes to One Field Device

- Integration of five forensic lab processes with disposable microfluidic technology.
- Automation allows DHS officers to process samples and receive final results.
- Two U.S. small businesses have commercial devices ready for purchase.



10 hours processing  
reduced to 90 min.



**Desktop  
Rapid-DNA Unit**

40% sample cost reduction.  
System \$250K vs. \$1M lab.



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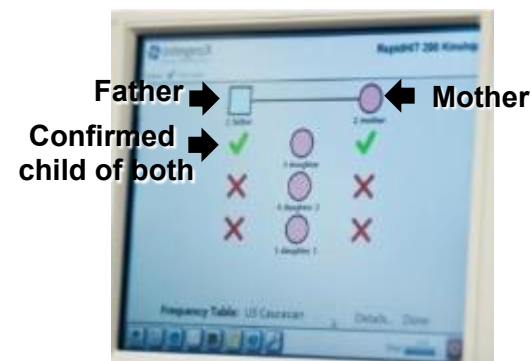
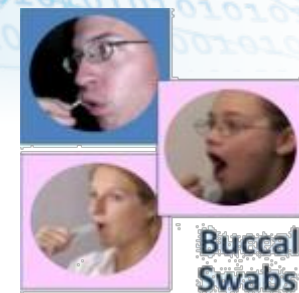
# Fully Integrated & Automated

## Enables the Decision Maker:

- A new and better capability – Only biometric that verifies family relationships at greater than a 99.5% probability.
- Unique capability – Provides ability in the field to check against criminal and known or suspected terrorist (KST) DNA databases.
- Analyzes trace evidence: (i.e., degraded bodies, bones, IED twist ties, tooth brush).
- Ready for field and officer use: ruggedized with an easy interface.
- Compatible with:
  - FBI, Interpol, ANSI/NIST and AABB data formats & reporting standards.

## Provides Screening at Speed:

- Microfluidics speeds processes and reduces costs.
- 'Privacy by Design' protection built into the system.
- No special training – fully automated DNA/kinship analysis.
- Disposable sample kit avoids run to run contamination.



Results screen showing family relationships



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# DNA Is The Most Powerful Biometric

More Effective, Informative, And Accurate

- Based on Objective Scientific Principles
- Supported by Statistics, Not Proprietary Algorithms
- Does Not Change Over Time
- Not Subject to Age Restrictions and Reliable After Death
- Effective in Trace Amounts
- Small Data Storage Footprint
- The ONLY Biometric to Verify Biological Relationships
- Privacy is protected by choosing DNA locations that do not reveal any physical traits, race, ethnicity, disease susceptibility, medical information or other sensitive information

INFORMATIVE

INCORRUPTIBLE

PERMANENT

KINSHIP



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# Potential DHS Applications

## Immigration:

- Enabling officers to rapidly screen applicant relationships:
  - During refugee interviews and against remote family members.
  - For those putting children up for overseas adoptions.
  - Against federal watch lists/criminal databases.



## Border and Enforcement:

- Countering human smuggling and trafficking by verifying relationships of suspicious persons and claimed families.
- Supporting investigations by linking objects to known persons.
- Screening arrested/detained persons against DNA watch lists.



## Disaster Recovery and Resilience:

- Medical Examiner daily use for morgue identifications reduces body storage costs and ensures medicolegal staff is ready to deploy.
- Helps resilient communities rapidly recover by identifying victims – even before decontamination – and reuniting families.
- FEMA grants can fund system and consumables purchase.



