

Probabilistic Genotyping Test No. 25-5901/2 Summary Report

Each participant received a sample pack consisting of two known bloodstains and two questioned stains which they were asked to analyze using their existing protocols. Data were returned from 80 participants and are compiled into the following tables:

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Appendix: Data Sheet

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.

Participant results are reported using a randomly assigned "WebCode". This code maintains participant's anonymity, provides linking of the various report sections, and will change with every report.

Manufacturer's Information

Each sample pack consisted of two known bloodstains provided on either white fabric or FTA[™] Micro Cards, and two questioned stains on colored fabric. Participants were asked to analyze these items using their existing protocols.

SAMPLE PREPARATION: Stains from different sources were prepared at separate times and were packaged once they were thoroughly dried into separate envelopes.

SAMPLE PACK ASSEMBLY: One of each item was placed into a pre-labeled sample pack envelope and sealed. The sealed envelopes were then packaged in pre-labeled heat seal envelopes and sealed. Completed sample packs were stored at -20°C until shipment.

VERIFICATION: Predistribution results were consistent with each other and the manufacturer's preparation information. Consistent allelic results were reported for all STR and YSTR loci.

ltem	Known/ Questioned	Substrate (Test No.)	Body Fluid	Volume (Mixture Ratio)	Sex	Donor Information
1	Known	White Fabric (5901)/ FTA™ Micro Card (5902)	Blood	50 μL/ 75 μL	Female	Victim
2	Known	White Fabric (5901)/ FTA™ Micro Card (5902)	Blood	50 μL/ 75 μL	Male	Suspect
3	Questioned	Tan Fabric	Blood/Blood/Semen	50 µL (1:1:1)	Female/Male/Male	Victim/Suspect/ Additional Male A
4	Questioned	Blue Fabric	Blood/Blood/Blood	50 μL (1:1:1)	Male/Male/Male	Suspect/Additional Male B/Additional Male C

Amelogenin and STR Results							
	Results co	mpiled from predi	stribution laborato	ries and a consensus o	of at least 10 pai	ticipants.	
ltem	D1\$1656	D2S1338	D2S441	D3S1358	D5\$818	D6S1043	
	D75820	D8\$1179	D10S1248	D12S391	D13S317	D16S539	
	D18\$51	D195433	D21511	D22S1045	Amelogenin	CSF1PO	
	FGA	Penta D	Penta E	SE33	TH01	TPOX	
	vWA	DYS391	DY\$570	DYS576	Y Indel		
	, /.3	17,25	,	16,18	10,12	,	
	10,12	13,15	13,13	16,17	8,12	9,12	
_	14,17	12,13	31.2,32.2	11,12	X,X	10,11	
	21,24	9,11	8,11	17,24.2	6,9.3	8,8	
	16,16	NM	NM	NM	<u>NM</u>		
2	16,17.3	17,18	14,14	16,16	11,13	12,13	
	9,11	10,13	13,13	18,19	12,12	9,14	
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12	
	20,23	9,12	5,10	18,28.2	7,7	11,14	
	15,17	10	*	*	2		
3-Blood	11,16,17.3	17,18,25	11,14	16,18	10,11,12,13	*	
	9,10,11,12	10,13,15	13	16,17,18,19	8,12	9,12,14	
	14,15,17,19	12,13,15.2	31.2,32.2,33,33.2	11,12,14,15	X,Y	10,11,12	
	20,21,23,24	9,11,12	5,8,10,11	17,18,24.2,28.2	6,7,9.3	8,11,14	
	15,16,17	10	*	*	2		
3-Semen	14,18.3	20,25	11,14	15,18	9,11	12,18	
	9,11	13,15	13,14	19,21	11,12	8,13	
	17,18	12,13	30,31.2	15,15	X,Y	10,11	
	20,23.2	10,13	8,12	15,28.2	6,9	8,10	
	16,17	10	*	*	2		
4	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	10,11,12,13,14	
	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14	
	13,15,17,19	12,13,13.2,14, 15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13	
	20,21,23,26	9,10,11,12,15	5,10,11,12,14	14,18,23.2,26.2,28.2	7,9.3	8,11,14	
	15,16,17	10,11	*	*	2		

Manufacturer's Information, continued

* Results were not received by a minimum of 10 participants for the loci indicated.

NM - Non-Male profile, YSTR results not expected.

YSTR Results									
	Results compiled from a consensus of at least 10 participants.								
ltem	DYF387S1	DYS19	DYS385	DYS389-I	DYS389-II	DYS390	DYS391	DYS392	DYS393
	DYS437	DYS438	DYS439	DYS448	DYS449	DYS456	DYS458	DYS460	DYS481
	DY\$518	DYS533	DYS549	DYS570	DYS576	DYS627	DYS635	DYS643	YGATAH4
2	35,37	13	16,17	13	30	24	10	11	13
	14	10	11	19	32	17	15	9	22
	40	12	*	20	17	22	21	*	11
3-Blood	35,37	13	16,17	13	30	24	10	11	13
	14	10	11	19	32	17	15	9	22
	40	12	*	20	17	22	21	*	11
3-Semen	35,36	14	11,13	14	30	24	10	13	13
	15	12	12	19	29	14	17	11	22
	38	12	*	18	19	20	24	*	12
4	35,36,37,38	13,14,15	11,15,16,17	12,13	28,29,30	22,24	10,11	10,11,14	13,14
	14,15,16	10,12	11,12	19,21	28,30,32	14,16,17	15,18,19	9,10,11	20,22,23
	37,38,40	10,12	*	17,20	16,17,21	20,22	21,23	*	11,13

* Results were not received by a minimum of 10 participants for the loci indicated.

Summary Comments

This test was designed to allow participants to assess their proficiency in the identification and comparison of dried stains by means of body fluid screening and/or DNA profiling utilizing Probabilistic Genotyping software. Participants were supplied with two "known" bloodstains (Items 1 and 2) and two "questioned" stains (Items 3 and 4). Item 1 was created using blood from a female donor. Item 2 was created using blood from a male donor. Item 3 was created by combining one part blood from the Item 1 female donor, one part blood from the Item 2 male donor, and one part semen from another male donor (Additional Male A) whose known standard was not provided. Item 4 was created by combining equal parts of blood from the Item 2 male donor, and two additional male donors (Additional Male B and Additional Male C), whose known standards were not provided. Refer to the Manufacturer's Information for preparation details.

Data were returned by 80 participants.

Screening Test Results

A total of 74 participants reported screening results for at least one body fluid (blood, semen, saliva), though not all of these participants returned screening results for every fluid across both questioned items.

For Item 3, all participants reported "Positive" for the presence of blood. For the presence of semen, all participants reported "Positive." For the presence of saliva, all but one participant reported "Negative." The remaining participant reported "Positive." For Human Origin Screening and Y-Screening, all participants reported "Positive."

For Item 4, all participants reported "Positive" for the presence of blood. For the presence of semen, all participants reported "Negative." For the presence of saliva, all participants reported "Negative." For Human Origin Screening, all but three participants reported "Positive." The remaining participants reported "Negative." For Y-Screening, all participants reported "Positive."

DNA Analysis

All participants reported DNA results. Only allelic results not containing the minimum expected alleles are highlighted as inconsistent.

For STR results, all but twenty-four participants reported consistent results for one or more items. The majority of participants were missing alleles for Item 3-Blood (3e), with over half missing one or more alleles at Locus D5S818. The majority of these missing alleles were attributable to the Item 1 female donor.

For YSTR results, all but six participants reported consistent results for one or more items.

DNA Interpretations

For Item 3, all but one participant included the victim (Item 1) as a possible contributor to the stain. The remaining participant reported "Inconclusive." All but one participant included the suspect (Item 2) as a possible contributor to the stain. The remaining participant reported "Inconclusive." All participants identified the presence of three contributors in the epithelial fraction. For the sperm fraction, a consensus was not achieved. A total of 44 participants (68%) identified one contributor, 15 participants (23%) identified two contributors, and six participants (9%) identified three contributors. An additional participant did not perform a differential extraction on this item, but reported the presence of three contributors.

For Item 4, all participants excluded the victim (Item 1) as a possible contributor to the stain. All but one participant included the suspect (Item 2) as a possible contributor. The remaining participant reported "Inconclusive" as to whether the suspect was a contributor. All but one participant identified the presence of three contributors. The remaining participant reported the presence of four contributors.

Key for Screening Tests Used

Participants were asked to use, where possible, the following chart of abbreviated screening test names. This was not an all inclusive list and was not designed to determine what tests should be performed. Participants were advised that tests not on this list may be used for screening.

Test	Abbreviation
Acid Phosphatase	AP
Kastle Meyer	КМ
Leucomalachite Green	lmg
Microscopic	Micro
Ortho-tolidine	O-tol
Phenolphthalein-Tetramethyl benzidine	PTMB
Prostate Specific Antigen	PSA
Quantiblot	QB
Quantifiler	QF
Tetramethyl benzidine	ТМВ

Serology Screening Results

Indicate the results of any screening tests performed on the questioned stains (Items 3 & 4).

TABLE 1a						
Blood Screening Results						
WebCode - Test	Item 3	Poc PTMP				
2QGMC2 - 5902						
34G3B3 - 5902	Pos KM					
37J2CU - 5901	Pos KM	Pos KM				
3MQZUY - 5902	Pos KM	Pos KM				
3WHFJ3 - 5902	Pos KM	Pos KM				
44PDRD - 5901	Pos KM	Pos KM				
4BEUJW - 5902	Pos TMB, Hematrace	Pos TMB, Hematrace				
4FBJY4 - 5902	Pos LMG	Pos LMG				
4UQG64 - 5901	Pos KM, Hematrace	Pos KM, Hematrace				
4XQRZ4 - 5901	Pos KM, Hematrace	Pos KM, Hematrace				
62K9Q4 - 5901	Pos KM, Hematrace	Pos KM, Hematrace				
63VBC2 - 5901	Pos KM, Hematrace	Pos KM, Hematrace				
6PTH6Y - 5902	Pos PTMB	Pos PTMB				
77A6RY - 5901	Pos IR, pth, Hematrace	Pos pth				
786NUZ - 5901	Pos KM, HemeDirect	Pos KM, HemeDirect				
7H4JRV - 5902	Pos PTMB	Pos PTMB				
8Y8DYV - 5902	Pos ptmb	Pos PTMB				
8YUL9Z - 5901	Pos TMB, FOB, Seratec PMB, Lumin	nol Pos TMB, FOB, Seratec PMB, Luminol				
9CLYK9 - 5901	Pos KM	Pos KM				
9MBR8Q - 5901	Pos TMB	Pos TMB				
9WLTTR - 5902	Pos KM, Hematrace	Pos KM, Hematrace				
B7W38V - 5901	Pos PTH, HEMATRACE, IR	Pos PTH				
BDR7Z7 - 5901	Pos KM, HemDirect	Pos KM, HemDIrect				
BKHHBU - 5901	Pos PTH, IR	Pos PTH, HemaTrace				
BNGT8U - 5902	Pos KM	Pos KM				
CDTG9M - 5901	Pos Hemastix, RSID	Pos Hemastix, RSID				
CKECAQ - 5902	Pos PTMB	Pos PTMB				
CRBAWU - 5901	Pos KM	Pos KM				
CRVQCW - 5901	Pos KM, Hematrace	Pos KM, Hemtrace				
CXYRCQ - 5902	Pos PTMB	Pos PTMB				
CY6DP6 - 5901	Pos KM	Pos KM				
CZ6EPV - 5901	Pos KM, HemaTrace	Pos KM, HemaTrace				

TABLE 1a

Blood Screening Results						
WebCode - Test Item 3 Item 4						
E34ANN - 5902	Pos PTMB	Pos PTMB				
EHXZJK - 5902	Pos TMB, Hematrace	Pos TMB, Hematrace				
ELJJPP - 5902	Pos PTMB	Pos PTMB				
EY6YTN - 5902	Pos PTMB	Pos PTMB				
F7H8UP - 5902	Pos KM	Pos KM				
F8XQDM - 5902	Pos TMB,KM	Pos TMB,KM				
FPXE6L - 5902	Pos KM	Pos KM				
GCH7HK - 5902	Pos PTMB	Pos PTMB				
H484WJ - 5902	Pos PTMB	Pos PTMB				
HEQHYL - 5902	Pos PTMB	Pos PTMB				
HJX7CN - 5901	Pos KM	Pos KM				
J37QNH - 5902	Pos Hemastix, Hematrace	Pos Hemastix, Hematrace				
J772JG - 5902	Pos TMB (Hemastix), Hematrace	Pos TMB (Hemastix), Hematrace				
JHLQHM - 5902	Pos Hemastix, OBTI	Pos Hemastix, OBTI				
JPGUBX - 5901	Pos KM	Pos KM				
K9ECCE - 5902	Pos KM, Hematrace	Pos KM, Hematrace				
KPDQAK - 5902	Pos KM	Pos KM				
L67JVL - 5901	Pos KM, Hematrace	Pos KM, Hematrace				
L8XEJE - 5901	Pos TMB	Pos TMB				
LBAKVF - 5902	Pos KM	Pos KM; ABAcard				
LG3PNJ - 5901	Pos LMG	Pos LMG				
LHUN3H - 5902	Pos TMB, PHE	Pos TMB, PHE				
LUPUWF - 5901	Pos KM	Pos KM				
LYKQJL - 5901	Pos KM, HemaTrace	Pos KM, HemaTrace				
MW99ZF - 5902	Pos TMB,PHE	Pos TMB,PHE				
NWNRKD - 5901	Pos KM	Pos KM				
P3QH3A - 5901	Pos TMB	Pos TMB				
P8KEPF - 5902	Pos KM	Pos KM				
Q99BQE - 5901	Pos Seratec HEM	Pos Seratec HEM				
R3MFPB - 5901	Pos KM	Pos KM				
R9RDYE - 5901	Pos KM, Hematrace	Pos KM, Hematrace				
RCPXZD - 5901	Pos KM	Pos KM				
TMDEV4 - 5902	Pos KM	Pos KM				
TZBZVA - 5901	Pos KM	Pos KM				

TABLE 1a

Blood Screening Results					
WebCode - Test	Item 3	Item 4			
UNVR99 - 5902	Pos PTMB	Pos PTMB			
VFHZHA - 5902	Pos KM	Pos KM			
VK9834 - 5902	Pos RSID Blood	Pos RSID Blood			
X28843 - 5901	Pos TMB	Pos TMB			
XCU6U9 - 5902	Pos LMG	Pos LMG			
YGJ6L9 - 5901	Pos KM, Hematrace	Pos KM, Hematrace			
YJMUQ3 - 5901	Pos O-tol	Pos O-tol			
Z8LUQ4 - 5902	Pos PTMB	Pos PTMB			

Table 1a: Serology Screening Response Summary - Blood

Participants: 74

This summary table excludes the count of participants who did not report or reported "Not Tested" for Item 3 and/or Item 4. Therefore, participant total may not align with totals shown below.

