

DNA Parentage Test No. 15-5870 Summary Report

This proficiency test was sent to 21 participants. Each participant received a sample pack consisting of the standard paternity trio, collected from a mother, daughter, and alleged father. Participants were requested to analyze the samples using their existing protocols. Data were returned from 19 participants (90% response rate) and are compiled into the following tables:

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This report contains the data received from the participants in this test. Since these participants are located in many countries around the world, and it is their option how the samples are to be used (e.g., training exercise, known or blind proficiency testing, research and development of new techniques, etc.), the results compiled in the Summary Report are not intended to be an overview of the quality of work performed in the profession and cannot be interpreted as such. The Summary Comments are included for the benefit of participants to assist with maintaining or enhancing the quality of their results. These comments are not intended to reflect the general state of the art within the profession.

Manufacturer's Information

Each sample set was a collection of known blood samples, provided on FTA Micro cards, from three individuals (Items 1-3), a mother, daughter and a potential father. Participants were requested to analyze these items using their existing protocols. Also included in the Data sheet was a kinship exercise that consisted of autosomal DNA profiles of two individuals for comparison. Participants were requested to determine, if a sibling relationship claim was supported following the review of these profiles.

SAMPLE PREPARATION: All stains were prepared from human whole blood which was drawn into EDTA tubes. Item 1 (75 μ I) was created using blood collected from a female (mother) donor, Item 2 (75 μ I) was from a female (daughter) donor and Item 3 (75 μ I) was created using blood collected from one male (the actual biological father) donor. Each different Item was prepared at separate times and were packaged once they were thoroughly dried. Completed sample sets were stored at -20°C until shipment on January 19, 2015.

SAMPLE SET ASSEMBLY: For each sample set, all three Items (1-3) were placed in a pre-labeled sample pack envelope. The sealed sample pack envelopes were then packaged in pre-labeled Heat Seal envelopes and sealed. This process was repeated until all of the sample sets were prepared.

KINSHIP EXERCISE: This exercise included allelic results representing a full sibling relationship.

VERIFICATION: Laboratories that conducted predistribution analysis of the samples reported consistent results and associations.

		Ame	logenin and S	STR Results						
	Results compiled by predistribution laboratories and a consensus of at least 10 participants.									
<u>Item</u>	D1S1656 D8S1179 D19S433 Penta D	D2S1338 D10S1248 D21S11 Penta E	D2S441 D12S391 D22S1045 SE33	D3S1358 D13S317 Amelogenin TH01	D55818 D165539 CSF1PO TPOX	D7S820 D18S51 FGA vWA				
1	* 10,16 14,14.2	19,20 * 30,31.2	* *	15,16 11,12 X	12,13 12 12	10 14 20,23				
	*	*	*	7,8	11,12	16				
2	* 10,17 14 *	19 * 30,31.2 *	* * *	15,16 11,12 X 7	11,13 10,12 11,12 11	10,11 12,14 23,24 16,17				
3	* 14,17 14 *	19,23 * 30,30.2 *	* * *	15 12 X,Y 6,7	11 10,11 10,11	11,13 12,18 24,25 17,18				

^{*} Results were not received by a minimum of 10 participants for STR loci indicated.

Summary Comments

The 15-5870 DNA Parentage test was designed to allow participants to assess their proficiency in the analysis and interpretation of a standard trio of blood stains on FTA Micro cards. Item 1 was blood collected from a female donor (mother), Item 2 was blood collected from a female donor (daughter) and Item 3 was blood collected from a male donor (father of Item 2). Participants were requested to analyze the samples and provide allelic and statistical results and relationship conclusions regarding the potential father. Sample sets also included a kinship exercise provided on the data sheets where participants were requested to evaluate the provided DNA profiles and report the kinship index and relationship conclusions. [Refer to the Manufacturer's Information for preparation details]

For Paternity DNA Statistics & Conclusions (Tables 5 & 6), a variety of responses were provided for the combined paternity index. However, all 19 responding participants reported that the potential father (Item 3) could not be excluded as the biological parent of the daughter (Item 2).

For Kinship DNA Statistics (Table 7), a variety of responses were provided for the kinship index. However, all 11 responding participants reported that the genetic evidence supported the claim of a full sibling relationship. Statistical responses for both paternity and kinship questions varied due to the use of different population databases. However, consistencies were noted amongst the indices reported from participants using the same database.

Only one participant reported allelic results that differed from the consensus/predistribution results. This participant reported inconsistent autosomal alleles for two loci in Item 3. A consensus of at least ten participants was not obtained for the YSTR allelic loci therefore no inconsistencies were determined.

Amelogenin & STR Results

TABLE 1

WebCode	Item	D1S1656 D8S1179 D19S433 Penta D	D2S1338 D10S1248 D21S11 Penta E	D2S441 D12S391 D22S1045 SE33	D3S1358 D13S317 Amelogenin TH01	D5S818 D16S539 CSF1PO TPOX	D7S820 D18S51 FGA vWA
				- Amelogenin 8			
4TWZUA	Powe	erPlex® 21	110111 1	7 unologomin c	2 0 110 10000113		
411120/1			10.00		15.17	10.10	10.10
	1	16,17.3	19,20	10.01	15,16	12,13	10,10
		10,16		19,21	11,12	12,12	14,14
		14,14.2	30,31.2		X,X	12,12	20,23
		10,11	5,18		7,8	11,12	16,16
7LCGP2	lden	tifiler®					
	1		19,20		15,16	12,13	10
		10,16			11,12	12	14
		14,14.2	30,31.2		X	12	20,23
					7,8	11,12	16
92EXNX	Iden	ifiler® plus kit					
	1		19,20		15,16	12,13	10,10
		10,16			11,12	12,12	14,14
		14,14.2	30,31.2		X,X	12,12	20,23
					7,8	11,12	16,16
97BL66	Powe	erPlex® 21					•
. 5200	1	16,17.3	19,20		15,16	12,13	10,10
	1	10,17.3	17,20	19,21	11,12	12,12	14,14
		14,14.2	30,31.2	19,21	X,X	12,12	20,23
		10,11	5,18		7,8	11,12	16,16
			3,10		7,0	11,12	10,10
4293HX		ifiler®					
	1		19,20		15,16	12,13	10
		10,16			11,12	12	14
		14,14.2	30,31.2		X	12	20,23
					7,8	11,12	16
ARL7GW	Powe	erPlex® PowerPlex	: 16 HS				
	1				15,16	12,13	10
		10,16			11,12	12	14
			30,31.2		Χ	12	20,23
		10,11	5,18		7,8	11,12	16
BUUGAU	lden	tifiler® Plus					
	1		19,20		15,16	12,13	10,10
		10,16			11,12	12,12	14,14
		14,14.2	30,31.2		X,X	12,12	20,23
					7,8	11,12	16,16
FWI P7Y	lden	tifiler® Plus					
WLP7Y			10.00		15,16	12,13	10,10
EWLP7Y	1		19.70				
EWLP7Y	1	10.16	19,20				
EWLP7Y	1	10,16 14,14.2	30,31.2		11,12 X,X	12,12 12,12	14,14 20,23