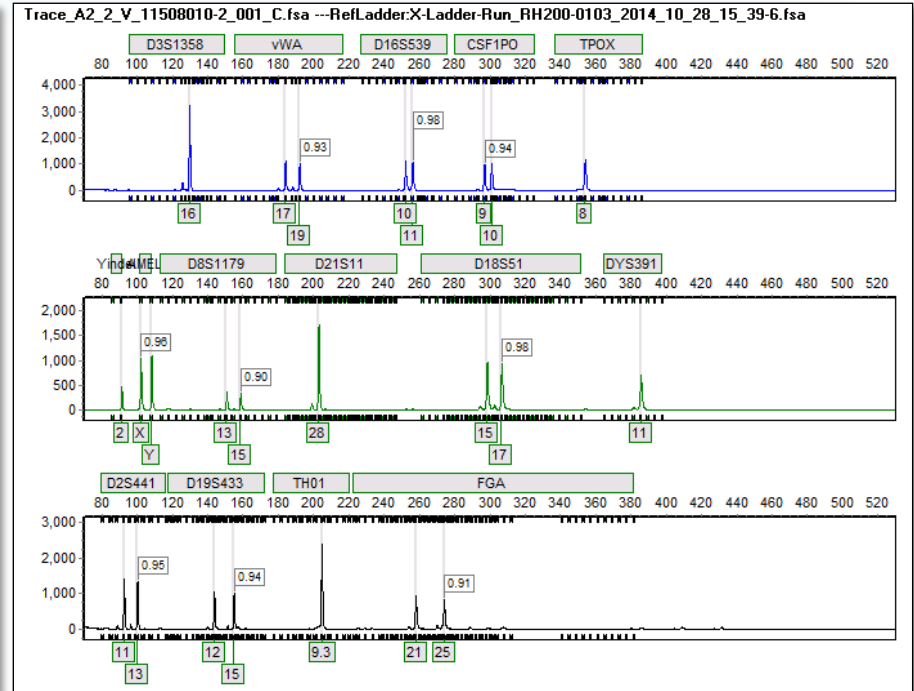
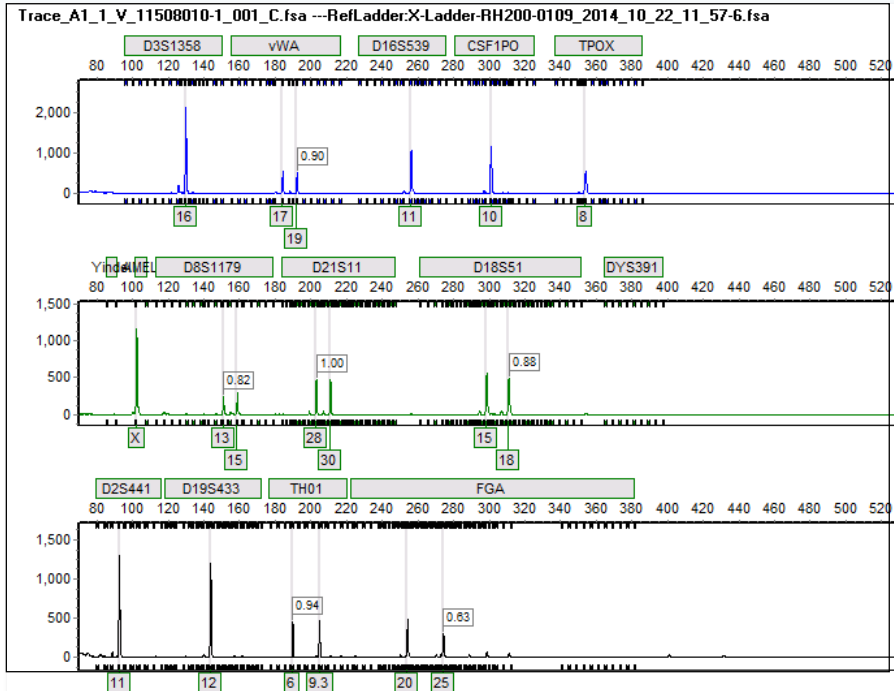
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# What a DNA Profile Looks Like

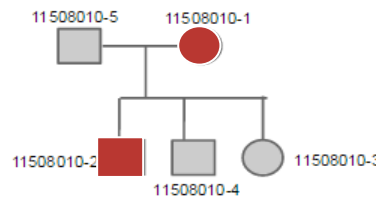
Mother (10-1)

Child (10-2)



DNA Profile: 16, 16; 17,19; 11, 11; 10,10; 8, 8;  
-, -; X, X; 13,15; 28, 30; 15, 18; -, -;  
11, 11; 12, 12; 6, 9.3; 20, 25

DNA Profile: 16, 16; 17,19; 10, 11; 9,10; 8, 8;  
2, 2; X, Y; 13,15; 28, 28; 15, 17; 11, 11;  
11, 13; 12, 15; 9.3, 9.3; 21, 25



Note: Only a portion of the profiles are shown

Probability of Maternity = 99.99999996%



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# Rapid-DNA Program Metrics

## Operational Requirements:

1. Fully automated swab to answer
2. Disposable consumables, pre-loaded with reagents and sealed
3. System cost  $\leq$  \$275K, (currently \$250K)
4. Cost/sample  $\leq$  \$100  
(NetBio currently \$270/sample)  
(IntegenX currently \$225/sample)
5. Ruggedized for transportation
6.  $\leq$  6 cu. ft., no side  $>$  30 in.,  $<$  110 lbs  
(IntegenX currently 180 lbs.)
7.  $<$ 1 hr of training for field user
8. Two person setup in  $\leq$  15 min
9. Reliability of system as specified  $\geq$  1 month routine support interval
10. No routine alignment or calibration
11. 120V, drawing  $<$  5 amps

## Sample Analysis:

1. Kinship Verification  $>$  99.5%
2. Answer in  $<$ 1 hr., Goal = 45 m  
(NetBio 82 min.)  
(IntegenX 115 min.)
3. Simultaneous processing of 5 samples, Goal = 15 samples  
(IntegenX capable of 7)
4. Reagent stable  $>$ 3 mo. at 20-300C, Goal = 6 mo. at -10-500C  
(NetBio  $>$ 9 mo. stability)  
(IntegenX = 6 mo. stability but must be refrigerated)
5. DNA extraction & purification comparable to lab methods
6. FBI CODIS compatibility and quality ( $\geq$  16 loci, single base-pair resolution, 500 bp length)
7. DHS Goal: 24 loci for kinship  
(NetBio currently 16 loci)  
(IntegenX currently 24 loci)
8. Processes fresh/dried buccal swabs & other DNA samples prepared manually

## Data Analysis:

1. Create & export profiles compatible with CODIS, ANSI/NIST, and expert systems
2. Raw & processed DNA profile data must be provided & stored
3. Bar code reader & GPS receiver that relay position & time to onboard computer
4. Generate data file for sample tracking with unique identifier information
5. Software: Windows XP, network connection capability, automated STR allele calling & profile generation, Comms, System control
6. ~~Wireless~~, wired, & USB network connections