	Item 3	Item 4
Positive	74	74
Negative	0	0
Inconclusive	0	0

Serology Screening Results

Indicate the results of any screening tests performed on the questioned stains (Items 3 & 4).

TABLE 1b					
Wah Casla Tast	Seme	n Screening Results			
20GMC2 - 5902	Pos ALS AP Micro				
34G3B3 - 5902	Pos ALS AP p30 Micro	Neg ALS, AP, p30, Micro			
37/2011 - 5901	Pos AP Micro	Neg AP			
3MQ7UY 5902		Neg AP			
	D ₁ = AD A(:				
3WHFJ3 - 5902	POS AP, Micro	INEG AP			
44PDRD - 5901	Pos ALS, AP, Micro	Neg ALS, AP			
4BEUJW - 5902	Pos AP, PSA, Micro	Neg AP, PSA			
4FBJY4 - 5902	Pos AP, Micro	Neg AP, micro			
4UQG64 - 5901	Pos AP, PSA, Micro	Neg AP			
4XQRZ4 - 5901	Pos AP, PSA, Micro	Neg AP			
62K9Q4 - 5901	Pos AP, Micro, PSA	Neg AP			
63VBC2 - 5901	Pos AP, PSA, Micro	Neg AP			
6PTH6Y - 5902	Pos ALS, AP, Micro	Neg ALS, AP			
77A6RY - 5901	Pos ALS, AP, Micro, PSA	Neg AP, Micro			
786NUZ - 5901	Pos ALS, AP, PSA, Micro	Neg ALS, AP, PSA, Micro			
7H4JRV - 5902	Pos ALS, AP, Micro	Neg AP			
8Y8DYV - 5902	Pos AP, Micro, ALS	Neg AP, Micro, ALS			
8YUL9Z - 5901	Pos ALS, AP, RSID Semen, Se	ratec PAM Neg ALS, AP, RSID Semen, Seratec PAM			
9CLYK9 - 5901	Pos ALS, AP	Neg ALS, AP			
9MBR8Q - 5901	Pos AP, Micro	Neg AP			
9WLTTR - 5902	Pos Alternative Light Source, Microscopy	AP, Semenogelin, Neg Alternative Light Source, AP			
B7W38V - 5901	Pos ALS, AP, MICRO, PSA	Neg AP			
BDR7Z7 - 5901	Pos AP, Micro, PSA	Neg AP			
BKHHBU - 5901	Pos AP, PSA, micro, ALS	Neg AP, micro			
BNGT8U - 5902	Pos AP, p30, Micro	Neg AP, p30, Micro			
CDTG9M - 5901	Pos AP, RSID, Micro	Neg AP, RSID			
CKECAQ - 5902	Pos ALS, AP, MICRO				
CRBAWU - 5901	Pos AP	Neg AP			
CRVQCW - 5901	Pos AP, PSA, micro	Neg AP			
CXYRCQ - 5902	Pos ALS, AP, Micro	Neg ALS, AP			

TABLE 1b

		Semen Screening Results		
WebCode - Test	Ite	m 3	lte	m 4
CY6DP6 - 5901	Pos	ALS, AP	Neg	ALS, AP
CZ6EPV - 5901	Pos	AP, PSA, Micro	Neg	AP
E34ANN - 5902	Pos	ALS, AP, Micro	Neg	ALS, AP
EHXZJK - 5902	Pos	AP, PSA, Micro	Neg	AP, PSA, Micro
ELJJPP - 5902	Pos	ALS, AP, Micro, PSA	Neg	ALS, AP, Micro, PSA
EY6YTN - 5902	Pos	ALS, AP, Micro	Neg	ALS, AP
F7H8UP - 5902	Pos	AP, MICRO	Neg	AP, MICRO
F8XQDM - 5902	Pos	ALS,AP, Micro	Neg	ALS,AP,Micro
FPXE6L - 5902	Pos	AP, PSA	Neg	AP
GCH7HK - 5902	Pos	ALS, AP, Micro	Neg	ALS, AP
H484WJ - 5902	Pos	ALS, AP, micro		
HEQHYL - 5902	Pos	ALS, AP, Micro	Neg	ALS, AP
HJX7CN - 5901	Pos	P30	Neg	P30
J37QNH - 5902	Pos	AP, PSA, Micro	Neg	AP
J772JG - 5902	Pos	AP, Micro, PSA (p30)	Neg	AP, Micro, PSA (p30)
JHLQHM - 5902	Pos	PSA	Neg	PSA
JPGUBX - 5901	Pos	ALS, AP	Neg	ALS, AP
K9ECCE - 5902	Pos	AP, RSID Semen	Neg	AP
KPDQAK - 5902	Pos	AP, Micro	Neg	AP
L67JVL - 5901	Pos	AP, Micro, PSA	Neg	AP
L8XEJE - 5901	Pos	AP	Neg	AP
LBAKVF - 5902	Pos	AP, PSA	Neg	AP
LG3PNJ - 5901	Pos	PSA	Neg	PSA
LHUN3H - 5902	Pos	ALS, AP, Micro	Neg	ALS, AP, Micro
LUPUWF - 5901	Pos	AP, Micro	Neg	AP
LYKQJL - 5901	Pos	AP, Micro, PSA	Neg	AP
MW99ZF - 5902	Pos	ALS, AP, Micro	Neg	ALS, AP, Micro
NWNRKD - 5901	Pos	AP, Micro	Neg	AP
P3QH3A - 5901	Pos	Місгозсору, АР	Neg	Microscopy, P30 (PSA), AP
P8KEPF - 5902	Pos	AP, micro	Neg	AP
Q99BQE - 5901	Pos	Seratec PSA + Sperm Tracker	Neg	Seratec PSA + Sperm Tracker
R3MFPB - 5901	Pos	AP, micro	Neg	AP

TABLE 1b

Participants: 74

Semen Screening Results							
WebCode - Test	lte	m 3	Ite	m 4			
R9RDYE - 5901	Pos	AP, Micro, PSA	Neg	AP			
RCPXZD - 5901	Pos	AP, PSA, Micro	Neg	AP			
TMDEV4 - 5902	Pos	AP, Micro	Neg	AP			
TZBZVA - 5901	Pos	AP, Micro	Neg	AP			
UNVR99 - 5902	Pos	ALS, AP, Micro	NT				
VFHZHA - 5902	Pos	AP, micro	Neg	AP			
VK9834 - 5902	Pos	Acid phosfatase, RSID Semen, Sperm Hylighter Express	Neg	Acid phosfatase, RSID Semen, Sperm Hylighter Express			
X28843 - 5901	Pos	AP, Microscopy	Neg	AP			
XCU6U9 - 5902	Pos	p30, sperm search	Neg	р30			
YGJ6L9 - 5901	Pos	AP, PSA, Micro	Neg	AP			
YJMUQ3 - 5901	Pos	AP, PSA, Micro	Neg	AP, PSA, Micro			
Z8LUQ4 - 5902	Pos	ALS, AP, Micro	Neg	ALS, AP			

Table 1b: Serology Screening Response Summary - Semen

This summary table excludes the count of participants who did not report or reported "Not Tested" for Item 3 and/or Item 4. Therefore, participant total may not align with totals shown below.

	Item 3	ltem 4	
Positive	74	0	
Negative	0	71	
Inconclusive	0	0	

Serology Screening Results

Indicate the results of any screening tests performed on the questioned stains (Items 3 & 4).

TABLE 1c							
		Saliva Screeni	ng Results				
WebCode - Test	lte	m 3	Ite	Item 4			
3MQZUY - 5902	Neg	Phadebas	Neg	Phadebas			
3WHFJ3 - 5902	Neg	Phadebas	Neg	Phadebas			
4BEUJW - 5902	Neg	RSID Saliva	Neg	RSID Saliva			
77A6RY - 5901	Neg	SALIgAE	NT				
786NUZ - 5901	Neg	Seratec Amalase Test	Neg	Seratec Amalase Test			
8YUL9Z - 5901	Neg	ALS, RSID Saliva, Seratec PAM	Neg	ALS, RSID Saliva, Seratec PAM			
B7W38V - 5901	Neg	SALIGAE	NT				
BKHHBU - 5901	Neg	SaligAE	NT				
CDTG9M - 5901	Neg	RSID	Neg	RSID			
CRBAWU - 5901	Pos	Amylase	Neg	Amylase			
EHXZJK - 5902	Neg	RSID saliva	Neg	RSID saliva			
F7H8UP - 5902	Neg	PHADEBAS	Neg	PHADEBAS			
FPXE6L - 5902	Neg	Phadebas press	Neg	Phadebas press			
J37QNH - 5902	NT		Neg	RSID			
J772JG - 5902	Neg	RSID saliva	Neg	RSID saliva			
JHLQHM - 5902	Neg	RSID Saliva	Neg	RSID Saliva			
KPDQAK - 5902	Neg	Phadebas	Neg	Phadebas			
LBAKVF - 5902	Neg	Phadebas press	Neg	Phadebas press			
LUPUWF - 5901	Neg	Phadebas	Neg	Phadebas			
NWNRKD - 5901	Neg	Phadebas	Neg	Phadebas			
P8KEPF - 5902	Neg	Phadebas	Neg	Phadebas			
R3MFPB - 5901	Neg	Phadebas	Neg	Phadebas			
TZBZVA - 5901	Neg	Phadebas	Neg	Phadebas			
VFHZHA - 5902	Neg	Phadebas	Neg	Phadebas			
VK9834 - 5902	Neg	RSID Saliva	Neg	RSID Saliva			
XCU6U9 - 5902	Neg	Amylase	Neg	amylase			

Table 1c: Serology Screening Response Summary - Saliva

Participants: 26

This summary table excludes the count of participants who did not report or reported "Not Tested" for Item 3 and/or Item 4. Therefore, participant total may not align with totals shown below.

	Item 3	Item 4	
Positive	1	0	
Negative	24	23	
Inconclusive	0	0	

Serology Screening Results

Indicate the results of any screening tests performed on the questioned stains (Items 3 & 4).

TABLE 1d							
	Human Origin Screening F	Results					
WebCode - Test	Item 3	Item 4					
2QGMC2 - 5902	Pos HemaTrace, p30	NT					
3MQZUY - 5902	Pos HemaTrace	Pos HemaTrace					
44PDRD - 5901	Pos Hematrace	Pos Hematrace					
4BEUJW - 5902	Pos Quantifiler trio, PP21	Pos Quantifiler trio, PP21					
6PTH6Y - 5902	Pos P30	Pos Hematrace					
786NUZ - 5901	Pos Ouchterlony Double Immunodiffusion	Pos Ouchterlony Double Immunodiffusion					
7H4JRV - 5902	Pos p30	Pos Hematrace					
8Y8DYV - 5902	Pos Hematrace	Neg P30					
8YUL9Z - 5901	Pos FOB, Seratec PMB, RSID Semen, Seratec PAM	Pos FOB, Seratec PMB					
CDTG9M - 5901	Pos RSID-Blood	Pos RSID-Blood					
CKECAQ - 5902	Pos HEMATRACE, P30						
CXYRCQ - 5902	Pos P30	Pos Hematrace					
E34ANN - 5902	Pos Hematrace, PSA	NT					
EHXZJK - 5902	Pos Quant trio, PowerPlex 21	Pos Quant trio, PowerPlex 21					
ELJJPP - 5902	NT	Pos Hematrace					
EY6YTN - 5902	Pos pSA	Pos Hematrace					
F8XQDM - 5902	Pos PSA,HemaTrace	Neg PSA					
FPXE6L - 5902	Pos ABACard Hematrace	NT					
GCH7HK - 5902	Pos PSA	Pos Hematrace					
H484WJ - 5902	Pos PSA	Pos Hematrace					
HEQHYL - 5902	Pos Hematrace, P30	NT					
HJX7CN - 5901	Pos Hema Trace ABA Card	Pos Hema Trace ABA Card					
J37QNH - 5902	Pos QT, PP21	Pos QT, PP21					
J772JG - 5902	Pos PowerPlex21, YFilerPlus, Quant Trio	Pos PowerPlex21, YFilerPlus, Quant Trio					
JHLQHM - 5902	Pos Obti, PSA	Pos Obti, PSA					
K9ECCE - 5902	Pos Quantifiler Trio	Pos Quantifiler Trio					
LBAKVF - 5902	NT	Pos ABAcard					
LG3PNJ - 5901	Pos Quantiplex Pro RGQ, STR	Pos Quantiplex Pro RGQ, STR					
LHUN3H - 5902	NT	Neg P30					
LUPUWF - 5901	Pos PowerQuant	Pos PowerQuant					
MW99ZF - 5902	Pos p30	Pos p30					
NWNRKD - 5901	Pos PowerQuant	Pos PowerQuant					
R3MFPB - 5901	Pos PowerQuant	Pos PowerQuant					

Test 25-5901/2

Participants: 37

TABLE 1d

Human Origin Screening Results					
WebCode - Test	Item 3	Item 4			
TZBZVA - 5901	Pos PowerQuant	Pos PowerQuant			
UNVR99 - 5902	Pos Hematrace, P30	NT			
VK9834 - 5902	Pos Quantifiler Trio	Pos Quantifiler Trio			
Z8LUQ4 - 5902	Pos PSA	Pos Hematrace			

Table 1d: Serology Screening Response Summary - Human Origin

This summary table excludes the count of participants who did not report or reported "Not Tested" for Item 3 and/or Item 4. Therefore, participant total may not align with totals shown below.