TABLE 1

IABLE I								
WebCode I	tem	D1\$1656 D8\$1179	D2S1338 D10S1248	D2S441 D12S391	D3S1358 D13S317	D5S818 D16S539	D7\$820 D18\$51	
		D19\$433	D21511	D22S1045	Amelogenin	CSF1PO	FGA	
		Penta D	Penta E	SE33	TH01	TPOX	vWA	
			Item 1	- Amelogenin 8	& STR Results			
KA7CHK	NGΛ	A SElect						
	1	16,17.3	19,20	12,14	15,16			
		10,16	14,15	19,21		12,12	14,14	
		14,14.2	30,31.2	15,17	X,X		20,23	
				17,24.2	7,8		16,16	
NWB3LP	Powe	erPlex® 21						
	1	16,17.3	19,20		15,16	12,13	10,10	
	•	10,16	.,,	19,21	11,12	12,12	14,14	
		14,14.2	30,31.2	,	X,X	12,12	20,23	
		10,11	5,18		7,8	11,12	16,16	
NZEZWG	Ident	tifiler® Plus	•		,	,		
,	1	CI © 1 103	19,20		15,16	12,13	10,10	
	ı	10,16	17,∠∪					
		10,16	30,31.2		11,12 X,X	12,12 12,12	14,14 20,23	
		17,14.4	50,51.2		7,8	11,12	16,16	
14.070.0		(r)			7,0	11,12	10,10	
JAG7DC		tifiler® Plus						
	1		19,20		15,16	12,13	10	
		10,16			11,12	12	14	
		14,14.2	30,31.2		X	12	20,23	
					7,8	11,12	16	
/89DWH	ldeni	tifiler® Plus						
	1		19,20		15,16	12,13	10	
		10,16			11,12	12	14	
		14,14.2	30,31.2		X,X	12	20,23	
					7,8	11,12	16	
H6JPG	Powe	erPlex® 21						
	1	16,17.3	19,20		15,16	12,13	10,10	
		10,16	.,,	19,21	11,12	12,12	14,14	
		14,14.2	30,31.2	, <u>, –</u> .	X,X	12,12	20,23	
		10,11	5,18		7,8	11,12	16,16	
NXJ4C	Ident	tifiler® Plus and D			·		*	
	1		19,20		15,16	12,13	10	
		10,16	17,20		11,12	12,13	14	
		14,14.2	30,31.2		X	12	20,23	
		,	30,31.2		7,8	11,12	16	
WN6AA	Da	erPlex® PP16HS			,,0	11,14	10	
MADNIVV		ennex® FF10H3			3537	10.10	1.0	
	1	10.17			15,16	12,13	10	
		10,16	20.21.2		11,12	12	14	
		10.11	30,31.2		X	12	20,23	
		10,11	5,18		7,8	11,12	16	
R8JD9	Powe	erPlex® HS						
	1				15,16	12,13	10	
		10,16			11,12	12	14	
			30,31.2		Χ	12	20,23	
		10,11	5,18		7,8	11,12	16	

DNA Parentage Test 15-5870 TABLE 1

			IADLL	I		
WebCode Item	D151656 D851179 D195433 Penta D	D2S1338 D10S1248 D21S11 Penta E	D2S441 D12S391 D22S1045 SE33	D3S1358 D13S317 Amelogenin TH01	D55818 D165539 CSF1PO TPOX	D7S820 D18S51 FGA vWA
		Item 1	- Amelogenin 8	& STR Results		
Z6822D Ide	entifiler® Plus					
1		19,20		15,16	12,13	10,10
	10,16			11,12	12,12	14,14
	14,14.2	30,31.2		X,X	12,12	20,23
				7,8	11,12	16,16
ZVFHN6 Ide	entifiler®					
1		19,20		15,16	12,13	10
	10,16			11,12	12	14
	14,14.2	30,31.2		Χ	12	20,23
				7,8	11,12	16