Green = Demonstrated

Black = Not yet demonstrated

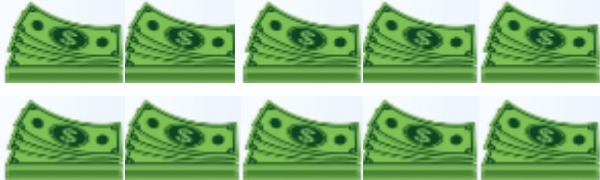
Red = Below target performance



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# Probability of Relationship Illustration

% Probability of Relationship	Likelihood of Claimed Relationship
50	 <p>Just as likely to be related to each other as to a random person</p>
99	 <p>100 times more likely to be related to each other than to a random person</p>
99.5	 <p>500 times more likely to be related to each other than to a random person</p>

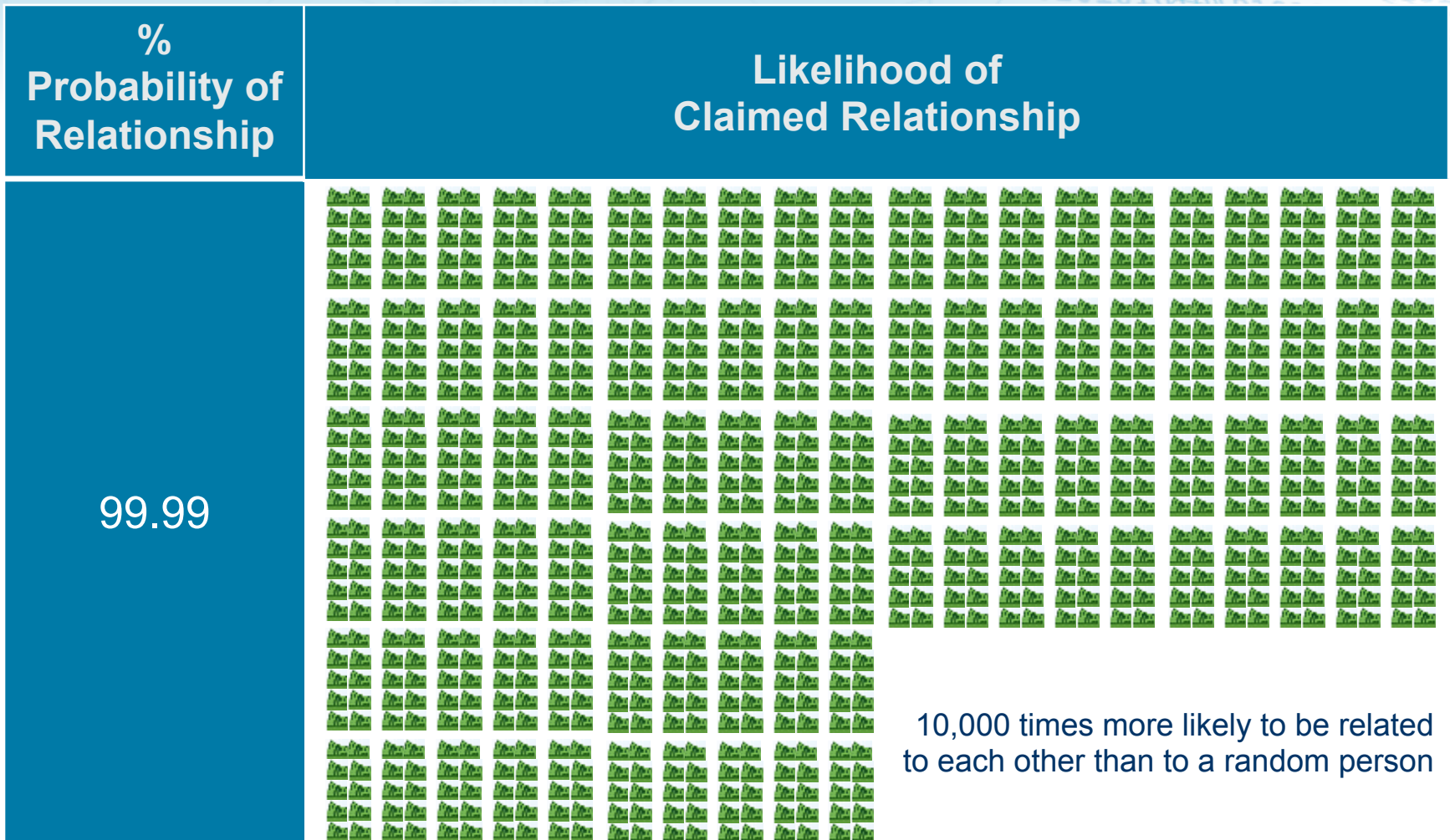


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# Probability of Relationship



10,000 times more likely to be related to each other than to a random person



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# Internal Validation for Kinship



## GENERATING PROFILES

- Evaluate that the Rapid DNA instrument produces accurate/reliable allele calls
- Evaluate that the Rapid DNA analysis software correctly identifies profile/data anomalies