	Item 3	Item 4
Positive	34	28
Negative	0	3
Inconclusive	0	0

Serology Screening Results

Indicate the results of any screening tests performed on the questioned stains (Items 3 & 4).

TABLE 1e							
		Y Screeni	ng Results				
WebCode - Test	lte	m 3	Ite	m 4			
3MQZUY - 5902	Pos	PlexorHY	Pos	PlexorHY			
44PDRD - 5901	Pos	Quant Trio	Pos	Quant Trio			
8Y8DYV - 5902	Pos	MDS & qPCR procedure	Pos	MDS & qPCR procedure			
9CLYK9 - 5901	Pos	Quant Trio	Pos	Quant Trio			
CY6DP6 - 5901	Pos	Quantifiler Trio	Pos	Quantifiler Trio			
EHXZJK - 5902	Pos	Quant trio, Yfiler Plus	Pos	Quant trio, Yfiler Plus			
F8XQDM - 5902	Pos	MDS & qPCR procedure	Pos	MDS & qPCR procedure			
FPXE6L - 5902	Pos	Plexor HY	Pos	Plexor HY			
H484WJ - 5902	Pos	MDS&qPCR procedure	Pos	MDS&qPCR procedure			
HJX7CN - 5901	Pos	Y-FILER-plus	Pos	Y-FILER-plus			
J37QNH - 5902	Pos	QT, PP21	Pos	QT, PP21			
J772JG - 5902	Pos	YFilerPlus, Quant Trio	Pos	YFilerPlus, Quant Trio			
JHLQHM - 5902	Pos		Pos				
JPGUBX - 5901	Pos	QuantTrio	NT				
K9ECCE - 5902	Pos	Quantifiler Trio	Pos	Quantifiler Trio			
LBAKVF - 5902	Pos	Plexor HY	Pos	Plexor HY			
LG3PNJ - 5901	Pos	Y-STR	Pos	Y-STR			
LUPUWF - 5901	Pos	PowerQuant	Pos	PowerQuant			
NWNRKD - 5901	Pos	PowerQuant	Pos	PowerQuant			
P8TP7Q - 5901	Pos	Quant Trio	Pos	Quant Trio			
R3MFPB - 5901	Pos	PowerQuant	Pos	PowerQuant			
TZBZVA - 5901	Pos	PowerQuant	Pos	PowerQuant			
VK9834 - 5902	Pos	Quantifiler Trio	Pos	Quantifiler Trio			
W2RLKZ - 5901	Pos	Quant Trio	Pos	Quant Trio			
WRGKYZ - 5901	Pos	Quantifiler Trio	Pos	Quantifiler Trio			
XCU6U9 - 5902	Pos	y-screen	Pos	y-screen			

Table 1e: Serology Screening Response Summary - Y Screening Participants: 26 This summary table excludes the count of participants who did not report or reported "Not Tested" for Item 3 and/or Item 4. Therefore, participant total may not align with totals shown below. Participant

	Item 3	Item 4
Positive	26	25
Negative	0	0
Inconclusive	0	0

Serology Screening Results

Indicate the results of any screening tests performed on the questioned stains (Items 3 & 4).

TABLE 1f							
Other Screening Results							
WebCode - Tes	t Ite	em 3	Ite	em 4			
8YUL9Z - 5901	Menstrual blood No	eg Seratec PMB	Menstrual blood N	leg Seratec PMB			
K9ECCE - 5902	Spermatozoa Po	DS Microscopy					
LG3PNJ - 5901	sperm cells Po	os Micro	sperm cells N	leg Micro			

DNA Interpretations

Based on results obtained from DNA analysis, could the Victim (Item 1) and/or the Suspect (Item 2) be a contributor to the questioned stains (Items 3 & 4)?

WebCode-	bCode- <u>Victim (Item 1)</u> <u>Suspect (Item 2)</u>		WebCode-	Victim (Item 1)		Suspect (Item 2)			
lest	Item 3	Item 4	Item 3	Item 4	lest	ltem 3	Item 4	ltem 3	Item 4
2QGMC2 - 5902	Yes	No	Yes	Yes	9CLYK9 - 5901	Yes	No	Yes	Yes
34G3B3 - 5902	Yes	No	Yes	Yes	9MBR8Q - 5901	Yes	No	Yes	Yes
37J2CU - 5901	Yes	No	Yes	Yes	9WLTTR - 5902	Yes	No	Yes	Yes
3MQZUY - 5902	Yes	No	Yes	Yes	B7W38V - 5901	Yes	No	Yes	Yes
3WHFJ3 - 5902	Yes	No	Yes	Yes	BDR7Z7 - 5901	Yes	No	Yes	Yes
44PDRD - 5901	Yes	No	Yes	Yes	BKHHBU - 5901	Yes	No	Yes	Yes
4BEUJW - 5902	Yes	No	Yes	Yes	BNGT8U - 5902	Yes	No	Yes	Yes
4FBJY4 - 5902	Yes	No	Yes	Yes	CDTG9M - 5901	Yes	No	Yes	Yes
4UQG64 - 5901	Yes	No	Yes	Yes	CKECAQ - 5902	Yes	No	Yes	Yes
4XQRZ4 - 5901	Yes	No	Yes	Yes	CRBAWU - 5901	Yes	No	Yes	Yes
62K9Q4 - 5901	Yes	No	Yes	Yes	CRVQCW - 5901	Yes	No	Yes	Yes
63VBC2 -	Yes	No	Yes	Yes	CXYRCQ - 5902	Yes	No	Yes	Yes
6PTH6Y -	Yes	No	Yes	Yes	CY6DP6 - 5901	Yes	No	Yes	Yes
77A6RY -	Yes	No	Yes	Yes	CZ6EPV - 5901	Yes	No	Yes	Yes
786NUZ -	Yes	No	Yes	Yes	E34ANN - 5902	Yes	No	Yes	Yes
7H4JRV -	Yes	No	Yes	Yes	EHXZJK - 5902	Yes	No	Yes	Yes
8DDAHX -	Yes	No	Yes	Yes	ELJJPP - 5902	Yes	No	Yes	Yes
8Y8DYV -	Yes	No	Yes	Yes	EY6YTN - 5902	Yes	No	Yes	Yes
8YUL9Z -	Yes	No	Yes	Yes	F7H8UP - 5902	Yes	No	Yes	Yes
5701					F8XQDM -	Yes	No	Yes	Yes

5902

TABLE 2

WebCode-	<u>Victim</u>	<u>Victim (Item 1)</u>		<u>(Item 2)</u>	WebCode-	<u>Victim (</u>	Victim (Item 1)		Suspect (Item 2)	
Test	Item 3	Item 4	Item 3	Item 4	Test	Item 3	Item 4	Item 3	Item 4	
FPXE6L - 5902	Yes	No	Yes	Yes	P8KEPF - 5902	Yes	No	Yes	Yes	
GCH7HK - 5902	Yes	No	Yes	Yes	P8TP7Q - 5901	Yes	No	Yes	Yes	
H484WJ - 5902	Yes	No	Yes	Yes	PD88ZF - 5902	Yes	No	Yes	Yes	
HEQHYL - 5902	Inc	No	Yes	Yes	Q99BQE - 5901	Yes	No	Yes	Yes	
HJX7CN - 5901	Yes	No	Yes	Yes	R3MFPB - 5901	Yes	No	Yes	Yes	
J37QNH - 5902	Yes	No	Yes	Yes	R9RDYE - 5901	Yes	No	Yes	Yes	
J772JG - 5902	Yes	No	Yes	Yes	RCPXZD - 5901	Yes	No	Yes	Yes	
JHLQHM - 5902	Yes	No	Yes	Yes	TMDEV4 - 5902	Yes	No	Yes	Yes	
JPGUBX - 5901	Yes	No	Yes	Yes	TZBZVA - 5901	Yes	No	Yes	Yes	
K9ECCE - 5902	Yes	No	Yes	Yes	UNVR99 - 5902	Yes	No	Yes	Yes	
KPDQAK - 5902	Yes	No	Yes	Yes	VFHZHA - 5902	Yes	No	Yes	Yes	
L67JVL - 5901	Yes	No	Yes	Yes	VK9834 - 5902	Yes	No	Yes	Yes	
L8XEJE - 5901	Yes	No	Yes	Yes	W2RLKZ - 5901	Yes	No	Yes	Yes	
LBAKVF - 5902	Yes	No	Yes	Yes	WRGKYZ - 5901	Yes	No	Yes	Yes	
LG3PNJ - 5901	Yes	No	Yes	Yes	WU8M28 - 5901	Yes	No	Yes	Yes	
LHUN3H - 5902	Yes	No	Yes	Yes	X28843 - 5901	Yes	No	Yes	Yes	
LUPUWF - 5901	Yes	No	Yes	Yes	XCU6U9 - 5902	No Interp	No Interp	Inc	Inc	
LYKQJL - 5901	Yes	No	Yes	Yes	YGJ6L9 - 5901	Yes	No	Yes	Yes	
MW99ZF - 5902	Yes	No	Yes	Yes	YJMUQ3 - 5901	Yes	No	Yes	Yes	
nwnrkd - 5901	Yes	No	Yes	Yes	Z8LUQ4 - 5902	Yes	No	Yes	Yes	
P3QH3A - 5901	Yes	No	Yes	Yes						

DNA Interpretation								
Response Summary Participants reporting DNA results:								
Based on results obtained from D the questioned stains (Items 3 & 4	NA analysis, 4)?	, could the Vict	im (Item 1) and/	or the Suspe	ect (Item 2) be a contributor to			
	Victim (Item 1		Suspect (Item 2)					
	Item 3	Item 4	Item 3	Item 4				
Yes	78	0	79	79				
No	0	79	0	0				
Inc	1	0	1	1				
No Interpretation	1	1	0	0				
No Response	0	0	0	0				

STR Amplification Kit(s) & Results

	TABLE 3							
WebCod	e - Test	Amplification Kits -	Probablistic Gene	otyping Software				
	D1S1656	D2S1338	D2S441	D3S1358	D5S818	D6S1043		
Item	D7S820	D8S1179	D10S1248	D12S391	D13S317	D16S539		
	D18551	D195433	D21511	D22S1045	Amelogenin	CSF1PO		
	FGA	Penta D	Penta E	SE33	TH01	трох		
	VWA	013371		Dission	i indei			
206402	2 - 5902	GlobalFiler™ Express- Tri	ITEM I - JIK	Results				
2001102	11.17.3	17.25		16.18	10.12			
1	10,12	13,15	13	16,17	8,12	9,12		
1	14,17	12,13	31.2,32.2	11,12	X,X	10,11		
	21,24		,	17,24.2	6,9.3	8,8		
	16,16							
34G3B3 -	- 5902	GlobalFiler™						
	11,17.3	17,25	11	16,18	10,12			
1	10,12	13,15	13	16,17	8,12	9,12		
	14,17	12,13	31.2,32.2	11,12	Х	10,11		
	21,24			17,24.2	6,9.3	8		
	16							
37J2CU -	5901	GlobalFiler™- STRMix™						
	11,17.3	17,25	11,11	16,18	10,12			
1	10,12	13,15	13,13	16,17	8,12	9,12		
	14,17	12,13	31.2,32.2	11,12	X,X	10,11		
	21,24			17,24.2	6,9.3	8,8		
	16,16							
3MQZUY	- 5902	Identifiler® Plus						
		17,25		16,18	10,12			
1	10,12	13,15			8,12	9,12		
	14,17	12,13	31.2,32.2		X,X	10,11		
	21,24				6,9.3	8,8		
	16,16							
3WHFJ3 -	5902	GlobalFiler™- STRMix™ V	/2.9.1					
	11,17.3	17,25	11	16,18	10,12			
1	10,12	13,15	13	16,17	8,12	9,12		
	14,17	12,13	31.2,32.2	11,12	X	10,11		
	21,24			17,24.2	6,9.3	8		
	16							
44PDRD -	5901	GlobalFiler™		1 (1)	10.10			
	11,17.3	17,25	11	16,18	10,12	0.10		
1	10,12	13,15	13	16,17	8,12	9,12		
	14,1/	12,13	31.2,32.2	11,12	Χ	10,11		
	21,24			17,24.2	0,7.3	ŏ		
	10							