TABLE 1

				IABLE	<u> </u>		
WebCode	ltem	D1S1656 D8S1179 D19S433	D2S1338 D10S1248 D21S11	D2S441 D12S391 D22S1045	D3S1358 D13S317 Amelogenin	D5S818 D16S539 CSF1PO	D7S820 D18S51 FGA
		Penta D	Penta E	SE33	TH01	TPOX	vWA
				- Amelogenin 8			
4TWZUA	Powe	erPlex® 21		orogoriiii c	5111 11000113		
2071	2	10,17.3	19,19		15,16	11,13	10,11
	2	10,17.3	17,17	18,19	11,12	10,12	12,14
		14,14	30,31.2	10,17	X,X	11,12	23,24
		10,13	12,18		7,7	11,11	16,17
7LCGP2	Idon	tifiler®	12,10		, ,,	,	10,17
		iiiilei ®	10		15.17	11.10	10.11
	2	10.17	19		15,16	11,13	10,11
		10,17 14	20.21.0		11,12	10,12	12,14
		14	30,31.2		X	11,12	23,24
005\4."/					7	11	16,17
		tifiler® plus kit					
	2		19,19		15,16	11,13	10,11
		10,17			11,12	10,12	12,14
		14,14	30,31.2		X,X	11,12	23,24
					7,7	11,11	16,17
97BL66	Powe	erPlex® 21					
	2	10,17.3	19,19		15,16	11,13	10,11
		10,17		18,19	11,12	10,12	12,14
		14,14	30,31.2		X,X	11,12	23,24
		10,13	12,18		7,7	11,11	16,17
A293HX	Iden	tifiler®					
	2		19		15,16	11,13	10,11
	-	10,17	• •		11,12	10,12	12,14
		14	30,31.2		X	11,12	23,24
			,		7	11	16,17
ARL7GW	Powe	erPlex® PowerPlex	16 HS				
,	2				15,16	11,13	10,11
	2	10,17			11,12	10,12	12,14
		10,17	30,31.2		X	11,12	23,24
		10,13	12,18		7	11	16,17
BUUGAU	12	tifiler® Plus	12,10		,		10,17
DOOGAU		iiiilei w FiUS	10.10		15.17	11.10	10.11
	2	10.17	19,19		15,16	11,13	10,11
		10,17	20.21.0		11,12	10,12	12,14
		14,14	30,31.2		X,X	11,12	23,24
					7,7	11,11	16,17
EWLP7Y		tifiler® Plus					
	2		19,19		15,16	11,13	10,11
		10,17			11,12	10,12	12,14
		14,14	30,31.2		X,X	11,12	23,24
					7,7	11,11	16,17
KA7CHK I	NGA	M SElect					
	2	10,17.3	19,19	10,12	15,16		
		10,17	13,14	18,19		10,12	12,14
		14,14	30,31.2	15,17	X,X		23,24
		,	•	,	·		,

TABLE 1

			TABLE	<u> </u>		
WebCode Ite	m D1S1656 D8S1179 D19S433 Penta D	D2S1338 D10S1248 D21S11 Penta E	D2S441 D12S391 D22S1045 SE33	D3S1358 D13S317 Amelogenin TH01	D5S818 D16S539 CSF1PO TPOX	D7S820 D18S51 FGA vWA
	Penia D				IPUX	VWA
		Item 2	- Amelogenin 8	& STR Results		
NWB3LP F	PowerPlex® 21					
2	•	19,19		15,16	11,13	10,11
	10,17		18,19	11,12	10,12	12,14
	14,14	30,31.2		X,X	11,12	23,24
	10,13	12,18		7,7	11,11	16,17
NZEZWG 16	dentifiler® Plus					
2		19,19		15,16	11,13	10,11
	10,17			11,12	10,12	12,14
	14,14	30,31.2		X,X	11,12	23,24
				7,7	11,11	16,17
UAG7DC	dentifiler® Plus					
2		19		15,16	11,13	10,11
	10,17			11,12	10,12	12,14
	14	30,31.2		X	11,12	23,24
				7	11	16,17
V89DWH	dentifiler® Plus					
2		19		15,16	11,13	10,11
2	10,17	17		11,12	10,12	12,14
	14	30,31.2		X,X	11,12	23,24
	17	30,31.2		7	11	16,17
VH6JPG F	PowerPlex® 21			,		10,17
		10.10		15.17	11.10	10.11
2	•	19,19	10.10	15,16	11,13	10,11
	10,17	20.21.0	18,19	11,12	10,12	12,14
	14,14	30,31.2		X,X	11,12	23,24
	10,13	12,18		7,7	11,11	16,17
	dentifiler® Plus and D					
2		19		15,16	11,13	10,11
	10,17			11,12	10,12	12,14
	14	30,31.2		Χ	11,12	23,24
				7	11	16,17
VWN6AA F	PowerPlex® PP16HS					
2				15,16	11,13	10,11
	10,17			11,12	10,12	12,14
		30,31.2		Χ	11,12	23,24
	10,13	12,18		7	11	16,17
XR8JD9 F	PowerPlex® HS					
2				15,16	11,13	10,11
	10,17			11,12	10,12	12,14
		30,31.2		X	11,12	23,24
	10,13	12,18		7	11	16,17
Z6822D la	dentifiler® Plus					
2		19,19		15,16	11,13	10,11
2	10,17	17,17		11,12	10,12	12,14
	14,14	30,31.2		X,X	11,12	23,24
	1 1/1 7	50,01.2		7,7	11,11	16,17
				1,1	11,11	10,17

DNA Parentage Table 1

WebCode Ite	m D1S1656 D8S1179 D19S433 Penta D	D2S1338 D10S1248 D21S11 Penta E	D2S441 D12S391 D22S1045 SE33	D3S1358 D13S317 Amelogenin TH01	D5S818 D16S539 CSF1PO TPOX	D7S820 D18S51 FGA vWA
		Item 2	- Amelogenin 8	& STR Results		
ZVFHN6 10	dentifiler®					
2		19		15,16	11,13	10,11
	10,17			11,12	10,12	12,14
	14	30,31.2		Χ	11,12	23,24
				7	11	16,17