## EXPORTING PROFILES


- Evaluate the DNA profiles export into the interpretation software correctly



## KINSHIP ANALYSIS

- Evaluate the kinship software generates & reports correct statistical calculations



	<b>Kinship Cases Exchanged with Accredited Laboratory</b>  <b>31 Samples</b>	<b>Allele Call Concordance Study</b>  <b>200 Determinations/ Electropherograms</b>	<b>Artifact/Anomaly Concordance Study</b>  <b>100 Electropherograms</b>	<b>Statistical Concordance Study Spanning DNA Locations/ Alleles/Formulae</b>  <b>~10 Cases</b>
Does Microfluidic Technology Produce Accurate/ Reliable DNA Profiles?	✓			
Does the Expert System Call Alleles Correctly?	✓	✓	✓	
Does the Expert System Accurately Identify Artifacts?		✓	✓	
Are Profiles Exported Properly Into the Kinship Software?	✓	✓	✓	✓
Does the Kinship Software Generate & Report Correct Statistics?	✓			✓

**Kinship Buccal Swab Set**

**DNA Profile Data Test Set**

**Kinship Data Test Set**



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# Kinship Buccal Validation Set

- 31 Donor Samples collected by Paternity Testing Corporation (PTC) Laboratories (an accredited AABB lab).
- Includes four family groups covering diverse biological associations (e.g. grandparent, aunt/uncle, half-siblings). See next slide.
- Six buccal swabs collected from each person and barcoded with a unique identifier to protect privacy.
- Samples tested at PTC and NIST using conventional DNA methods.
  - DNA data and kinship reports provided to DHS S&T.
- Results presented are from Rapid DNA platform
  - Conventional and Rapid DNA results were concordant.

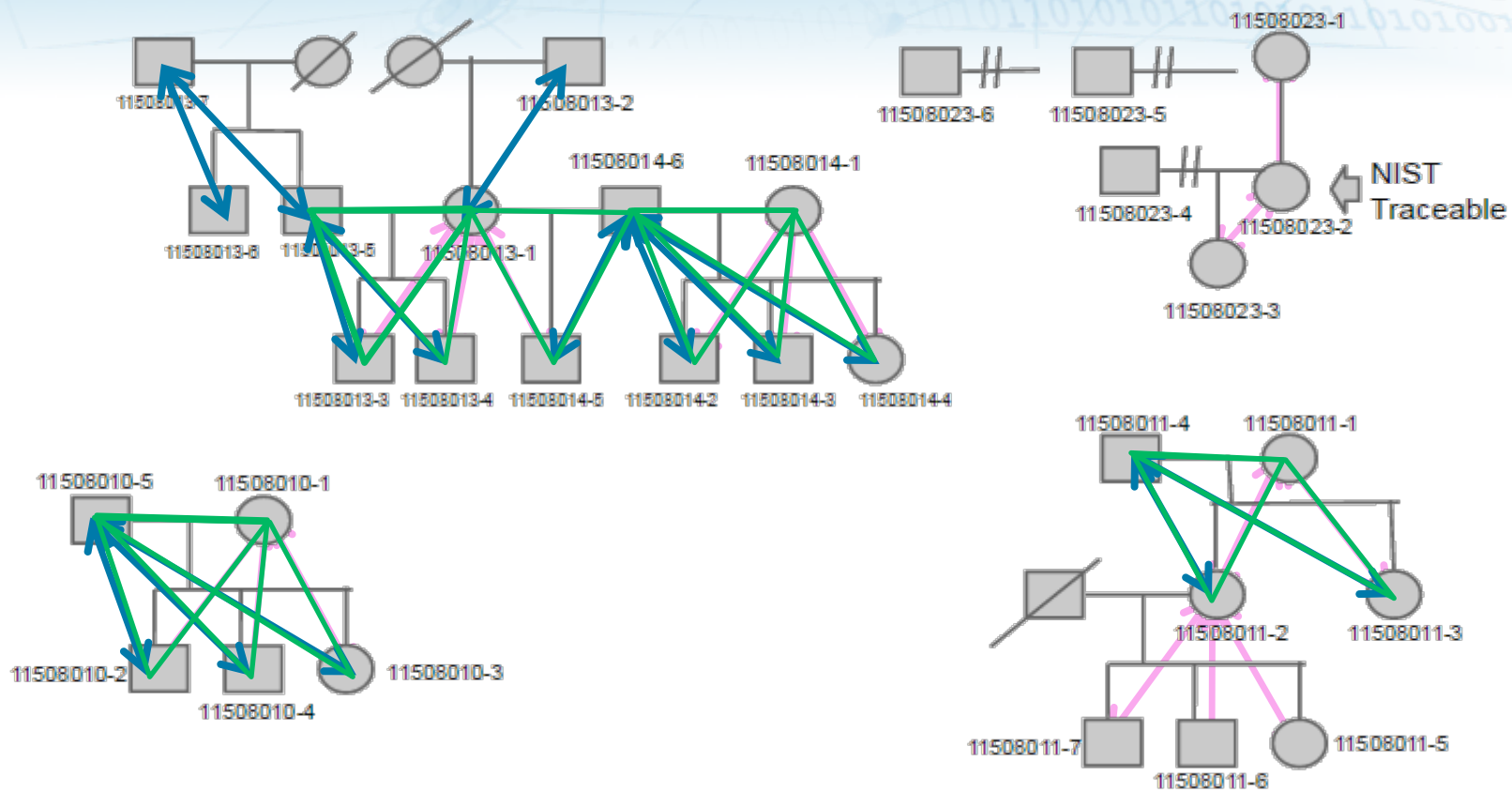


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# Kinship Validation Samples