			TABLE	3		
WebCo	de - Test	Amplification Kits - P	robablistic Gen	otyping Software		
	D1\$1656	D251338	D25441	D3\$1358	D55818	D651043
ltem	D75820	D8S1179	D10S1248	D125391	D135317	D165539
	D18S51	D19S433	D21S11	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	vWA	DYS391	DYS570	DYS576	Y Indel	
	5000		Item I - SIR	Results		
4BEUJW	- 5902	PowerPlex® 21- 51 KMix ***	2.10.0	14.10	10.10	
,	11,17.3	17,25		16,18	10,12	0.10
I	10,12	13,15	21.0.20.0	10,17	8,12	9,12
	14,17	12,13	31.2,32.2		Χ,Χ	10,11
	21,24	9,11	8,11		0,9.3	8,8
	10,10					
4FBJY4 -	5902	PowerPlex® Fusion 6C - L	JINAXS	14.10	10.10	
,	11,17.3	17,25	11,11	16,18	10,12	0.10
I	10,12	13,15	13,13	10,17	8,12	9,12
	14,17	0 1 1	9.11	17.24.2	۸,۸ ۵.0.3	10,11
	16.16	9,11	0,11	17,24.2	0,7.3	0,0
411004	4 5001					
40000	4 - 3901		11	14 10	10.10	
1	10.12	12.15	12	16,10	10,12	0.10
I	10,12	10,10	21 0 20 0	10,17	0,12	9,12
	14,17	12,13	31.2,32.2	17.24.2	A 4 0 2	0,11
	16			17,24.2	0,7.5	0
	5001					
47QNZ4	- 3701		11	16 19	10.12	
1	10.12	12.15	12	16,17	R 10	0 1 2
I	14.17	12.13	31 2 32 2	11.12	ν	10.11
	21.24	12,10	01.2,02.2	17.24.2	693	8
	16			17721.2	0,7.0	3
628904	- 5901	GlobalEiler™- STRMix™				
02107004	11 17 3	17.25	11	16.18	10.12	
1	10.12	13 15	13	16.17	8.12	912
	14.17	12.13	31.2.32.2	11.12	X	10.11
	21.24	,		17.24.2	6,9,3	8
	16			,	,	
63VBC2	- 5901	GlobalFiler™				
-	11,17,3	17.25	11	16,18	10,12	
1	10,12	13,15	13	16,17	8,12	9,12
	14,17	12,13	31.2,32.2	11,12	X	10,11
	21,24			17,24.2	6,9.3	8
	16					
6PTH6Y	- 5902	GlobalFiler™ Express- True	eAllele® VUler Rele	ase 2022b		
	11,17.3	17,25	11,11	16,18	10,12	
1	10,12	13,15	13	16,17	8,12	9,12
	14,17	12,13	31.2,32.2	11,12	X	10,11
	21,24			17,24.2	6,9.3	8,8
	16,16					

			TABLE	3		
WebCod	de - Test	Amplification Kits -	Probablistic Gen	otyping Software		
	D151656	D251338	D25441	D351358	D55818	D651043
ltem	D75820	D851179	D10S1248	D125391	D135317	D165539
	D18S51	D195433	D21S11	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	vWA	DYS391	DYS570	DYS576	Y Indel	
774401/	5001	ClabalE:lau™	Item I - SIR	Results		
//AORT-	- 5901		11 11	14 10	10.10	
1	10.12	17,25	1212	16,17	9.12	0 1 2
I	10,12	12.13	31 2 32 2	10,17	0,12	10.11
	21.24	12,10	51.2,52.2	17 24 2	693	8.8
	16.16			17,27.2	0,7.0	0,0
786NU7	- 5901	Identifiler® Plus - EuroEc	or Mix			
7001102	0701	17 25		16 18	10.12	
1	10.12	13.15		10,10	8.12	9.12
	14,17	12,13	31.2,32.2		X,X	10,11
	21,24	,	,		6,9.3	8,8
	16,16					
7H4JRV -	- 5902	GlobalFiler™ Express- Tr	ueAllele® VUler Rele	ase 2022b		
	11,17.3	17,25	11	16,18	10,12	
1	10,12	13,15	13	16,17	8,12	9,12
	14,17	12,13	31.2,32.2	11,12	Х	10,11
	21,24			17,24.2	6,9.3	8,8
	16,16					
8DDAHX	- 5901	GlobalFiler™- STRMix™	2.8			
	11,17.3	17,25	11	16,18	10,12	
1	10,12	13,15	13	16,17	8,12	9,12
	14,17	12,13	31.2,32.2	11,12	Х	10,11
	21,24			17,24.2	6,9.3	8
	16					
8Y8DYV	- 5902	GlobalFiler™- TrueAllele	® VUler Release 202	22b		
	11,17.3	17,25	11,11	16,18	10,12	
1	10,12	13,15	13,13	16,17	8,12	9,12
	14,17	12,13	31.2,32.2	11,12	X,X	10,11
	21,24			17,24.2	6,9.3	8,8
	16,16					
8YUL9Z -	- 5901	PowerPlex® Fusion 6C, I	NGM Detect - LRmix			
	11,17.3	17,25	11,11	16,18	10,12	
1	10,12	13,15	13,13	16,17	8,12	9,12
	14,17	12,13	31.2,32.2	11,12	X,X	10,11
	21,24	9,11	8,11	17,24.2	6,9.3	8,8
	16,16		-	-	-	
YCLYKY -	- 5901	GlobalFiler - 51 KMix		1/ 10	10.10	
1	11,17.3	17,25	12	16,18	10,12	0.10
I	10,12	13,15	31.0.20.0	10,17	8,12 V	9,12
	- 21-24	12,13	31.2,32.2	17.24.2	۸ ۸	Ω
	16	NR		17,24.2	NR	0
	10	1.315			1.115	

			TABLE	3		
WebCode	e - Test	Amplification Kits -	Probablistic Gen	otyping Software		
	D161656	D251228	D25441	D251259	D55919	D6510/2
Item	D75820	D251338	D1051248	D125391	D135317	D165539
	D18S51	D195433	D21511	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	vWA	DYS391	DY\$570	DYS576	Y Indel	
			ltem 1 - STR	Results		
9MBR8Q -	5901	PowerPlex® 21				
	11,17.3	17,25		16,18	10,12	11,11
1	10,12	13,15		16,17	8,12	9,12
	14,17	12,13	31.2,32.2		X,X	10,11
	21,24	9,11	8,11		6,9.3	8,8
011/1 770	16,16					
9WLIIR - 5	5902	PowerPlex® 21				
	11,17.3	17,25		16,18	10,12	,
	10,12	13,15	21.0.20.0	16,17	8,12	9,12
	14,17	12,13	31.2,32.2		λ,λ	10,11
	21,24	9,11	0,11		0,7.3	0,0
	5001					
D/ WJOV -	11 17 2		11 11	14 10	10.10	
1	10.12	12.15	12.12	16,10	0,12	0.10
	14.17	12.13	31 2 32 2	10,17	0,12	9,12
	21.24	12,10	51.2,52.2	17.24.2	693	8.8
	1616			17,27.2	0,7.0	0,0
	5901	PowerPlex® Eusion 6C	GlobalEiler™ Express			
	11 17 3	17 25	11	16 18	10.12	
1	10.12	13 15	13	16,17	8.12	912
	14,17	12,13	31.2.32.2	11,12	X	10,11
	21,24	9,11	8,11	17,24.2	6,9.3	8
	16					
BKHHBU -	5901	GlobalFiler™				
	11,17.3	17,25	11,11	16,18	10,12	
1	10,12	13,15	13,13	16,17	8,12	9,12
	14,17	12,13	31.2,32.2	11,12	X,X	10,11
	21,24			17,24.2	6,9.3	8,8
_	16,16					
BNGT8U -	5902	GlobalFiler™				
	11,17.3	17,25	11	16,18	10,12	
1	10,12	13,15	13	16,17	8,12	9,12
_	14,17	12,13	31.2,32.2	11,12	Х	10,11
	21,24			17,24.2	6,9.3	8
	16					
CDTG9M -	- 5901	GlobalFiler™				
	11,17.3	17,25	11,11	16,18	10,12	-
1	10,12	13,15	13,13	16,17	8,12	9,12
	14,17	12,13	31.2,32.2	11,12	X,X	10,11
	21,24	-	-	17,24.2	6,9.3	8,8
	16,16	-	-	-	-	

			TABLE	3		
WebCoo	de - Test	Amplification Kits -	Probablistic Gen	otyping Software		
	D151656	D251338	D25441	D3\$1358	D5\$818	D651043
Item	D7S820	D8S1179	D1051248	D125391	D13S317	D16S539
	D18S51	D19S433	D21S11	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	VWA	DYS391	DYS570	DYS576	Y Indel	
CVECAO	5002	ClabalE:lar™ Everage Tr	Item I - SIK	Kesults		
CRECAQ	11 17 2	Giobali lier Express- In		14 19	10.10	
1	10.12	17,25	13 13	16,17	9.12	0.10
I	10,12	10.13	21 0 20 0	11.12	0,12	10.11
	21.24	12,15	51.2,52.2	17.24.2	693	8.8
	16.16			17,27.2	0,7.0	0,0
	10,10	PowerPlay® FSI 17 Fast	LiRa v3 0			
CREATIO	11 17 3	17 25	11 11	16.18		
1	11,17.5	13 15	13 13	16,17		912
I	14 17	12 13	31 2 32 2	11 12	ХX	/,12
	21.24	12,10	01.2,02.2	17.24.2	6.9.3	
	16,16			,	-,	
	V - 5901	GlobalFiler™				
	11.17.3	17.25	11	16.18	10.12	
1	10,12	13,15	13	16,17	8,12	9,12
	14,17	12,13	31.2,32.2	11,12	X	10,11
	21,24			17,24.2	6,9.3	8
	16					
CXYRCQ	- 5902	GlobalFiler™ Express				
	11,17.3	17,25	11,11	16,18	10,12	
1	10,12	13,15	13	16,17	8,12	9,12
	14,17	12,13	31.2,32.2	11,12	Х	10,11
	21,24			17,24.2	6,9.3	8,8
	16,16					
CY6DP6	- 5901	GlobalFiler™				
	11,17.3	17,25	11	16,18	10,12	
1	10,12	13,15	13	16,17	8,12	9,12
	14,17	12,13	31.2,32.2	11,12	Х	10,11
	21,24			17,24.2	6,9.3	8
	16					
CZ6EPV -	- 5901	GlobalFiler™				
	11,17.3	17,25	11	16,18	10,12	
1	10,12	13,15	13	16,17	8,12	9,12
	14,17	12,13	31.2,32.2	11,12	Х	10,11
	21,24			17,24.2	6,9.3	8
	16					
E34ANN	- 5902	GlobalFiler [™] Express- Tru	ueAllele® VUler Rele	ease 2022b		
	11,17.3	17,25	11,11	16,18	10,12	
1	10,12	13,15	13,13	16,17	8,12	9,12
	14,17	12,13	31.2,32.2	11,12	X,X	10,11
	21,24			17,24.2	6,9.3	8,8
	16,16					

			TABLE	3		
WebCod	e - Test	Amplification Kits -	Probablistic Gen	otyping Software		
	D1S1656	D251338	D2S441	D3\$1358	D5\$818	D651043
ltem	D75820	D8S1179	D10S1248	D125391	D135317	D165539
	D18S51	D19S433	D21S11	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	vWA	DYS391	DYS570	DYS576	Y Indel	
	5000			Results		
ΕΠΧΖΙΚ -	3902	PowerPiex® 21- 51 KMIX	2.10	17.10	10.10	
1	10.12	17,25		16,16	0,12	0.12
	14.17	10.03	31 0 30 0	10,17	0,12	9,12
	21.24	0 11	8 11		693	8.8
	16.16	7,11	0,11		0,7.5	0,0
ELLIPP 5	002	GlobalEilar™ Express Tr	ua Allala® VI llar Pala	asa 2022h		
	11 17 3	17.25		16.18	10.12	
1	10.12	13 15	13	16,17	8.12	912
	14 17	12 13	31 2 32 2	11 12	X X	10 11
	21.24	12,10	01.2,02.2	17.24.2	6.9.3	8.8
	16,16			,	-,	-/-
EY6YTN -	5902	GlobalFiler™ Express- Tr	ueAllele® VUler Rele	ease 2022b		
2.0	11.17.3	17.25	11.11	16.18	10.12	
1	10.12	13,15	13,13	16,17	8.12	9,12
	14,17	12,13	31.2,32.2	11,12	X,X	10,11
	21,24	,	,	17,24.2	6,9.3	8,8
	16,16					
F7H8UP -	5902	GlobalFiler™- STRMix™	v2.9.1			
	11,17.3	17,25	11	16,18	10,12	
1	10,12	13,15	13	16,17	8,12	9,12
	14,17	12,13	31.2,32.2	11,12	Х	10,11
	21,24			17,24.2	6,9.3	8
	16					
F8XQDM	- 5902	GlobalFiler™- TrueAllele	® VUler Release 202	22b		
	11,17.3	17,25	11,11	16,18	10,12	
1	10,12	13,15	13,13	16,17	8,12	9,12
	14,17	12,13	31.2,32.2	11,12	X,X	10,11
	21,24			17,24.2	6,9.3	8,8
	16,16					
FPXE6L - 5	5902	Identifiler® Plus				
		17,25		16,18	10,12	
1	10,12	13,15			8,12	9,12
_	14,17	12,13	31.2,32.2		X,X	10,11
	21,24				6,9.3	8,8
	16,16					
GCH7HK	- 5902	GlobalFiler™ Express- Tr	ueAllele® VUler Rele	ease 2022b		
	11,17.3	17,25	11,11	16,18	10,12	
1	10,12	13,15	13,13	16,17	8,12	9,12
	14,17	12,13	31.2,32.2	11,12	X,X	10,11
	21,24			17,24.2	6,9.3	8,8
	16,16					