TABLE 1

				TABLE			
WebCode Ii	lem	D1S1656 D8S1179 D19S433 Penta D	D2S1338 D10S1248 D21S11 Penta E	D25441 D125391 D2251045 SE33	D3S1358 D13S317 Amelogenin TH01	D5S818 D16S539 CSF1PO TPOX	D7S820 D18S51 FGA vWA
				- Amelogenin 8			
4TWZUA	Powe	rPlex® 21		7			
	3	10,16	19,23		15,15	11,11	11,13
	J	14,17	17,25	18,21	12,12	10,11	12,18
		14,17	30,30.2	10,21	X,Y	10,11	24,25
		12,13	12,12		6,7	11,11	17,18
71.6000		•	12,12		0,7	11,11	17,10
LCGP2		ifiler®					
	3		19,23		15	11	11,13
		14,17			12	10,11	12,18
		14	30,30.2		X,Y	10,11	24,25
					6,7	11	17,18
2EXNX	ldenti	ifiler® plus kit					
	3	<u></u>	19,23		15,15	11,11	11,13
	Γ	14,14			12,12	10,11	12,18
	ſ	14,17	30,30.2		X,Y	10,11	24,25
	L				6,7	11,11	17,18
7BL66	Pove	rPlex® 21			- / ·	.,	,
7 BLOO							
	3	10,16	19,23		15,15	11,11	11,13
		14,17	00.00	18,21	12,12	10,11	12,18
		14,14	30,30.2		X,Y	10,11	24,25
		12,13	12,12		6,7	11,11	17,18
293HX	ldenti	ifiler®					
	3		19,23		15	11	11,13
		14,17			12	10,11	12,18
		14	30,30.2		X,Y	10,11	24,25
					6,7	11	17,18
RL7GW	Powe	rPlex® PowerPlex	16 HS				
	3				15	11	11,13
		14,17			12	10,11	12,18
		•	30,30.2		X,Y	10,11	24,25
		12,13	12		6,7	11	1 <i>7,</i> 18
UUGAU	Identi	ifiler® Plus					*
200/10	3	5. 5 1 105	10.00		15 15	11 11	11 10
	S	14 17	19,23		15,15 12,12	11,11 10,11	11,13
		14,17 14,14	30,30.2		12,12 X,Y	10,11	12,18 24,25
		17,14	50,50.2		6,7	11,11	24,25 17,18
\		·(i) @ DI			0,/	11,11	17,10
WLP7Y		ifiler® Plus					
	3		19,23		15,15	11,11	11,13
		14,17	00.5		12,12	10,11	12,18
		14,14	30,30.2		X,Y	10,11	24,25
					6,7	11,11	17,18
A7CHK	NGN	1 SElect					
	3	10,16	19,23	10,11	15,15		
		14,17	13,16	18,21		10,11	12,18
		14,14	30,30.2	15,16	X,Y		24,25

TABLE 1

				TABLE	<u> </u>		
WebCode l	ltem	D1S1656 D8S1179	D2S1338 D10S1248	D2S441 D12S391	D3S1358 D13S317	D5\$818 D16\$539	D7S820 D18S51
		D19S433	D21S11	D22S1045	Amelogenin	CSF1PO	FGA
		Penta D	Penta E	SE33	TH01	TPOX	vWA
			Item 3	- Amelogenin 8	& STR Results		
NWB3LP	Powe	erPlex® 21					
	3	10,16	19,23		15,15	11,11	11,13
		14,17		18,21	12,12	10,11	12,18
		14,14	30,30.2		X,Y	10,11	24,25
		12,13	12,12		6,7	11,11	17,18
nzezwg	ldent	ifiler® Plus					
	3		19,23		15,15	11,11	11,13
		14,17	•		12,12	10,11	12,18
		14,14	30,30.2		X,Y	10,11	24,25
					6,7	11,11	17,18
UAG7DC	ldent	ifiler® Plus					
	3		19,23		15	11	11,13
	J	14,17	17,20		12	10,11	12,18
		14	30,30.2		X,Y	10,11	24,25
			/		6,7	11	17,18
V89DWH	Ident	ifiler® Plus					
VO/DVVII		illiel © 1 los	10.02		1.5	1.1	11 12
	3	14,17	19,23		15 12	11	11,13
		14,17	30,30.2		X,Y	10,11 10,11	12,18 24,25
		14	30,30.2		6,7	11	17,18
VH6JPG	Davis	erPlex® 21			0,7		17,10
VIIOJEG			10.00		15.15	11.11	11.10
	3	10,16	19,23	10.01	15,15	11,11	11,13
		14,17 14,14	20.20.2	18,21	12,12	10,11	12,18
		12,13	30,30.2 12,12		X,Y 6,7	10,11 11,11	24,25 17,18
1 A DVI 4 C					0,7	11,11	17,10
VNXJ4C		ifiler® Plus and D					
	3		19,23		15	11	11,13
		14,17	00.00		12	10,11	12,18
		14	30,30.2		X,Y	10,11	24,25
	_				6,7	11	17,18
VWN6AA	Powe	erPlex® PP16HS					
	3				15	11	11,13
		14,17			12	10,11	12,18
		-	30,30.2		X,Y	10,11	24,25
		12,13	12		6,7	11	17,18
XR8JD9	Powe	erPlex® HS					
	3				15	11	11,13
		14,17			12	10,11	12,18
			30,30.2		X,Y	10,11	24,25
		12,13	12		6,7	11	17,18
Z6822D	ldent	ifiler® Plus					
	3		19,23		15,15	11,11	11,13
		14,17	•		12,12	10,11	12,18
		14,14	30,30.2		X,Y	10,11	24,25
					6,7	11,11	17,18

DNA Parentage Table 1 Test 15-5870

WebCode Ite	m D1S1656 D8S1179 D19S433	D2S1338 D10S1248 D21S11	D2S441 D12S391 D22S1045	D3S1358 D13S317 Amelogenin	D5S818 D16S539 CSF1PO	D7\$820 D18\$51 FGA
	Penta D	Penta E	SE33 TH01	TPOX	vWA	
		Item 3	- Amelogenin 8	& STR Results		
ZVFHN6 10	dentifiler®					
3		19,23		15	11	11,13
	14,17			12	10,11	12,18
	14	30,30.2		X,Y	10,11	24,25
				6,7	11	17,18