- ↔ = 16 Mother to Child Relationship
- ↔ = 14 Father to Child Relationship
- ↔ = 11 Mother/Father/Child Relationships
- +12 Full Siblings, 5 Half Siblings,  
12 Grandparent/Grandchild, and  
5 Aunt/Uncle – Niece/Nephew



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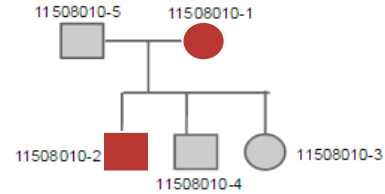
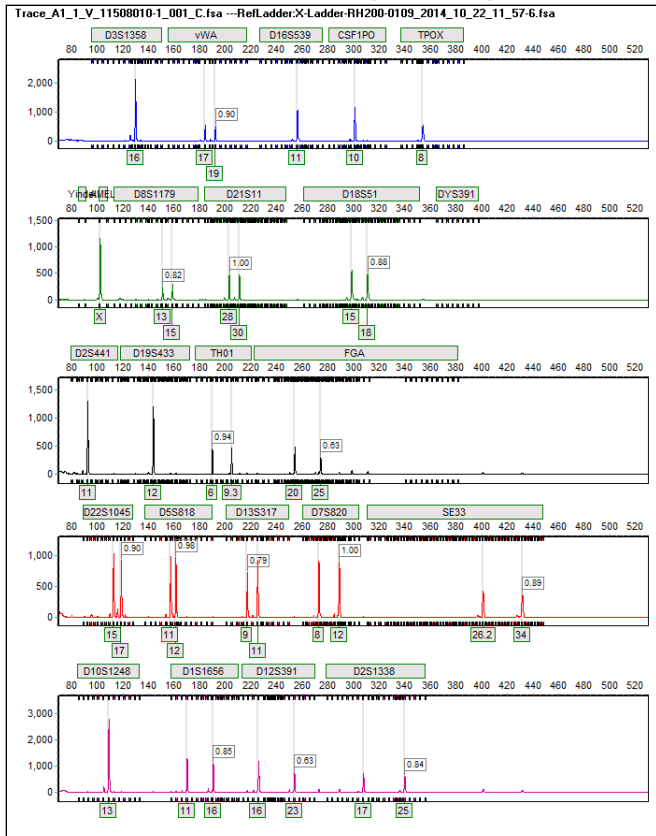
# Kinship Associations

Kinship	# Comparisons	# Kinship Evaluations
Mother – Child	16	42
Father – Child	14	
Grandmother – Grandchild	4	
Grandfather – Grandchild	8	
<i>Family Trio (Paternity)</i>	11	33
<i>Full Siblings</i>	12	
<i>Half Siblings</i>	5	
<i>Aunt/Uncle – Niece/Nephew</i>	5	

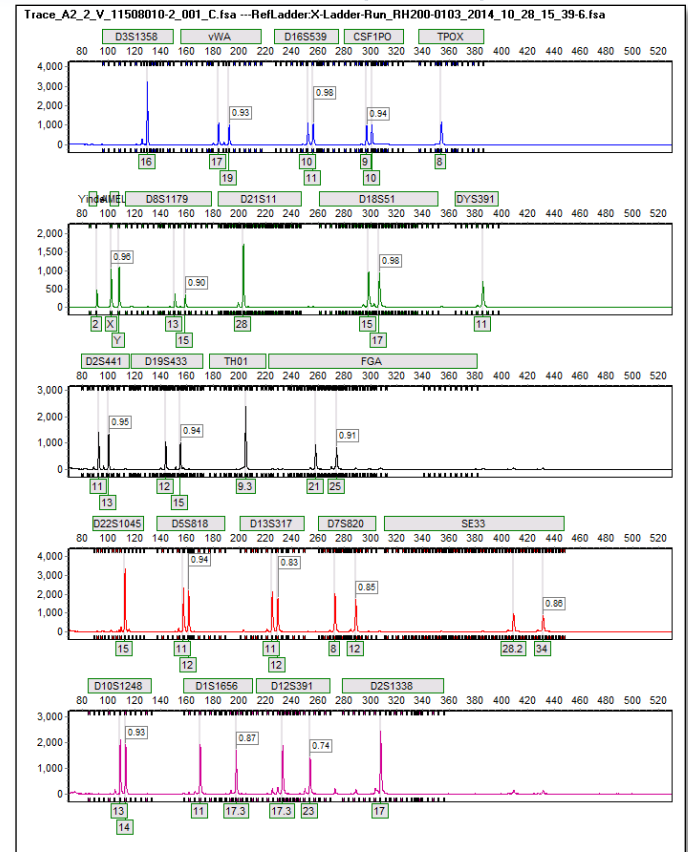


# Half of Child's DNA from Each Parent

Mother (10-1)