			TABLE	3		
WebCoo	de - Test	Amplification Kits -	Probablistic Gen	otyping Software		
	D151656	D251338	D25441	D3\$1358	D5\$818	D651043
ltem	D75820	D8S1179	D10S1248	D12S391	D135317	D165539
	D18S51	D195433	D21511	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	vWA	DYS391	DYS570	DYS576	Y Indel	
	5000		Item I - SIR	Results		
H484WJ	- 5902	GlobalFiler Express- Iru	eAllele® VUler Kele	ase 2022b	10.10	
1	11,17.3	17,25	11,11	16,18	10,12	0.10
I	10,12	13,15	13,13	10,17	8,12	9,12
	21.24	12,13	31.2,32.2	17.24.2	۸,۸	10,11
	16.16			17,24.2	0,7.5	0,0
	5002	GlabalFilar™ Everage Tru	Allala® VI llar Pala	2022b		
HLQIIIL	11 17 3	17.25		14 19	10.12	
1	10.12	17,25	13.13	16,17	8.12	0 1 2
I	14 17	12.13	31 2 32 2	11 12	X X	10.11
	21 24	12,10	01.2,02.2	17 24 2	693	8.8
	16.16			17721.2	0,7.0	0,0
HIX7CN	- 5901	GlobalFiler™ IQC				
	11 17 3	17 25	11 11	16 18	10.12	
1	10.12	13.15	13.13	16,17	8.12	9.12
	14,17	12,13	31.2.32.2	11,12	X.X	10.11
	21.24	,		17.24.2	6,9.3	8,8
	16,16			,		,
J37QNH	- 5902	PowerPlex® 21- STRMix™	v2.10.0			
	11,17.3	17,25		16,18	10,12	11,11
1	10,12	13,15		16,17	8,12	9,12
	14,17	12,13	31.2,32.2		X,X	10,11
	21,24	9,11	8,11		6,9.3	8,8
	16,16					
J772JG -	- 5902	PowerPlex® 21- STRMix™	2.10			
	11,17.3	17,25		16,18	10,12	11,11
1	10,12	13,15		16,17	8,12	9,12
	14,17	12,13	31.2,32.2		X,X	10,11
	21,24	9,11	8,11		6,9.3	8,8
	16,16					
JHLQHM	- 5902	GlobalFiler™ - DNAxs				
	11,17.3	17,25	11,11	16,18	10,12	
1	10,12	13,15	13,13	16,17	8,12	9,12
	14,17	12,13	31.2,32.2	11,12	X,X	10,11
	21,24			17,24.2	6,9.3	8,8
	16,16					
JPGUBX	- 5901	GlobalFiler™				
	11,17.3	17,25	11	16,18	10,12	
1	10,12	13,15	13	16,17	8,12	9,12
	14,17	12,13	31.2,32.2	11,12	Х	10,11
	21,24			17,24.2	6,9.3	8
	16					

			TABLE	3		
WebCode	e - Test	Amplification Kits - F	Probablistic Gen	otyping Software		
	D1S1656	D251338	D25441	D3\$1358	D5\$818	D651043
ltem	D7S820	D8S1179	D10S1248	D12S391	D135317	D16S539
_	D18S51	D195433	D21S11	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	VWA	D12391			t Indel	
KOECCE	5002	PowerPlay® 21	Item I - SIR	Results		
R/LCCL -	11 17 3	17 25		16 18	10.12	11 11
1	10.12	13.15		16,17	8.12	9.12
	14.17	12.13	31.2.32.2	,.,	X.X	10.11
	21,24	9,11	8,11		6,9.3	8,8
	16,16					
KPDQAK -	5902	GlobalFiler™- STRMix™ v2	2.9.1			
	11,17.3	17,25	11	16,18	10,12	
1	10,12	13,15	13	16,17	8,12	9,12
_	14,17	12,13	31.2,32.2	11,12	Х	10,11
	21,24			17,24.2	6,9.3	8
	16					
L67JVL - 59	901	GlobalFiler™- STRMix™				
	11,17.3	17,25	11	16,18	10,12	
1	10,12	13,15	13	16,17	8,12	9,12
_	14,17	12,13	31.2,32.2	11,12	Х	10,11
	21,24			17,24.2	6,9.3	8
	16					
L8XEJE - 59	901	PowerPlex® 21				
_	11,17.3	17,25		16,18	10,12	11,11
1	10,12	13,15		16,17	8,12	9,12
	14,17	12,13	31.2,32.2		X,X	10,11
	21,24	9,11	8,11		6,9.3	8,8
	16,16					
LBAKVF - 5	902	Identifiler® Plus		17.10	10.10	
	10.10	17,25		10,18	0,12	0.10
	14.17	13,13	31 2 32 2		0,12	9,12
	21.24	12,10	01.2,02.2		693	8.8
	16.16				0,7.0	0,0
IG3PNI-	5901	PowerPlex® ESX17 Fast Sv	vstem- STRMix™ V2	511		
2001113	11.17.3	17.25	11.11	16.18		
1		13,15	13,13	16,17		9,12
	14,17	12,13	31.2,32.2	11,12	X,X	
	21,24			17,24.2	6,9.3	
	16,16					
LHUN3H -	5902	GlobalFiler™ Express- Tru	eAllele® VUler Rele	ase 2022b		
	11,17.3	17,25	11,11	16,18	10,12	
1	10,12	13,15	13	16,17	8,12	9,12
	14,17	12,13	31.2,32.2	11,12	Х	10,11
	21,24			17,24.2	6,9.3	8,8
	16,16					

			TABLE	3		
WebCo	de - Test	Amplification Kits -	Probablistic Gen	otyping Software		
	D151656	D251338	D25441	D3\$1358	D5\$818	D651043
ltem	D75820	D8S1179	D10S1248	D12S391	D135317	D165539
	D18S51	D195433	D21511	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	vWA	DYS391	DYS570	DYS576	Y Indel	
	5001		Item I - SIR	Results		
LUPUWF	- 5901	Identifiler® Plus, PowerPle	ex® Fusion 6C, Min	ifiler	10.10	
,	11,17.3	17,25	10	16,18	10,12	0.10
I	10,12	13,15	13	10,17	8,12	9,12
	21.24	0.11	SI.2,SZ.Z	17.24.2	A 0 3	10,11
	1.4	7,11	0,11	17,24.2	0,7.3	0
	5001					
LINQJL -	5901		11	14 10	10.10	
1	10.12	17,25	12	16,10	0,12	0.10
I	10,12	10,10	31 0 30 0	10,17	0,12	9,12
	21.24	12,15	51.2,52.2	17.24.2	693	8
	16			17,27.2	0,7.0	0
۸ <i>۸</i> \۸/007F	5002	GlobalFilar™ Express Tri	Allala® VI llar Pala	ugeo 2022h		
101007721	- 3702	17.25		16 18	10.12	
1	10.12	17,25	13.13	16,17	8.12	0 1 2
I	10,12	12.13	31 2 32 2	11,12	0,12 X X	10.11
	21.24	12,15	01.2,02.2	17.24.2	693	8.8
	16.16			17,27.2	0,7.0	0,0
	5901	Idantifilar® Plus PowarPl	NR Fusion 6C Min	Filor		
	11 17 3	17 25		16.18	10.12	
1	10.12	17,25	13	16,17	8.12	0 1 2
I	14.17	12,13	31 2 32 2	11 12	X	10.11
	21.24	9.11	8 1 1	17.24.2	693	8
	16	,,	0,		0,7.0	
	- 5901	PowerPlex® 21_ STRMiv™	28			
	11 17 3	17 25	2.0	16 18	10.12	11 11
1	10.12	13.15		16,17	8.12	9.12
	14,17	12,13	31.2.32.2	/	X.X	10,11
	21.24	9,11	8,11		6,9.3	8,8
	16,16	,			,	,
P8KEPF -	5902	GlobalFiler™- STRMix™				
	11,17,3	17.25	11	16,18	10,12	
1	10,12	13,15	13	16,17	8,12	9,12
	14,17	12,13	31.2,32.2	11,12	X	10,11
	21,24			17,24.2	6,9.3	8
	16					
P8TP7Q	- 5901	GlobalFiler™				
	11,17.3	17,25	11	16,18	10,12	
1	10,12	13,15	13	16,17	8,12	9,12
	14,17	12,13	31.2,32.2	11,12	X	10,11
	21,24			17,24.2	6,9.3	8
	16					

			TABLE	3		
WebCo	de - Test	Amplification Kits - F	Probablistic Gen	otyping Software		
	D151656	D251338	D25441	D3\$1358	D5\$818	D651043
ltem	D75820	D8S1179	D10S1248	D125391	D13S317	D16S539
	D18S51	D19S433	D21S11	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	vWA	DYS391	DYS570	DYS576	Y Indel	
	5000			Results		
PD88ZF	- 5902		2.8	17.10	10.10	
,	11,17.3	17,25	11	16,18	10,12	0.10
I	10,12	13,15	13	10,17	8,12	9,12
	14,17	12,13	31.2,32.2	11,12	λ,λ	10,11
	21,24			17,24.2	0,7.3	0
	= 5001					
Q99DQI	- 5901		11 11	14 10	10.10	
1	10,12	17,25	12.12	16,18	10,12	0.10
I	10,12	10,10	31 2 22 2	10,17	0,12	9,12
	21.24	12,15	51.2,52.2	17.24.2	^,^ 6 0 3	8.8
	16.16			17,24.2	0,7.5	0,0
	5001	Idantifilar® Plue PowerPla	w® Eusion 6C Min	ifilor		
K3IVII F D	11 17 2	17 25		16 19	10.12	
1	10.12	12.15	12	16,17	R 10	0 1 2
I	10,12	13,13	31 2 32 2	11 12	0,12 X	10.11
	21.24	0 11	8 1 1	17.24.2	693	8
	16	7,11	0,11	17,27.2	0,7.0	0
	5901	GlobalFilor™				
RANDIE	11 17 3	17.25	11	16.18	10.12	
1	10.12	13 15	13	16,17	8.12	912
Į.	14 17	12 13	31 2 32 2	11 12	X	10 11
	21.24	.27.0	02/02.12	17.24.2	6.9.3	8
	16			,	-,	-
	- 5901	Investigator® 24plex QS				
Renze	11.17.3	17.25	11.11	16.18	10.12	
1	10.12	13.15	13.13	16.17	8.12	9.12
	14,17	12,13	31.2,32.2	11,12	X,X	10,11
	21,24			17,24.2	6,9.3	8,8
	16,16					
TMDEV4	- 5902	GlobalFiler™ Express				
	11,17.3	17,25	11,11	16,18	10,12	
1	10,12	13,15	13,13	16,17	8,12	9,12
	14,17	12,13	31.2,32.2	11,12	X,X	10,11
	21,24			17,24.2	6,9.3	8,8
	16,16					
TZBZVA	- 5901	Identifiler® Plus, PowerPle	ex® Fusion 6C, Min	iFiler™		
	11,17.3	17,25	11	16,18	10,12	
1	10,12	13,15	13	16,17	8,12	9,12
	14,17	12,13	31.2,32.2	11,12	X	10,11
	21,24	9,11	8,11	17,24.2	6,9.3	8
	16					

			TABLE	3		
WebCod	e - Test	Amplification Kits - Pr	obablistic Gen	otyping Software		
	D151656	- D251338	D25441	D351358	D55818	D651043
Item	D75820	D8S1179	D10S1248	D125391	D135317	D16S539
	D18S51	D195433	D21511	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	VWA	DYS391		DY5576	Y Indel	
	5002	GlobalFilor™ Evoraça, Truc	Item I - SIK	Kesults		
0110177 -	11 17 3	17.25		16 18	10.12	
1	10.12	13 15	13	16,17	8.12	912
·	14,17	12,13	31 2.32 2	11.12	XX	10.11
	21,24	/ . 0	0112/0212	17.24.2	6,9,3	8,8
	16,16			,		,
VFHZHA -	5902	GlobalFiler™- STRMix™ v2.	9.1			
	11,17.3	17,25	11	16,18	10,12	
1	10,12	13,15	13	16,17	8,12	9,12
	14,17	12,13	31.2,32.2	11,12	Х	10,11
	21,24			17,24.2	6,9.3	8
_	16					
VK9834 -	5902	PowerPlex® Fusion 6C- ST	RMix™ v2.10			
	11,17.3	17,25	11	16,18	10,12	
1	10,12	13,15	13	16,17	8,12	9,12
_	14,17	12,13	31.2,32.2	11,12	Х	10,11
	21,24	9,11	8,11	17,24.2	6,9.3	8
	16					
W2RLKZ -	5901	GlobalFiler™				
	11,17.3	17,25	11	16,18	10,12	
1	10,12	13,15	13	16,17	8,12	9,12
	14,17	12,13	31.2,32.2	11,12	X	10,11
	21,24	ND		17,24.2	6,9.3	8
	5001				INK	
WKGKTZ	- 3901		11	14 10	10.10	
1	10.12	17,25	13	16,18	8.12	0 1 2
	14 17	12.13	31 2 32 2	11 12	0,12 X	10.11
	21.24	12,10	01.2,02.2	17.24.2	6.9.3	8
	16				0,7.0	
WU8M28	- 5901	GlobalFiler™- STRMix™ 2.8	3			
	11,17.3	17.25	11	16,18	10,12	
1	10,12	13,15	13	16,17	8,12	9,12
	14,17	12,13	31.2,32.2	11,12	Х	10,11
	21,24			17,24.2	6,9.3	8
-	16					
X28843 -	5901	PowerPlex® 21- STRMix™ \	/2.8.0			
	11,17.3	17,25		16,18	10,12	11,11
1	10,12	13,15		16,17	8,12	9,12
_	14,17	12,13	31.2,32.2		X,X	10,11
	21,24	9,11	8,11		6,9.3	8,8
	16,16					

			TABLE	3		
WebCod	e - Test	Amplification Kits - P	robablistic Gene	otyping Software		
	D1S1656	D2S1338	D2S441	D3S1358	D5\$818	D6S1043
Item	D7S820	D8S1179	D10S1248	D12S391	D13S317	D16S539
	D18S51	D195433	D21S11	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	vWA	DY\$391	DY\$570	DY\$576	Y Indel	
			ltem 1 - STR	Results		
YGJ6L9 -	5901	GlobalFiler™- STRMix™				
	11,17.3	17,25	11	16,18	10,12	
1	10,12	13,15	13	16,17	8,12	9,12
	14,17	12,13	31.2,32.2	11,12	Х	10,11
	21,24			17,24.2	6,9.3	8
	16					
YJMUQ3	- 5901	PowerPlex® Fusion 6C				
	11,17.3	17,25	11,11	16,18	10,12	
1	10,12	13,15	13,13	16,17	8,12	9,12
	14,17	12,13	31.2,32.2	11,12	X,X	10,11
	21,24	9,11	8,11	17,24.2	6,9.3	8,8
	16,16	ND	ND	ND		
Z8LUQ4	- 5902	GlobalFiler™- TrueAllele®	VUler Release 202	2b		
	11,17.3	17,25	11,11	16,18	10,12	
1	10,12	13,15	13,13	16,17	8,12	9,12
	14,17	12,13	31.2,32.2	11,12	X,X	10,11
	21,24			17,24.2	6,9.3	8,8
	16,16					