Item 3 Paternity Index Results

				IADLL .				
WebCode	ltem	D1S1656 D8S1179 D19S433 Penta D	D2S1338 D10S1248 D21S11 Penta E	D25441 D125391 D2251045 SE33	D3S1358 D13S317 Amelogenin TH01	D5S818 D16S539 CSF1PO TPOX	D7S820 D18S51 FGA vWA	
			ltem (3 - Paternity Inc	dex Results			
4TWZUA	NIST	Γ-STRBASE						
	3PI	47.2	2.5934		1.6620	2.5654	1.8018	
	011	47.2	2.0704	2.8090	2.2056	3.3245	4.3706	
		2.8265	1.3408	2.0070	1	1.7876	3.5236	
		3.4698	5.7571		1.6858	3.9939	2.0342	
7LCGP2	FBI I	PopStats			.,,,,,,,			
	3PI	1 -	1.91894		1.44636	2.37428	1.72729	
	511	40.6	1.71074		2.38823	2.88884	4.72099	
		3.1209	1.20117		2.00020	1.88288	3.98026	
		J. 12U7	1.2011/		1.48227	3.66663	2.25561	
92EXNX	NIST	Γ-STRBASE				2.22300	2.20001	
•	3PI		2.8		1.73	2.86	1.94	
	511	70.03			2.19	4.24	4.24	
		2.67	1.35			1.71	3.33	
			1.55		1.79	3.64	2.30	
97BL66	NIST-STRBASE							
	3PI	15.055	2.206		1.549	2.488	1.822	
		16.006	_,	2.689	1.947	3.098	3.867	
		2.516	1.207			1.809	3.251	
		3.21	4.77		1.567	3.238	2.03	
A293HX	FBI I	PopStats						
	3PI		1.92		1.45	2.37	1.73	
		40.60			2.39	2.89	4.72	
		3.12	1.20			1.88	3.98	
					1.48	3.67	2.26	
ARL7GW	labo	oratory specific date	abase					
	3PI				1.4992	2.8735	1.9230	
		41.6666			2.2779	3.2894	4.6728	
			1.3623		N/A	1.6611	2.6315	
		2.7932	5.2910		1.7361	4.0816	1.8726	
BUUGAU	loca	l database						
	3PI		2.73		1.55	3.15	1.36	
		32.05			2.71	4.00	12.62	
		4.10	1.40		-	2.04	3.13	
					1.93	3.73	1.98	

TABLE 2

				IABLE	_		
WebCode	Item	D1S1656 D8S1179 D19S433	D2\$1338 D10\$1248 D21\$11	D2S441 D12S391 D22S1045	D3\$1358 D13\$317 Amelogenin	D5S818 D16S539 CSF1PO	D7S820 D18S51 FGA
		Penta D	Penta E	SE33	TH01	TPOX	vWA
			ltem :	3 - Paternity Inc	dex Results		
EWLP7Y	FBI F	PopStats					
	3PI		1.9189		1.4463	2.3742	1.7271
	0	40.650	,,		2.3878	2.8885	4.7214
		3.1209	1.2011		2.557.5	1.8825	3.9809
		0.1207	1.2011		1.4824	3.6670	2.2553
КА7СНК	FBI PopStats						
	3PI	47.170	2.5934	1.4841	1.662		
	011	47.170	1.8295	2.809	1.002	3.3245	4.3706
		2.8265	1.3408	0.96731		0.0240	3.5236
		2.0200	1.0400	47.170	1.6858		2.0342
NWB3LP	NIST	-STRBASE					
	3PI	15.0519	2.2063		1.5491	2.4880	1.8217
	•	15.9946		2.6892	1.9471	3.0986	3.8661
		2.5162	1.2065	2.55,2		1.8088	3.2510
		3.2101	4.7705		1.5674	3.2385	2.0304
NZEZWG	FBI PopStats						
	3PI		1.9189		1.4463	2.3742	1.7271
	0	40.650	,,		2.3878	2.8885	4.7214
		3.1209	1.2011		2.557.5	1.8825	3.9809
			.,		1.4824	3.6670	2.2553
JAG7DC	Labo	oratory specific da	abase				
	3PI		2.1968		1.5183	2.5549	1.7367
		58.0000			2.2747	3.1867	4.8355
		3.1525	1.3182			1.7680	3.2594
					1.4872	3.6710	2.3201
/89DWH	[Cou	untry] Caucasian F	op Database				
	3PI		4.81		1.90	2.75	2.40
		135.14			1.71	8.71	3.47
		2.87	1.44			1.63	3.60
					2.65	3.71	1.87
VH6JPG	NIST	-STRBASE					
	3PI	15.052	2.206		1.549	2.488	1.822
		15.995		2.690	1.947	3.099	3.866
		2.516	1.207			1.809	3.251
		3.210	4.771		1.567	3.239	2.030
VNXJ4C	NIST	-STRBASE					
	3PI		2.59		1.66	2.57	1.80
		118.00			2.21	3.32	4.37
		2.83	1.34			1.79	3.52
					1.69	3.93	2.03

			I ADLL	_		
WebCode Ite	D1\$1656 D8\$1179 D19\$433	D2S1338 D10S1248 D21S11	D2S441 D12S391 D22S1045	D3S1358 D13S317 Amelogenin	D5S818 D16S539 CSF1PO	D7\$820 D18\$51 FGA
	Penta D	Penta E	SE33	TH01	TPOX	vWA
		Item 3	3 - Paternity Inc	dex Results		
VWN6AA	Laboratory specific da	tabase				
3	PI			1.4992	2.8735	1.9230
	41.6666			2.2779	3.2894	4.6728
		1.3623			1.6611	2.6315
	2.7932	5.2910		1.7361	4.0816	1.8726
XR8JD9	Laboratory Specific Do	atabase				
3	PI			1.4992	2.8735	1.9230
	41.6666			2.2779	3.2894	4.6728
		1.3623			1.6611	2.6315
	2.7932	5.2910		1.7361	4.0816	1.8726
Z6822D	FBI PopStats					
3	PI	1.9189		1.4463	2.3742	1.7271
	40.650			2.3878	2.8885	4.7214
	3.1209	1.2011			1.8825	3.9809
				1.4824	3.6670	2.2553
ZVFHN6	FBI PopStats					
3	PI	1.9189		1.4463	2.3742	1.7271
	40.650			2.3878	2.8885	4.7214
	3.1209	1.2011			1.8825	3.9809
				1.4824	3.6670	2.2553

YSTR Results

TABLE 3

WebCode	Item	DYS19 DYS437	DYS385 DYS438	DYS389-I DYS439	DYS389-II DYS448	DYS390 DYS456	DYS391 DYS458	DYS392 DYS481	DYS393 DYS533
		DYS549	DYS570	DYS576	DYS635	DYS643	Y GATA H4	YIndel	
				Item 3 - Y	STR Results				
EWLP7Y	Yfiler®								
	3	14	11,14	13	29	24	10	13	13
		14	12	13	18	18	18		
					23		11		
NZEZWG	Yfiler®								
	3	14	11,14	13	29	24	10	13	13
		14	12	13	18	18	18		
					23		11		
Z6822D	Yfiler®								
	3	14	11,14	13	29	24	10	13	13
		14	12	13	18	18	18		
					23		11		