Child (10-2)



Probability of Maternity  
99.99999996%

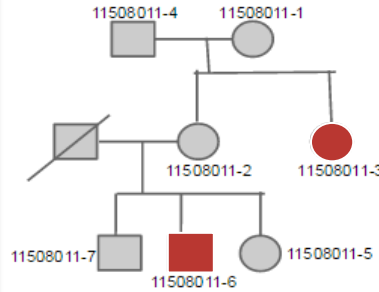
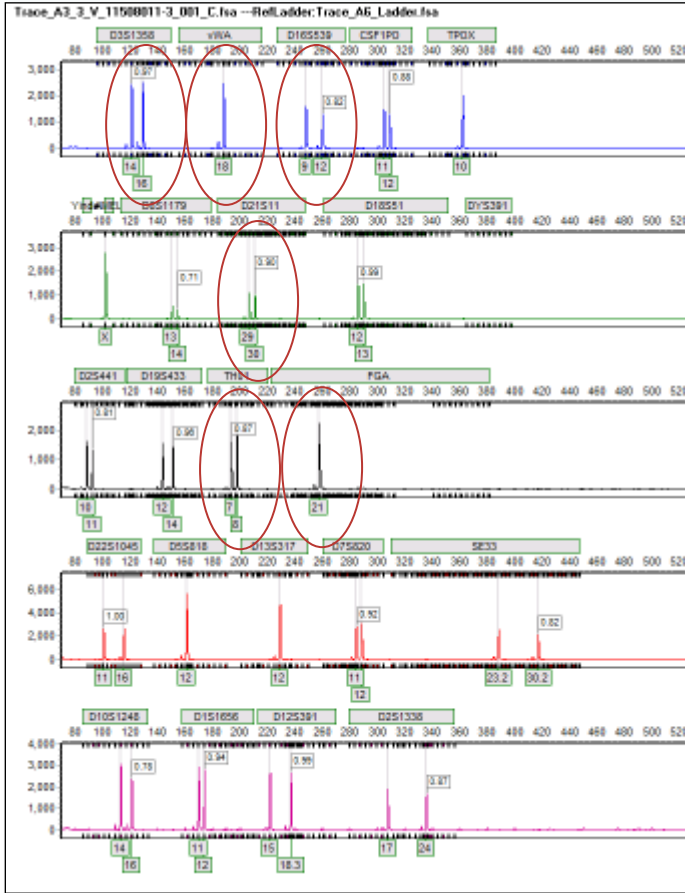


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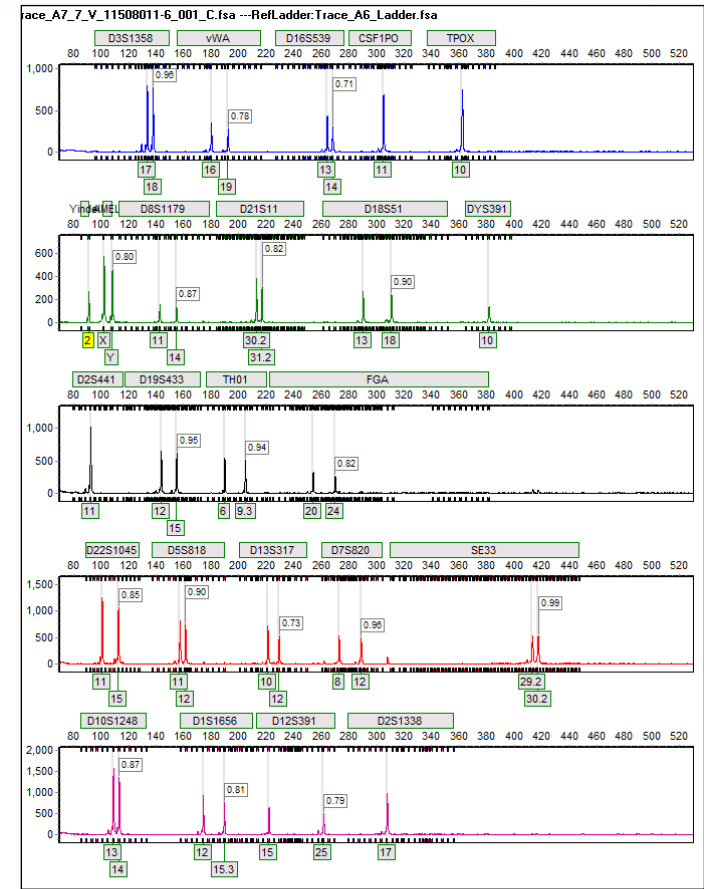
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# Aunt Can't Falsely Claim to be Mother

Aunt (11-3)



Child (11-6)



Excluded Match  
(Six locations  
don't match):

- D3
- vWA
- D16
- D21
- TH01
- FGA



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# DHS Kinship - True Claims

PTC Kinship Samples	# of tests	Average Relationship Result	Minimum Result
Mother – Child	16	99.999999997%	99.99995%
Father – Child	14	99.99999998%	99.99998%
<i>Family Trio (Paternity)</i>	11	99.999999998%	99.99998%
<i>Full Siblings</i>	12	99.999999997%	94.84%
<i>Half Siblings</i>	5	99.78%	95.53%
Grandmother – Grandchild	4	99.93%*	91.66%
Grandfather – Grandchild	8	98.93%*	8.12%
<i>Aunt/Uncle – Niece/Nephew</i>	5	99.21%	30.02%