TABLE 3							
WebCode - Test Amplification Kits - Probablistic Genotyping Software							
	D151656	D251338	D25441	D351358	D55818	D651043	
ltem	D75820	D8S1179	D1051248	D125391	D135317	D165539	
	D18S51	D19S433	D21S11	D22S1045	Amelogenin	CSF1PO	
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ	
	VWA	DYS391	DYS570	DYS576	Y Indel		
2000	2 5002	ClabelE:leu™ European Tra	Item 2 - SIR	Kesults			
ZQGMC	14 17 2			edse 2022b	11 12		
2	0 11	17,10	14,14	18,10	12.12	014	
Z	15 19	12 15 2	33 33 2	14.15	X Y	10.12	
	20.23	12,13.2	33,33.2	18 28 2	7 7	11.14	
	15.17	10		10,20.2	2	11,17	
346383	5902	GlobalFiler™			L		
040000	16 17 3	17 18	14	16	11 13		
2	9 11	10.13	13	18 19	12	914	
2	15.19	12,15,2	33.33.2	14.15	X.Y	10.12	
	20,23	,	00,00.2	18.28.2	7	11,14	
	15,17	10		,	2	,	
37.12CU	- 5901	GlobalFiler™- STRMix™					
0,0200	16,17.3	17,18	14,14	16,16	11,13		
2	9,11	10,13	13,13	18,19	12,12	9,14	
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12	
	20,23			18,28.2	7,7	11,14	
	15,17	10			2		
3MQZU1	(- 5902	Identifiler® Plus					
		17,18		16,16	11,13		
2	9,11	10,13			12,12	9,14	
	15,19	12,15.2	33,33.2		X,Y	10,12	
	20,23				7,7	11,14	
	15,17						
3WHFJ3 - 5902		GlobalFiler™- STRMix™ V	2.9.1				
	16,17.3	17,18	14	16	11,13		
2	9,11	10,13	13	18,19	12	9,14	
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12	
	20,23			18,28.2	7	11,14	
	15,17	10			2		
44PDRD	- 5901	GlobalFiler™					
	16,17.3	17,18	14	16	11,13		
2	9,11	10,13	13	18,19	12	9,14	
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12	
	20,23			18,28.2	7	11,14	
	15,17	10			2		
4BEUJW	- 5902	PowerPlex® 21- STRMix™	2.10.0				
	16,17.3	17,18		16,16	11,13	12,13	
2	9,11	10,13		18,19	12,12	9,14	
	15,19	12,15.2	33,33.2		Х,Ү	10,12	
	20,23	9,12	5,10		7,7	11,14	
	15,17						

TABLE 3						
WebCo	de - Test	Amplification Kits - P	robablistic Gen	otyping Software		
	D151656	D251338	D25441	D351358	D55818	D651043
ltem	D75820	D8S1179	D1051248	D125391	D135317	D165539
	D18S51	D19S433	D21511	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	vWA	DYS391	DYS570	DYS576	Y Indel	
	5000		Item 2 - STR	Results		
4FBJY4 -	5902	PowerPlex® Fusion 6C - L	INAxs			
0	16,17.3	17,18	14,14	16,16	11,13	0.1.4
2	9,11	10,13	13,13	18,19	12,12	9,14
	15,19	12,15.2	33,33.2	14,15	Χ,Υ	10,12
	20,23	9,12	5,10	18,28.2	/ ,/	11,14
	15,17		20,20	17,17		
4UQG64	4 - 5901	GlobalFiler . STRMix .				
	16,17.3	17,18	14	16	11,13	
2	9,11	10,13	13	18,19	12	9,14
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12
	20,23	10		18,28.2	/	11,14
	15,17				2	
4XQRZ4	- 5901	GlobalFiler . STRMix .				
	16,17.3	17,18	14	16	11,13	
2	9,11	10,13	13	18,19	12	9,14
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12
	20,23	10		18,28.2	/	11,14
	15,17				2	
62K9Q4	- 5901	GlobalFiler . STRMix .	- /			
	16,17.3	17,18	14	16	11,13	0.1.4
2	9,11	10,13	13	18,19	12	9,14
	15,19	12,15.2	33,33.2	14,15	Χ,Υ	10,12
	20,23	10		18,28.2	/	11,14
(0) (0.00	15,17				Z	
63VBC2	- 5901	GlobalFiler		1.4	11.10	
0	16,17.3	17,18	14	16	11,13	0.1.4
2	9,11	10,13	13	18,19	12	9,14
	15,19	12,15.2	33,33.2	14,15	λ, ř	10,12
	20,23	10		18,28.2	7	11,14
	15,17				2	
6P1H6Y	- 5902	GlobalFiler - TrueAllele®	VUler Release 202	(2b	11.10	
0	16,17.3	17,18	14,14	16,16	11,13	0.1.4
2	9,11	10,13	13,13	18,19	12,12	9,14
	15,19	12,15.2	33,33.2	14,15	۸, i ד ד	10,12
	20,23	10		10,20.2	/,/	11,14
	5001				Z	
//A6KY	- 3701	GiobalFiler	14.14	1 / 1 /	11.10	
0	16,17.3	1/,18	14,14	16,16	11,13	0.1.4
2	9,11	10,13		18,19	12,12	9,14
	10,19	12,13.2	<u>کی,کک</u>	14,10	Λ, Ϊ	10,12
	15 17	10		10,20.2	· · · · · · · · · · · · · · · · · · ·	11,14
	10,17	10			<u> </u>	

TABLE 3								
WebCode - Test Amplification Kits - Probablistic Genotyping Software								
	D151656	D251338	D25441	D3\$1358	D5\$818	D651043		
ltem	D75820	D8S1179	D10S1248	D12S391	D135317	D165539		
	D18S51	D19S433	D21S11	D22S1045	Amelogenin	CSF1PO		
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ		
	vWA	DYS391	DYS570	DYS576	Y Indel			
	5001		$\frac{11}{2}$	Results				
700INUZ	- 5901		VIIX	14 14	11.12			
2	0 1 1	1/,10		10,10	12.12	014		
Z	9,11	10,150	33 33 0		12,12	9,14		
	20.23	12,13.2	55,55.2		7.7	10,12		
	15.17				/ ,/	11,14		
7H/IRV	5002	GlobalEilar™ TruoAllala®	VIIIar Pologeo 202	20h				
/ 1 1 4 JI(v -	16 17 3			16	11 13			
2	9 11	10.13	13 13	18 19	12 12	914		
Z	15 19	12 15 2	33 33 2	14 15	X Y	10.12		
	20.23	12,10.2	00,00.2	18.28.2	7.7	11.14		
	15,17	10		,	2	,		
	- 5901	GlobalFiler™- STRMix™ 2	8					
000,000	16.17.3	17.18	14	16	11.13			
2	9,11	10,13	13	18,19	12	9,14		
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12		
	20,23			18,28.2	7	11,14		
	15,17	10		,	2	,		
8Y8DYV -	- 5902	GlobalFiler™- TrueAllele®	VUIer Release 202	22b				
	16,17.3	17,18	14,14	16,16	11,13			
2	9,11	10,13	13,13	18,19	12,12	9,14		
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12		
	20,23			18,28.2	7,7	11,14		
	15,17	10			2			
8YUL9Z -	5901	PowerPlex® Fusion 6C, NGM Detect - LRmix						
	16,17.3	17,18	14,14	16,16	11,13			
2	9,11	10,13	13,13	18,19	12,12	9,14		
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12		
	20,23	9,12	5,10	18,28.2	7,7	11,14		
	15,17	10	20	17	2			
9CLYK9 - 5901		GlobalFiler™- STRMix™						
	16,17.3	17,18	14	16	11,13			
2	9,11	10,13	13	18,19	12	9,14		
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12		
	20,23			18,28.2	7	11,14		
	15,17	10			2			
9MBR8Q	- 5901	PowerPlex® 21						
	16,17.3	17,18		16,16	11,13	12,13		
2	9,11	10,13		18,19	12,12	9,14		
	15,19	12,15.2	33,33.2		X,Y	10,12		
	20,23	9,12	5,10		7,7	11,14		
	15,17							

TABLE 3							
WebCode - Test Amplification Kits - Probablistic Genotyping Software							
	D151656	D251338	D25441	D351358	D55818	D651043	
Item	D75820	D8S1179	D10S1248	D125391	D135317	D16S539	
	D18S51	D195433	D21511	D22S1045	Amelogenin	CSF1PO	
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ	
	VWA	D12391		D15576	t Indel		
	5002	PowerPlay® 21	Item 2 - STR	Results			
7 VVLIIK -	16 17 3	17 18		16 16	11 13	12.13	
2	9 11	10.13		18.19	12 12	9 14	
2	15 19	12 15 2	33 33 2	10,17	X Y	10.12	
	20.23	9.12	5.10		7.7	11,14	
	15.17	,,	0,10		. ,.	,	
B7W38V	. 5901	GlobalEiler™					
2,11001	16.17.3	17.18	14.14	16.16	11.13		
2	9.11	10.13	13.13	18.19	12.12	9.14	
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12	
	20,23			18,28.2	7,7	11,14	
	15,17	10			2		
BDR7Z7 -	5901	PowerPlex® Fusion 6C, I	ElexPlex 27				
	16,17.3	17,18	14	16	11,13	12,13	
2	9,11	10,13	13	18,19	12	9,14	
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12	
	20,23	9,12	5,10	18,28.2	7	11,14	
	15,17	10	20	17			
BKHHBU - 5901		GlobalFiler™					
	16,17.3	17,18	14,14	16,16	11,13		
2	9,11	10,13	13,13	18,19	12,12	9,14	
-	15,19	12,15.2	33,33.2	14,15	X,Y	10,12	
	20,23			18,28.2	7,7	11,14	
	15,17	10			2		
BNGT8U - 5902		GlobalFiler™					
	16,17.3	17,18	14	16	11,13		
2	9,11	10,13	13	18,19	12	9,14	
_	15,19	12,15.2	33,33.2	14,15	X,Y	10,12	
	20,23			18,28.2	7	11,14	
	15,17	10			2		
CDTG9M	- 5901	GlobalFiler™					
_	16,17.3	17,18	14,14	16,16	11,13	-	
2	9,11	10,13	13,13	18,19	12,12	9,14	
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12	
	20,23	-	-	18,28.2	7,7	11,14	
	15,17	10	-	-	2		
CKECAQ	- 5902	GlobalFiler™- TrueAllele	® VUler Release 202	22b			
_	16,17.3	17,18	14,14	16,16	11,13		
2	9,11	10,13	13,13	18,19	12,12	9,14	
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12	
	20,23			18,28.2	7,7	11,14	
	15,17	10			2		

Printed: May 12, 2025
			TABLE	3		
WebCode	e - Test	Amplification Kits -	Probablistic Gene	otyping Software		
	D1\$1656	- D251338	D25441	D351358	D55818	D651043
Item	D75820	D8\$1179	D1051248	D125391	D135317	D165539
	D18S51	D195433	D21S11	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	vWA	DYS391	DYS570	DYS576	Y Indel	
			ltem 2 - STR	Results		
CRBAWU -	- 5901	PowerPlex® ESI-17 Fast				
	16,17.3	17,18	14,14	16,16		
2		10,13	13,13	18,19		9,14
	15,19	12,15.2	33,33.2	14,15	X,Y	
	20,23			18,28.2	7,7	
	15,17					
CRVQCW	- 5901	GlobalFiler™				
	16,17.3	17,18	14	16	11,13	
2	9,11	10,13	13	18,19	12	9,14
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12
	20,23			18,28.2	7	11,14
	15,17	10			2	
CXYRCQ -	5902	GlobalFiler™				
	16,17.3	17,18	14,14	16,16	11,13	
2	9,11	10,13	13,13	18,19	12,12	9,14
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12
	20,23			18,28.2	7,7	11,14
	15,17	10			2	
CY6DP6 -	5901	GlobalFiler™				
_	16,17.3	17,18	14	16	11,13	
2	9,11	10,13	13	18,19	12	9,14
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12
	20,23			18,28.2	7	11,14
	15,17	10			2	
CZ6EPV - S	5901	GlobalFiler™				
_	16,17.3	17,18	14	16	11,13	
2	9,11	10,13	13	18,19	12	9,14
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12
	20,23			18,28.2	7	11,14
	15,17	10			2	
E34ANN -	5902	GlobalFiler™ Express- Tru	⊎eAllele® VUler Rele	ase 2022b		
	16,17.3	17,18	14,14	16,16	11,13	
2	9,11	10,13	13,13	18,19	12,12	9,14
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12
	20,23			18,28.2	7,7	11,14
	15,17	10			2	
EHXZJK - 5	5902	PowerPlex [®] 21- STRMix [™]	2.10			
_	16,17.3	17,18		16,16	11,13	12,13
2	9,11	10,13		18,19	12,12	9,14
_	15,19	12,15.2	33,33.2		X,Y	10,12
	20,23	9,12	5,10		7,7	11,14
	15,17					