Additional DNA & PI Results

Locus	WebCode	Item 1	Item 2	Item 3	Item 3 Paternity Index
D6S1043	4TWZUA	14,14	12,14	12,13	2.4331
	97BL66	14,14	12,14	12,13	2.376
	NWB3LP	14,14	12,14	12,13	2.3764
	VH6JPG	14,14	12,14	12,13	2.376

Paternity DNA Statistics

TABLE 5

WebCode	Combined Paternity Index	Probability of Paternity	Population Database
4TWZUA	62 billion	99.999% (Not routinely reported)	NIST-STRBASE
7LCGP2	5200000	99.99998	FBI PopStats
92EXNX	22792890.00	99.99999 %	NIST-STRBASE
97BL66	1.57 x 10 ⁹	NA	NIST-STRBASE
A293HX	5200000	99.99998%	FBI PopStats
ARL7GW	14.1 million	99.9%	laboratory specific database
BUUGAU	42,595,735	99.9999%	local database
EWLP7Y	5.188 MILLION	99.99998072%	FBI PopStats
KA7CHK	2,221,000,000	99.9999995498	FBI PopStats
NWB3LP	1.57 x 10 ⁹	Not performed by the laboratory	NIST-STRBASE
NZEZWG	5,188,000	99.99998072	FBI PopStats
UAG7DC	9.12 million	99.9%	Laboratory specific database
V89DWH	1.88 x 10^8 (188,924,012)	greater than 99.99% (99.9999947%)	[Country] Caucasian Pop Database
VH6JPG	1.57 x 10 ⁹	Not determined	NIST-STRBASE
VNXJ4C	24,246,002	99.9999%	NIST-STRBASE
VWN6AA	14.1 million	99.9%	Laboratory specific database
XR8JD9	14.1 million	99.9%	Laboratory Specific Database
Z6822D	5,188,000	99.99998072	FBI PopStats
ZVFHN6	5,188,000	>99.999%	FBI PopStats

Paternity Conclusions

ReportCode	Conclusions	
4TWZUA	Not Excluded	
7LCGP2	Not Excluded	
92EXNX	Not Excluded	
97BL66	Not Excluded	
A293HX	Not Excluded	
ARL7GW	Not Excluded	
BUUGAU	Not Excluded	
EWLP7Y	Not Excluded	
KA7CHK	Not Excluded	
NWB3LP	Not Excluded	
nzezwg	Not Excluded	
UAG7DC	Not Excluded	
V89DWH	Not Excluded	
VH6JPG	Not Excluded	
VNXJ4C	Not Excluded	
VWN6AA	Not Excluded	
XR8JD9	Not Excluded	
Z6822D	Not Excluded	
ZVFHN6	Not Excluded	

Response Summary						
es es	Not Excluded	19				
sponse	Excluded	0				
Resp	Inconclusive	0				

Kinship DNA Statistics

Is the claim of a full sibling relationship supported by the genetic evidence?

WebCode	Population Database	Kinship Index	Claim Supported?
4TWZUA	NIST-STRBASE	25 Thousand	Yes
7LCGP2	FBI PopStats	1700	Yes
97BL66	NIST-STRBASE	75,000	Yes
A293HX	FBI PopStats	1700	Yes
EWLP7Y	FBI PopStats	1,691	Yes
KA7CHK	FBI PopStats	950,300	Yes
NWB3LP	NIST-STRBASE	75,000	Yes
NZEZWG	FBI PopStats	1,691	Yes
VH6JPG	NIST-STRBASE	75,000	Yes
Z6822D	FBI PopStats	1,691	Yes
ZVFHN6	FBI PopStats	1,691	Yes

Additional Kinship Statistical Results

WebCode	Additional Statistical Results
7LCGP2	Our inhouse kinship worksheet only provides stats on identifiler loci
97BL66	The kinship index was determined through using the caucasian population database with an Fst value of 0.02.
A293HX	Calculated kinship index using only Identifiler loci.
КА7СНК	DNA profiles A and B are approximately nine-hundred and fifty thousand times more likely to be observed if they came from individuals who are full siblings rather than from individuals who are unrelated to each other. It has been assumed that the parents of individuals A and B are unrelated to each other and that a double first cousin relationship between individuals A and B is not a possibility. The next most closely supported possible retaionship[sic] is a Half-Sib, Uncle/Aunt- Nephew/Niece, Grandparent-Grandchild relationship. Nevertheless, the profiles of individuals A and B are approximately one hundred and thirty times more likely to be observed if they came from full siblings rather than from, for example, half siblings.
NWB3LP	A theta value 0.02 was used in this calculation.
VH6JPG	Kinship index has been determined using a theta value of 0.02.
ZVFHN6	The statistical results are based on the STR loci of the Identifiler PCR amplification kit: D8S1179, D21S11, D7S820, CSF1PO, D3S1358, TH01, D13S317, D16S539, D2S1338, D19S433, vWA, TPOX, D18S51, D5S818, and FGA using FBI Caucasian population database.