\*When additional family is added for grandparent-grandchild relationships in 8 available tests, the average probability of relationship is 99.99997% with a low of 99.4%



# DHS Kinship - False Claims

PTC Kinship Samples False Claims	# of tests	Average Relationship Result	Maximum Result
Aunt – Child ( <i>Mother Claim</i> )	3	Exclusion	NA
Uncle – Child ( <i>Father Claim</i> )	2	Exclusion	NA
Aunt – Uncle – Child ( <i>Mother/Father/Child Claim</i> )	3	Exclusion	NA
Half Siblings ( <i>Full Siblings Claim</i> )	5	98.60%*	99.65%
Aunt – Niece ( <i>Grandmother – Grandchild Claim</i> )	3	98.34%**	98.48%
Uncle – Nephew ( <i>Grandfather – Grandchild Claim</i> )	2	99.52%	98.98%

\* False Half Siblings averaged 0.019% when additional siblings are added across 2 tests

\*\* False Grandmother-Grandchild with the mother included averaged 36.44% across 3 tests



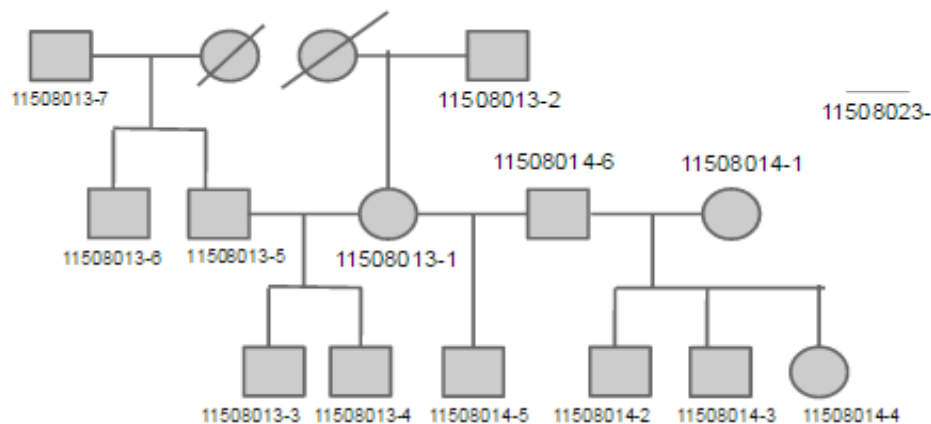
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# Comparing Population Databases

- Individual 20: 1/2 Asian 1/2 Caucasian (Father)
- Individual 23: 1/4 Asian and 3/4 Caucasian (Daughter)

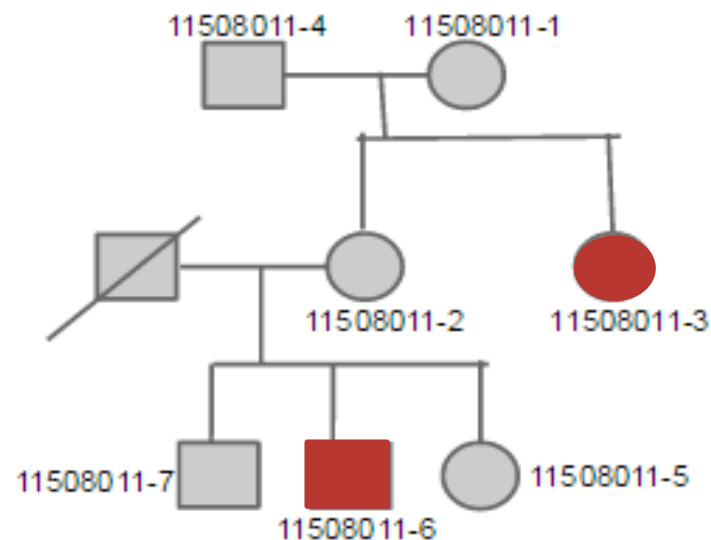
Population Group	Probability of Relationship
Asian	99.999998%
African American	99.999998%
Caucasian	99.9999996%
Hispanic	99.99999992%



# Comparing Population Databases

- Individual 9: Caucasian (Aunt)
- Individual 11: Caucasian (Nephew)

Population Group	Probability of Relationship
Caucasian	98.2%
African American	99.5%
Hispanic	99.3%
Asian	99.92%