			TABLE	3		
WebCod	de - Test	Amplification Kits -	Probablistic Gen	otyping Software		
	D151656	D251338	D25441	D3\$1358	D5\$818	D651043
Item	D75820	D8S1179	D10S1248	D125391	D135317	D165539
	D18S51	D19S433	D21S11	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	vWA	DYS391	DYS570	DYS576	Y Indel	
			Item 2 - SIR	Results		
ELJJPP - 5	902	GlobalFiler - TrueAllele	® VUler Kelease 202	22b	11.10	
0	16,17.3	17,18	14,14	16,16	11,13	014
Z	9,11	10,13	13,13	18,19	12,12	9,14
	15,19	12,15.2	33,33.Z	14,15	λ, Ϊ	10,12
	20,23	10		10,20.2	2	11,14
	5000			00001	Z	
ETOTIN -	- 5902	GlobalFiler Express- Ir		ase 2022b	11.10	
0	16,17.3	17,18	14,14	16,16	11,13	0.1.4
Z	9,11	10,13	13,13	18,19	12,12	9,14
	15,19	12,15.2	33,33.2	14,15	λ, f	10,12
	15.17	10		10,20.2	2	11,14
	5000				Ζ	
F/HOUP	- 5902		VZ.9.1	17	11.10	
0	10,17.3	17,10	14	10 10	11,13	0.14
Z	9,11	10,15	22.22.0	10,19		9,14
	15,19	12,13.2	33,33.2	14,15	7	10,12
	15.17	10		10,20.2	2	11,14
	5002	ClabalEilar™ True Allala	VIIIar Palaraa 200	206	Z	
TONQUIN	1- 3702	17 18		16.16	11 13	
2	0.11	17,10	14,14	10,10	10.10	014
Z	15 19	12 15 2	33 33 2	14 15	X Y	10.12
	20.23	12,10.2	00,00.2	18 28 2	7 7	11 14
	15,17	10		10,20.2	2	
EPXE61	5902	Identifiler® Plus				
TIMEOL	0702	17 18		1616	11 13	
2	911	10.13		10,10	12 12	914
_	15,19	12,15.2	33.33.2		X.Y	10,12
	20,23				7.7	11,14
	15,17				,	,
GCH7HK	(- 5902	GlobalFiler™- TrueAllele	® VUler Release 202	22b		
	16.17.3	17.18	14.14	16.16	11.13	
2	9,11	10,13	13,13	18,19	12,12	9,14
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12
	20,23			18,28.2	7,7	11,14
	15,17	10			2	
H484W.J	- 5902	GlobalFiler™- TrueAllele	® VUler Release 202	22b		
	16,17.3	17,18	14,14	16,16	11,13	
2	9,11	10,13	13,13	18,19	12,12	9,14
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12
	20,23			18,28.2	7,7	11,14
	15,17	10			2	

			TABLE	3		
WebCoc	le - Test	Amplification Kits -	Probablistic Gen	otyping Software		
	D151656	- D251338	D25441	D351358	D55818	D651043
Item	D75820	D251350	D10S1248	D125391	D135317	D165539
	D18S51	D195433	D21S11	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	vWA	DYS391	DY\$570	DYS576	Y Indel	
	5000		Item 2 - SIR	Results		
heQhil	- 5902		VUIEr Kelease 202	2D	11.10	
0	10,17.3	17,18	14,14	10,10	10.10	0.14
Z	7,11	10,15	33 33 0	10,19	12,12	9,14
	20.23	12,13.2	33,33.2	18.28.2	7.7	10,12
	15.17	10		10,20.2	2	11,14
	5001	GlobalFilar™ IOC			Z	
	16 17 3	17.18	1414	16 16	11 13	
2	9 11	10.13	13.13	18.19	12.12	914
Z	15 19	12 15 2	33 33 2	14 15	X Y	10.12
	20.23	12,10.2	00,00.2	18.28.2	7.7	11,14
	15,17	10		/	2	,
137QNH	- 5902	PowerPlex® 21- STRMix™	×2100			
	16,17.3	17,18	.2	16,16	11,13	12,13
2	9,11	10,13		18,19	12,12	9,14
	15,19	12,15.2	33,33.2	, •	, Х,Ү	10,12
	20,23	9,12	5,10		7,7	11,14
	15,17					
J772JG -	5902	PowerPlex® 21- STRMix™	[™] 2.10			
	16,17.3	17,18		16,16	11,13	12,13
2	9,11	10,13		18,19	12,12	9,14
	15,19	12,15.2	33,33.2		X,Y	10,12
	20,23	9,12	5,10		7,7	11,14
	15,17					
JHLQHM	- 5902	GlobalFiler™ - DNAxs				
	16,17.3	17,18	14,14	16,16	11,13	
2	9,11	10,13	13,13	18,19	12,12	9,14
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12
	20,23			18,28.2	7,7	11,14
	15,17	10			2	
JPGUBX -	5901	GlobalFiler™				
	16,17.3	17,18	14	16	11,13	
2	9,11	10,13	13	18,19	12	9,14
	15,19	12,15.2	33,33.2	14,15	Х,Ү	10,12
	20,23			18,28.2	7	11,14
	15,17	10			2	
K9ECCE	- 5902	PowerPlex® 21				
	16,17.3	17,18		16,16	11,13	12,13
2	9,11	10,13	-	18,19	12,12	9,14
	15,19	12,15.2	33,33.2		X,Y	10,12
	20,23	9,12	5,10		/,/	11,14
	13.17					

			TABLE	3		
WebCod	e - Test	Amplification Kits - P	robablistic Gene	otyping Software		
	D1\$1656	D251338	D25441	D351358	D55818	D651043
ltem	D75820	D851179	D1051248	D125391	D135317	D165539
	D18S51	D195433	D21S11	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	vWA	DYS391	DYS570	DYS576	Y Indel	
			Item 2 - STR	Results		
KPDQAK	- 5902	GlobalFiler - STRMix v	2.9.1	- /		
0	16,17.3	17,18	14	16	11,13	0.1.4
2	9,11	10,13	13	18,19	12	9,14
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12
	20,23	10		18,28.2	/	11,14
	10,17				Z	
L6/JVL - 3	5901	GlobalFiler *** - STRMix ***	- /	- /		
0	16,17.3	17,18	14	16	11,13	0.1.4
2	9,11	10,13	13	18,19	12	9,14
	15,19	12,15.2	33,33.2	14,15	Χ,Υ	10,12
	20,23	10		18,28.2	7	11,14
	10,17				Z	
L8XEJE - 5	1/170	PowerPlex® 21		1 / 1 /	11.10	10.10
0	16,17.3	17,18		16,16	11,13	12,13
2	9,11	10,13	22.22.0	18,19	12,12	9,14
	15,19	12,15.2	5.10		λ, ř	10,12
	20,23	9,12	5,10		/ ,/	11,14
	15,17	LL víti e D				
LBAKVF -	5902	Identifiler® Plus		1 / 1 /	11.10	
0	0.11	17,18		16,16	11,13	0.1.4
Z	9,11	10,13	22.22.0		12,12	9,14
	15,19	12,13.2	33,33.2		7,1	10,12
	15 17				/ ,/	11,14
	5001	David and Diavid ESV17 Earst St		5 1 1		
LGSPINJ -	14172	rowerriex (ESAI / rast S)	14.14	14.14		
2	10,17.3	17,10	14,14	10,10		0.14
Z	15 19	12 15 2	33 33 2	14,15	XY	7,14
	20.23	12,13.2	00,00.2	18 28 2	7 7	
	15.17			10,20.2	, , , , , , , , , , , , , , , , , , ,	
	5902	GlobalFiler™ TrueAllele®	VI IIer Release 202	22h		
LITOROFT	16 17 3	17.18	14 14	16 16	11 13	
2	9 11	10.13	13.13	18.19	12.12	9.14
~	15.19	12.15.2	33,33.2	14.15	X.Y	10,12
	20.23			18.28.2	7.7	11.14
	15.17	10			2	
UPUWF	- 5901	Identifiler® Plus PowerPla	x® Eusion 6C Mini	filer		
_0.000	16.17.3	17.18	14	16	11.13	
2	9.11	10.13	13	18.19	12	9.14
-	15.19	12.15.2	33,33.2	14.15	X.Y	10,12
	20,23	9,12	5,10	18,28.2	7	11,14
	15,17	10	20	17		

			TABLE	3		
WebCod	de - Test	Amplification Kits - P	robablistic Gen	otyping Software		
	D1S1656	D2\$1338	D2S441	D3\$1358	D5\$818	D6S1043
ltem	D75820	D8S1179	D1051248	D12S391	D13S317	D16S539
	D18S51	D195433	D21511	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	VWA	D12241		DIS570	T Indei	
IYKQII -	5901	GlobalFiler™- STRMix™		Results		
21110002	16,17.3	17,18	14	16	11,13	
2	9,11	10,13	13	18,19	12	9,14
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12
	20,23			18,28.2	7	11,14
	15,17	10			2	
MW99ZF	- 5902	GlobalFiler™ Express- True	eAllele® VUler Rele	ase 2022b		
	16,17.3	17,18	14,14	16,16	11,13	
2	9,11	10,13	13	18,19	12,12	9,14
	15,19	12,15.2	33,33.2	14,15	Υ	10,12
	20,23			18,28.2	7,7	11,14
	15,17	10			2	
NWNRKE) - 5901	Identifiler® Plus, PowerPle	x® Fusion 6C, Min	iFiler		
	16,17.3	17,18	14	16	11,13	
2	9,11	10,13	13	18,19	12	9,14
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12
	20,23	9,12	5,10	18,28.2	7	11,14
	15,17		20	17		
P3QH3A	- 5901	PowerPlex® 21- STRMix™	2.8	1 / 1 /	11.10	10.10
0	16,17.3	17,18		16,16	11,13	12,13
Z	9,11	10,15	33 33 0	10,19	12,12	9,14
	20.23	0 12	5 10		7 7	10,12
	15.17	//12	0,10		, , , , , , , , , , , , , , , , , , ,	11/17
P8KFPF -	5902	GlobalFiler™- STRMix™				
I OKEI I	16.17.3	17.18	14	16	11.13	
2	9,11	10,13	13	18,19	12	9.14
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12
	20,23			18,28.2	7	11,14
	15,17	10			2	
P8TP7Q	- 5901	GlobalFiler™				
	16,17.3	17,18	14	16	11,13	
2	9,11	10,13	13	18,19	12	9,14
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12
	20,23			18,28.2	7	11,14
	15,17	10			2	
PD88ZF -	5902	GlobalFiler™- STRMix™ v2	2.8			
	16,17.3	17,18	14	16	11,13	
2	9,11	10,13	13	18,19	12	9,14
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12
	20,23			18,28.2	7	11,14
	15,17	10			2	

			TABLE	3		
WebCoo	de - Test	Amplification Kits - F	Probablistic Gene	otyping Software		
	D151656	D251338	D25441	D3\$1358	D5\$818	D651043
ltem	D75820	D8S1179	D10S1248	D12S391	D135317	D16S539
	D18S51	D195433	D21511	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	vWA	DYS391	DYS570	DYS576	Y Indel	
	5001		Item 2 - STR	Results		
QAARGE	- 5901		1414	1 / 1 /	11.10	
0	16,17.3	17,18	14,14	16,16	11,13	0.14
Z	9,11	10,13	13,13	18,19	12,12	9,14
	15,19	12,15.2	33,33.2	14,15	λ, Ϊ	10,12
	20,23	10		10,20.2	/,/	11,14
	15,17			() I	2	
K3MFPB	- 5901	Identifiler® Plus, PowerPle	x® Fusion 6C, Mini	filer	11.10	
0	16,17.3	17,18	14	16	11,13	0.1.4
2	9,11	10,13	13	18,19	12	9,14
	15,19	12,15.2	5.10	14,15	λ, Ϊ	10,12
	20,23	9,12	5,10	10,20.2	/	11,14
	15,17		20	17		
R9RDYE -	- 5901	GlobalFiler'			11.10	
2	16,17.3	17,18	14	16	11,13	0.1.4
	9,11	10,13	13	18,19	12	9,14
	15,19	12,15.2	33,33.2	14,15	Χ,Υ	10,12
	20,23	10		18,28.2	/	11,14
	15,17				Z	
RCPXZD	- 5901	Investigator® 24plex QS				
	16,17.3	17,18	4, 4	16,16	11,13	0.1.4
2	9,11	10,13	13,13	18,19	12,12	9,14
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12
	20,23	10		18,28.2	/,/	11,14
	15,17					
IMDEV4	- 5902	GlobalFiler [™] Express				
	16,17.3	17,18	4, 4	16,16	11,13	0.1.4
2	9,11	10,13	13,13	18,19	12,12	9,14
	15,19	12,15.2	33,33.2	14,15	Χ,Υ	10,12
	20,23	10		18,28.2	/,/	11,14
	15,17				2	
IZBZVA -	- 5901	Identifiler® Plus, PowerPle	x® Fusion 6C, Mini	Filer'''		
0	16,17.3	17,18	14	16	11,13	0.1.4
2	9,11	10,13	13	18,19	12	9,14
	15,19	12,15.2	53,33.2	14,15	λ, ř	10,12
	20,23	9,12	5,10	18,28.2		11,14
	15,17			17		
UNVK99	- 5902	GiobalFiler'" - TrueAllele®	VUIer Kelease 202	(Zb	11.10	
0	16,17.3	17,18	14,14	16,16	11,13	0.14
2	9,11	10,13	13,13	18,19	12,12	9,14
	15,19	12,15.2	33,33.2	14,15	Χ,Υ	10,12
	20,23	10		18,28.2	/,/	11,14
	10,17	10			Z	