Additional Comments

	IT IDEE 7
WebCode	Additional Comments
4TWZUA	Our statistical software uses a minimum allele frequency of 5 observations in the data set. Where alleles are rarer than this a frequency equivalent to 5 observations in the data set is used. This has occurred at D1 and D8 Part 1 [Table 2 - Item 3 Paternity Index Result]. Our PIs at these loci may differ from others as a result.
97BL66	The alleged father and mother were assumed to be from the same sub-population (hispanic) and an Fst value of 0.03 used in the calculation of the Paternity Index. The Probability of Paternity is not a calculation performed within this laboratory.
KA7CHK	Additional comments on Part II [Tables 5 & 6; Paternity DNA Statistics & Paternity Conclusions] of the test: The DNA profiles obtained from items 1, 2 and 3 are in excess of one thousand million times more likely to be observed if the alleged father and the mother are the parents of the child rather than if a man unrelated to the alleged father and the mother are the parents.
NWB3LP	For the Paternity Index calculated in Part 2 [Table 2] the following assumptions were made: The mother, child and alleged father were from the same sub-population (Hispanic). A Theta value of 0.03 was used.
V89DWH	Population database used was published in. (2007) Eckhoff, C. et al. "Population data from sub-populations of the [Country] for 15 autosomal short tandem repeats (STR) loci" Forensic Sci. Int 171:237-49
VH6JPG	Paternity Index calculated using a theta value of 0.03, and assuming Mother and Alleged Father are from the same Hispanic subpopulation. Note: Probability of Paternity is not calculated in our laboratory, so this has not been determined for this test.
ZVFHN6	The trio paternity comparison used the FBI Southwestern Hispanic population database. The sibling comparison used only those markers in the Identifiler amplification set (D8S1179, D21S11, D7S820, CSF1PO, D3S1358, TH01, D13S317. D16S539, D2S1338, D19S433, vWA, TPOX, D18S51, D5S818, and FGA) and results are based on FBI Caucasian population database.

Appendix: Data Sheet

Collaborative Testing Services ~ Forensic Testing Program

Test No. 15-5870: DNA Parentage

DATA MUST BE RECEIVED BY <u>March 23, 2015</u> TO BE INCLUDED IN THE REPORT Participant Code: WebCode:

Accreditation Release	Statement
CTS submits external proficiency test data directly to A Please select one of the following statements to ensur	
This participant's data is intended for submission to A (Accreditation Release section on the last page must be c	SCLD/LAB and/or ANSI-ASQ NAB/FQS. ompleted and submitted.)
This participant's data is NOT intended for submissio	n to ASCLD/LAB or ANSI-ASQ NAB/FQS.

Scenario:

A standard paternity trio case has been presented to your laboratory. Blood standards have been collected from the mother, daughter and alleged father. Your laboratory is tasked with examining the blood standards and comparing the DNA profiles.

Items Submitted (Sample Pack DNP1):

Item 1: Blood Sample from Known Parent (Mother)

Item 2: Blood Sample from Known Child (Daughter)

Item 3: Blood Sample from Alleged Father (Hispanic)

DNA Reporting Instructions:

Use the instructions below to complete the following DNA Analysis sections of this data sheet.

- * Report alleles in numerical order, separated by a comma.
- * Follow your laboratory procedures for reporting homozygotes (i.e. "14,14", "14,-", "14")
- * PI = Paternity Index; KI Kinship Index

Example	D1S1656	D2S1338	D2S441	D3S1358	D5S818	D7S820
STR	15,18	12,17	10	14	12	5,13
PI	1.65	3.01	3.16	4.12	2.45	5.65

Online Data Entry

Visit <u>www.cts-portal.com</u> to enter and/or upload your proficiency test results online. If you have any questions please do not hesitate to contact CTS.

Please return all pages of this data sheet.

Page 1 of 8

Participant Code: WebCode:

Part I: DNA ANALYSIS FOR ITEM 1

STR	Amp	olification Ki		he brands that apply f d (i.e. 16, Plus, Direct,	or this item and record of 16 HS, etc.).	any additional kit speci	fic naming in the
			us®			x®	
Щ	Iden						
		D1S1656	D2S1338	D2S441	D3S1358	D5S818	D7S820
S	TR						
	·						
		D8S1179	D10S1248	D12S391	D13S317	D16S539	D18S51
= 2	TR						
ITEM 1		D19S433	D21S11	D22S1045	Amelogenin	CSF1P0	FGA
	i	175400	DZ1311	DZZ51043	Amologomii		10/
S	TR						
	i			0500		TDOV	
		Penta D	Penta E	SE33	TH01	TPOX	vWA
S	TR						
<u>ADDI</u>	TIOI	NAL DNA RE	ESULTS FOR ITE	<u>M 1</u>			
(If add	itional	l space is needed	l, copy this page or atta	ch your own form	following this layout)		
ITEM 1 Alleles Alleles							
	7 HIGHS						
				<u> </u>			

Participant Code: WebCode:

Part I: DNA ANALYSIS FOR ITEM 2

	plification Kit	blank provide	the brands that apply for d (i.e. 16, Plus, Direct,		,	•
	filer®/Profiler Plus ntifiler®	s®			x®	
	D1S1656	D2S1338	D2S441	D3S1358	D5S818	D7S820
STR						
	D8S1179	D10S1248	D12S391	D13S317	D16S539	D18S51
ATS STR						
ΞE	D19S433	D21S11	D22S1045	Amelogenin	CSF1P0	FGA
STR						
	Penta D	Penta E	SE33	TH01	TPOX	vWA
STR						
		SULTS FOR ITE				
(If addition	al space is needed,	copy this page or atta ITEM 2	•	following this layout)	ITE	M 2
		Alleles			Alle	les

Page 4 of 8

Participant Code: WebCode:

Part I: DNA ANALYSIS FOR ITEM 3

<u>S</u> 1	STR Amplification Kit Used: Please check the brands that apply for this item and record any additional kit specific naming in the blank provided (i.e. 16, Plus, Direct, 16 HS, etc.).									
	Cofiler®/Profiler Plus® Identifiler®					Po				
					NA Statistics' s e to determin		is data s	heet	t regardin	g the
		DISI	656	D2S1338	D2S441	D3S135	8	D5S	818	D7S820
	STR PI						╛╘			
		D8S1	179	D10S1248	D12S391	D13S31	7	D169	5539	D18S51
× 3	STR PI									
ITEM		D19S	433	D21S11	D22S1045	Ameloger	nin	CSF	1PO	FGA
	STR PI		<u> </u>							
	•	Pent	ra D	Penta E	SE33	TH01		TP	L 0X	vWA
	STR									
	PI									
					proficiency cor				1.0	· · · · · · · · · · · · · · · · · · ·
<u>Y:</u>		iler®	<u>ition Kit Use</u>	in the blank	k all the brands the provided (i.e. Pluc Plex® Y	ıs, 23, etc.).	s item and r		any kif speci	fic information
		DYS19	DYS385	DYS389-I	DYS389-II	DYS390	 DYS391		DYS392	DYS393
۷ ع	D	YS437	DYS438	DYS439	DYS448	DYS456	DYS458	}	DYS481	DYS533
ITEM										
	D	YS549	DYS570	DYS576	DYS635	DYS643	Y GATA H	4	Y Indel	7
				TS FOR ITE		m following this	e lavout)			
(If additional space is needed, copy this page or attach your own form following this layout) ITEM 3 Alleles Paternity Index Alleles Paternity						FEM 3 Paternity Index				
				_						
				_	· .				. ———	

Please return all pages of this data sheet.