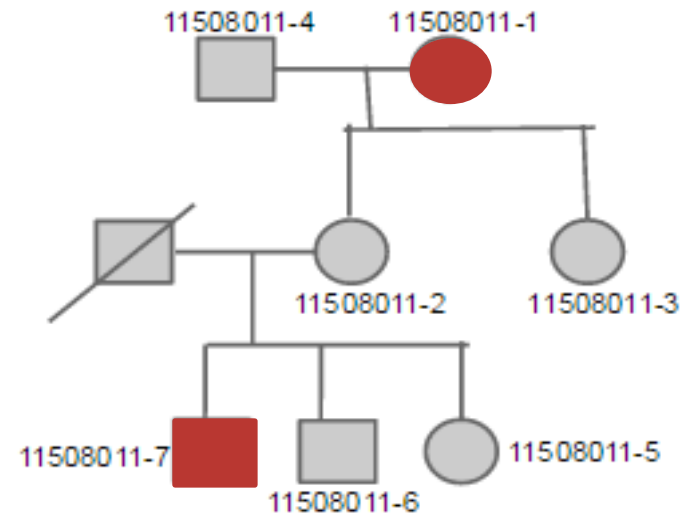




# Comparing Population Databases

- Individual 7: Caucasian (Grandmother)
- Individual 10: Caucasian (Grandson)

Population Group	Probability of Relationship
Caucasian	99.87%
Hispanic	99.95%
African American	99.992%
Asian	99.998%



# New Technology is More Robust

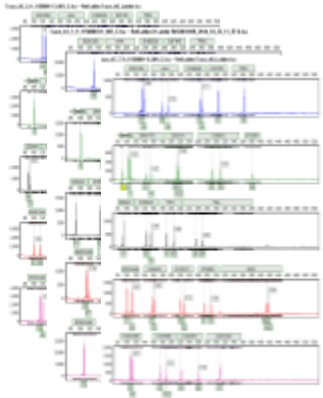
Relationship	# of tests	Average Probability of Relationship		
		CODIS 13 Loci	CODIS 20 Loci	24 Loci
Parent-Child	10	99.98%	99.99996%	99.999997%
Full Siblings	5	99.97%	99.99996%	99.999996%
Half Siblings	3	73.64%	73.1%	96.88%
Grandparent-Grandchild	7	74.10%	88.72%	95.88%
Aunt/Uncle-Niece/Nephew	3	23.18%	36.53%	74.82%





# DNA Profile Data Test Set

- S&T seeking to collaborate with Rapid DNA customers to build:
  - 200+ raw .fsa files and associated electropherograms
  - 100 profile anomalies
- Working with Rapid DNA vendors to ingest .fsa files into existing instrument pipeline
- Run every time expert software is updated/modified



Project: C:\RapidIT200\Results\Run\_RH200-0022\_2015\_10\_13\_09\_54\GM\_Analysis.sgf  
Panel: Globalfiler\_DKI  
Size: DY632  
Analysis Type: HID

	SE33	D10S1248
1 X-Ladder-Run_RH200-0109_2014_10_21_10_24-2.fsa		
2 Trace_A6_Ladder.fsa		
3 Trace_A1_1_20151013_10_13_09_54-fsa		
4 Trace_A2_2_20151013_10_13_09_54-fsa	14	28.2
5 Trace_A3_3_20151013_10_13_09_54-fsa	23.2	26.2
6 Trace_A4_4_20151013_10_13_09_54-fsa	14	23.2
7 Trace_A5_5_20151013_10_13_09_54-fsa	15	19
8 Trace_A7_7_20151013_10_13_09_54-fsa	14	14 **
9 Trace_A8_8_20151013_10_13_09_54-fsa	**	**



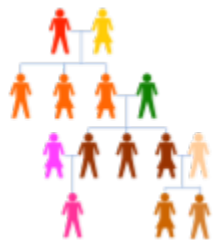
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# Kinship Data Test Set

- S&T developing 10+ artificial pedigrees (.cmf file format)
  - Comprised of biological relationship/kinship claims commonly encountered by USCIS and CBP (parent/child, grandparent/grandchild, siblings, etc)
  - Incorporates the 20 common paternity index formulas identified by AABB
  - Includes a wide number of alleles including off-ladder alleles and microvariants
  - Includes DNA profile complexities (e.g. parent-wrong gender, mutations, null alleles, and rare alleles)
- Enable quick and simple validation of Rapid DNA kinship software:
  - Verify allele frequencies & calculations
  - Verify correct reporting - inclusion/exclusion and probability of relationship
- Run every time expert software is updated/modified



STR Marker	Family Member	Allele	Probability	Phenotype
D3S1338	F1	15, 17	0.99	AC
D3S1338	F2	15, 17	0.99	AC
D3S1338	F3	15, 17	0.99	AC
D3S1338	F4	15, 17	0.99	AC
D3S1338	F5	15, 17	0.99	AC
D3S1338	F6	15, 17	0.99	AC
D3S1338	F7	15, 17	0.99	AC
D3S1338	F8	15, 17	0.99	AC
D3S1338	F9	15, 17	0.99	AC
D3S1338	F10	15, 17	0.99	AC
D3S1338	F11	15, 17	0.99	AC
D3S1338	F12	15, 17	0.99	AC
D3S1338	F13	15, 17	0.99	AC
D3S1338	F14	15, 17	0.99	AC
D3S1338	F15	15, 17	0.99	AC
D3S1338	F16	15, 17	0.99	AC
D3S1338	F17	15, 17	0.99	AC
D3S1338	F18	15, 17	0.99	AC
D3S1338	F19	15, 17	0.99	AC
D3S1338	F20	15, 17	0.99	AC



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# Technology is Field Ready!



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