Test 15-5870

Part II: PATERNITY DNA STATISTICS

For the purposes of consistency among reported statistical values, use the ethnicity listed for the alleged parent and choose one of the following population databases for all statistical calculations in this test:

- 1. **FBI Popstats:** If FBI Popstats is already available in your laboratory then you may select that option, otherwise use the population database below.
- 2. **NIST-STRBASE** is a publicly available U.S. population dataset at STRBASE on the following NIST web site: http://www.cstl.nist.gov/strbase/NISTpop.htm#Autosomal
- a. On the NIST web site, select the hyperlink labeled "Allele frequencies from autosomal STRs as Excel file" under the title "NIST 1036 U.S. Population Dataset".
- 3. If you unable to use one of the suggested population databases, report the population database used in the blank provided next to the "Other Pop. Database" option. Due to the tendency for allele frequencies to vary amongst different databases, no consensus value will be determined for this option. When reporting a population database name, please refrain from using terms that would allude to a laboratory specific name or location; general terms such as "local/state database" or "laboratory specific database" are preferred.

1) Ch	oose a Population Da	tabase:				
	FBI Popstats Pop	Database		NIST STRBASE	Pop. Database	
	Other Pop. Datal	pase:			_	
2) Red	cord the Combined Pa	ternity Index value:				_
3) Red	cord the Probability of	Paternity:				_
wordi		elect your response from the fa apt these conclusions as best				
	The Alleged pare	ent (Item 3) could not be exclu	ded as the b	piological parent of	the child (Item 2).	
	The Alleged pare	ent (Item 3) is excluded as a po	ossible biolo	gical parent of the	child (Item 2).	
		o whether the Alleged parent (It the reason in the Additional	•	•	. , ,	

Participant Code: WebCode:

Part III: KINSHIP DNA STATISTICS (NON-PARENTAGE)

To be completed if applicable to your laboratory.

The two DNA profiles below are presented as potential **Caucasian full siblings**. Compare these profiles to answer the questions and use the same population database used in previous sections of the data sheet, following the ethnicity listed above for this kinship scenario.

	D1S1656	D2S1338	D2S441	D3\$1358	D5\$818	D7\$820
Profile A	14,15	16,20	10,11.3	15,17	12,13	10,12
Profile B	13,15	16,18	14,14	15,17	12,13	10,11

	D8S1179	D10S1248	D12S391	D13S317	D16\$539	D18S51
Profile A	12,13	12,13	17,20	11,13	11,12	12,18
Profile B	12,13	13,14	17,20	11,13	11,12	15,16

	D19\$433	D21511	D22\$1045	Amelogenin	CSF1PO	FGA
Profile A	15,16.2	28,29	11,15	X,X	10,11	20,23
Profile B	13,15	29,31	11,11	X,Y	11,13	19,21

	Penta D	Penta E	SE33	TH01	трох	vWA
Profile A	9,10	7,7	13,18	6,9	8,8	17,19
Profile B	10,11	7,12	13,18	9,9.3	8,8	17,19

1) Evaluate profiles A and B and record the kinship index. 2) Is the claim of a full sibling relationship supported by the genetic evidence?					

Participant Code: WebCode:

Part IV: ADDITIONAL COMMENTS						
Comments regarding any part of this Parentage Test.						
Any interpretations based on the results obtained should be reported in the Paternity DNA Statistics designated section.						

Return Instructions: Data must be received via online data entry, fax (please include a cover sheet), or mail by March 23, 2015 to be included in the report.

ONLINE DATA ENTRY: www.cts-portal.com

+1-571-434-1937 FAX:

or Toll-Free: 1-866-FAX-2CTS (329-2287)

MAIL: Collaborative Testing Services, Inc.

P.O. Box 650820

EMAIL: forensics@cts-interlab.com

+1-571-434-1925 (8 am - 4:30 pm EST)

QUESTIONS?

www.ctsforensics.com Sterling, VA 20165-0820 USA

Collaborative Testing Services ~ Forensic Testing Program

RELEASE OF DATA TO ACCREDITATION BODIES

The following Accreditation Releases will apply only to:

Participant Code: WebCode:

for Test No. 15-5870: DNA Parentage

This release page must be completed and received by <u>March 23, 2015</u> to have this participant's submitted data included in the reports forwarded to the respective Accreditation Bodies.

ASCLI	D/LAB RELEASE
If your lab has been accredited by ASCLD/LAB ar proficiency test requirements, have the laboratory	nd you are submitting this data as part of their external
	s designated marviabal complete the following. Intirety for the results to be submitted to ASCLD/LAB.
ASCLD/LAB Legacy Certificate AS	CLD/LAB International Certificate No
Signature	Date
Laboratory Name	
Location (City/State)	
ANSI-ASQ	NAB/FQS RELEASE
If your laboratory maintains its accreditation thro form in its entirety to have your results forwarded	ugh ANSI-ASQ NAB/FQS, please complete the following
ANSI-ASQ NAB/FQS Certificate	
Signature and Title	Date
Laboratory Name	
Location (City/State)	

Accreditation Release

Return Instructions

Please submit the completed Accreditation Release at the same time as your full data sheet. See Data Sheet Return Instructions on the previous page.

Questions? Contact us 8 am-4:30 pm EST Telephone: +1-571-434-1925 email: forensics@cts-interlab.com

Please return all pages of this data sheet.

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