			TABLE	3		
WebCo	de - Test	Amplification Kits - P	robablistic Gen	otyping Software		
	D151656	- D251338	D25441	D351358	D55818	D651043
ltem	D75820	D851179	D1051248	D125391	D135317	D165539
	D18S51	D19S433	D21511	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	vWA	DYS391	DYS570	DYS576	Y Indel	
	5000		Item 2 - STR	Results		
VFHZHA	- 5902	GlobalFiler - STRMix v2				
0	16,17.3	1/,18	14	16	11,13	0.1.4
2	9,11	10,13	13	18,19	12	9,14
	15,19	12,15.2	33,33.2	14,15	Χ, Υ	10,12
	20,23	10		18,28.2	7	11,14
	15,17				Z	
VK9834	- 5902	PowerPlex® Fusion 6C- 51	RMIX *** v2.10		11.10	
0	16,17.3	17,18	14	16	11,13	0.1.4
2	9,11	10,13	13	18,19	12	9,14
	15,19	12,15.2	53,33.2	14,15	λ, ř	10,12
	20,23	9,12	3,10	10,20.2	/	11,14
	5001		20	17		
VVZKLKZ	- 5901	GlobalFiler	1.4	17	11.10	
0	10,17.3	17,10	14	10 10	11,13	0.14
Z	9,11	10,15	22.22.0	10,19	12	9,14
	20.23	12,13.2	33,33.2	14,15	~,1	10,12
	15.17	10		10,20.2	2	11,14
	7 5001	GlabalFilar™			L	
WKGKTZ	16 17 3	17 18	14	16	11 13	
2	0 1 1	10.13	14	18 10	12	01/
2	15.19	12 15 2	33 33 2	14 15	XY	10.12
	20.23	12,13.2	00,00.2	18 28 2	7	11 14
	15.17	10		10,20.2	2	,
WI 18M2	8 - 5901	GlobalFiler™- STRMix™ 2	8			
110011121	16.17.3	17.18	14	16	11.13	
2	9.11	10.13	13	18.19	12	9.14
_	15,19	12,15.2	33,33.2	14,15	X.Y	10,12
	20,23	·	,	18,28.2	7	11,14
	15,17	10			2	
X28843	- 5901	PowerPlex® 21- STRMix™	V2.8.0			
	16,17.3	17,18		16,16	11,13	12,13
2	9,11	10,13		18,19	12,12	9,14
	15,19	12,15.2	33,33.2		X,Y	10,12
	20,23	9,12	5,10		7,7	11,14
	15,17					
YGJ6L9	- 5901	GlobalFiler™- STRMix™				
-	16,17.3	17,18	14	16	11,13	
2	9,11	10,13	13	18,19	12	9,14
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12
	20,23			18,28.2	7	11,14
	15,17	10			2	

			TABLE	3						
WebCoo	WebCode - Test Amplification Kits - Probablistic Genotyping Software									
ltem	D1S1656	D251338	D2S441	D3S1358	D55818	D6S1043				
	D18551	D195433	D21511	D22S1045	Amelogenin	CSF1PO				
	FGA vWA	Penta D DYS391	Penta E DYS570	SE33 DYS576	TH01 Y Indel	ΤΡΟΧ				
	ltem 2 - STR Results									
YJMUQ3	- 5901	PowerPlex® Fusion 6C								
	16,17.3	17,18	14,14	16,16	11,13					
2	9,11	10,13	13,13	18,19	12,12	9,14				
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12				
	20,23	9,12	5,10	18,28.2	7,7	11,14				
	15,17	10	20	17						
Z8LUQ4	- 5902	GlobalFiler™- TrueAllele®	VUler Release 202	2b						
	16,17.3	17,18	14,14	16,16	11,13					
2	9,11	10,13	13,13	18,19	12,12	9,14				
	15,19	12,15.2	33,33.2	14,15	X,Y	10,12				
	20,23			18,28.2	7,7	11,14				
	15,17	10			2					

			TABLE	3					
WebCo	WebCode - Test Amplification Kits - Probablistic Genotyping Software								
	D1S1656	D2S1338	D2S441	D3S1358	D5S818	D6S1043			
ltem	D7\$820	D8\$1179	D10S1248	D12S391	D13S317	D16S539			
	D18S51	D195433	D21S11	D22S1045	Amelogenin	CSF1PO			
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ			
	vWA	DY\$391	DY\$570	DYS576	Y Indel				
			ltem 3 - STR	Results					
8YUL9Z -	- 5901 Powe	erPlex® Fusion 6C,	NGM Detect - LRmix						
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,10,11,12,13				
3	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14			
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 .2	11,12,14,15	X,Y	10,11,12			
	20,21,23,23.2,24	9,10,11,12,13	5,8,10,11,12	15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14			
	15,16,17	10	18,20	17,19	2				
LG3PNJ	- 5901 Powe	erPlex® ESX17 Fast	System- STRMix™ V2.	5.11					
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18					
3		10,13,15	13,14	16,17,18,19,21		8,9,12,13,14			
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 .2	11,12,14,15	X,Y				
	20,21,23,23.2,24			15,17,18,24.2,28.2	6,7,9,9.3				
	15,16,17								

			TABLE	3		
WebCo	de - Test 🛛 🗛 An	nplification Kits -	Probablistic Gene	otyping Software		
	D1S1656	D2\$1338	D2S441	D3S1358	D5\$818	D6S1043
ltem	D75820	D8S1179	D1051248	D125391	D135317	D16S539
	D18551	D195433	D21511	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	трох
	VWA	D13391		D15576	r indei	
2QGMC	2 - 5902 Glol	balFiler™- TrueAllele	® "VUler Release 20	22b		
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,10,11,13	
3e	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,33,33.2	11,12,14,15	X,Y	10,11,12
	20,21,23,23.2,24			15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
34G3B3	- 5902 Glol	balFiler™- STRMix™	2.8			
	11,13,14,15,16,16.3 ,17.3,18.3	16,17,18,19,20,24, 25	10,11,13,14,15	14,15,16,17,18	8,9,10,11,12,13	
3e	8,9,10,11,12	9,10,12,13,14,15	12,13,14	16,17,18,19,20,21	8,10,11,12	7,8,9,11,12,13,14
	14,15,16,17,18,19	11,12,13,14.2,15.2	29,30,30.2,31.2,32 ,32.2,33,33.2	11,12,13,14,15,16	X,Y	9,10,11,12
	19,20,21,22,22.2,2 3,23.2,24			14,14.2,15,16,16.2 ,17,17.2,18,23.2,2 4.2,27.2,28,28.2,2	5,6,7,8,9,9.3	8,10,11,13,14
	14,15,16,17	9,10		/.2	2	
37J2CU	- 5901 Glol	balFiler™- STRMix™				
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,10,11,12,13	
3e	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32,32.2,33 ,33.2	11,12,14,15	X,Y	10,11,12
	20,21,23,23.2,24			15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
3MQZU	Y - 5902 Iden	tifiler® Plus- STRMix	тм			
		17,18,20,25		15,16,18	9,10,11,12,13	
3e	9,10,11,12	10,13,15			8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33		X,Y	10,11,12
	20,21,23,23.2,24		.2		6,7,9,9.3	8,10,11,14
	15,16,17					
3WHFJ3	- 5902 Glol	balFiler™- STRMix™ `	√2.9.1			
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,10,11,12,13	
3e	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 .2	11,12,14,15	X,Y	10,11,12
	20,21,23,23.2,24			15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
44PDRD	- 5901 Glol	balFiler™- STRMix™				
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,10,11,12,13	
3e	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 .2	11,12,14,15	Х,Ү	10,11,12
	20,21,23,23.2,24			15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	

			TABLE	3		
WebCo	de - Test 🛛 An	nplification Kits -	Probablistic Gene	otyping Software		
ltom	D1S1656	D251338	D2S441	D3S1358	D5\$818	D6S1043
nem	D73620	D195433	D1031248	D125391 D22\$1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	TPOX
	vWA	DYS391	DY\$570	DYS576	Y Indel	
			ltem 3e - STR	Results		
4BEUJW	- 5902 Pow	erPlex® 21- STRMix	™ 2.10.0			
	11,14,16,17.3,18.3	17,18,25		15,16,18	10,11,12,13	12,13
3e	9,11	10,13,15		16,18,19	11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33		X,Y	10,12
	20,24	9,10,11,12,13	5,10		6,7,9.3	8,11,14
	15,16,17					
4FBJY4 -	5902 Pow	erPlex® Fusion 6C -	DNAxs			
	11	17,18,20,25	14,16,17.3,18.3	15,16,18	9,10,11,12,13	
3e	9,10,11,12	10,13,15	11,14	16,17,18,19,21	13,14	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33	11,12,14,15	X,Y	10,11,12
	20.21.23.23.2.24	9.10.11.12.13	.2	15,17,18,24,2,28,2	6,7,9,9.3	8,10,11,14
	15,16,17	10	18.20	17,19	-/. /. /	-//.
4UQG64	4 - 5901 Glo	balFiler™- STRMix™	,			
	11,14,15,16,17.3,18	17,18,20,24,25	11,14	15,16,18	9,10,11,12,13	
3e	9,10,11,12	10,12,13,15	12,13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33	11,12,14,15	X,Y	10,11,12
	20,21,23,23.2,24		.۲	15,17,18,24.2,27.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
4XQRZ4	- 5901 Glo	balFiler™- STRMix™				
	11,13,14,15,16,16.3 ,17.3,18.3	16,17,18,19,20,24, 25	10,11,13,14	14,15,16,17,18	9,10,11,12,13	
3e	9,10,11,12	10,12,13,14,15	12,13,14	16,17,18,19,20,21	8,10,11,12	8,9,12,13,14
	14,15,16,17,18,19	11,12,13,14.2,15.2	30,31.2,32.2,33,33	11,12,13,14,15,16	X,Y	10,11,12
	19,20,21,22,22.2,2 3,23.2,24		.2	14,15,16,17,18,24. 2,27.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
62K9Q4	- 5901 Glo	balFiler™- STRMix™				
	11,13,14,15,16,16.3 ,17.3,18.3	16,17,18,19,20,24, 25	10,11,13,14	14,15,16,17,18	9,10,11,12,13	
3e	9,10,11,12	10,12,13,14,15	12,13,14	16,17,18,19,21	8,10,11,12	8,9,12,13,14
	14,15,16,17,18,19	11,12,13,14.2,15.2	30,30.2,31.2,32.2, 33,33.2	11,12,13,14,15,16	X,Y	9,10,11,12
	19,20,21,23,23.2,2 4			14,15,16,17,18,24. 2,27.2,28.2	6,7,9,9.3	8,10,11,14
	15,16.17	10			2	

			TABLE	3		
WebCo	de - Test 🛛 Am	plification Kits -	Probablistic Gene	otyping Software		
	D1S1656	D2S1338	D2\$441	D3S1358	D5S818	D6S1043
ltem	D7\$820	D8S1179	D10S1248	D12S391	D135317	D16S539
	D18\$51	D195433	D21511	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	VWA	D12241		Pooulto	rindei	
63VBC2	- 5901 Glob	oalFiler™- STRMix™	liem Se - STK	Results		
	11,13,14,15,16,16.3 ,17.3,18.3	16,17,18,19,20,24, 25	10,11,13,14	15,16,17,18	9,10,11,12,13	
3e	9,10,11,12	10,12,13,14,15	12,13,14	16,17,18,19,21	8,10,11,12	8,9,12,13,14
	14,15,16,17,18,19	11,12,13,15.2	30,30.2,31.2,32.2, 33,33.2	11,12,13,14,15,16	X,Y	9,10,11,12
	20,21,22.2,23,23.2 ,24			14,15,16,17,18,24. 2,27.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
6PTH6Y	- 5902 Glob	oalFiler™- TrueAllele	® VUler Release 202	2b		
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,10,11,12,13	
3e	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 .2	11,12,14,15	X,Y	10,11,12
	20,21,23,23.2,24			15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
77A6RY	- 5901 Glob	oalFiler™- STRMix™				
	11,13,14,15,16,16.3 ,17.3,18.3	16,17,18,19,20,24, 25	10,11,13,14	15,16,17,18	9,10,11,12,13	
3e	8,9,10,11,12	9,10,12,13,14,15	12,13,14	16,17,18,19,20,21	8,10,11,12	8,9,12,13,14
	14,15,16,17,18,19	11,12,13,14.2,15.2	30,30.2,31.2,32,32 .2,33,33.2	11,12,13,14,15,16	Х,Ү	9,10,11,12
	19,20,21,22,22.2,2 3,23.2,24			14,15,16,17,18,24. 2,27.2,28,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	9,10			2	
786NUZ	- 5901 Ident	tifiler® Plus - EuroFo	orMix			
		17,18,20,25		15,16,18	9,10,11,12,13	
3e	9,10,11,12	10,13,15			8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33		X,Y	10,11,12
	20,21,23,23.2,24		۷.		6,7,9,9.3	8,10,11,14
	15,16,17					
7H4JRV	- 5902 Glob	alFiler™- TrueAllele	® VUler Release 202	2b		
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,10,11,12,13	
3e	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33	11,12,14,15	X,Y	10,11,12
	20,21,23,23.2,24			15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
8DDAHX	Glob	oalFiler™- STRMix™	2.8			
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,10,11,12,13	
3e	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 .2	11,12,14,15	X,Y	10,11,12
	20,21,23,23.2,24			15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	

			TABLE	3		
WebCo	de - Test 🛛 An	nplification Kits -	Probablistic Gen	otyping Software		
	D1\$1656	D251338	D2S441	D3\$1358	D5\$818	D6S1043
ltem	D7\$820	D8S1179	D10S1248	D12S391	D13S317	D16S539
	D18551	D195433	D21511	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	VWA	D12391			t Indel	
8Y8DYV	- 5902 Glob	oalFiler™- TrueAllele	R VUler Release 202	A Kesuits 22b		
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,11,12,13	
3e	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33	11,12,15	X,Y	10,11,12
	20,21,23,23.2,24		.2	15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
9CLYK9	- 5901 Glob	oalFiler™- STRMix™				
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,10,11,12,13	
3e	9,10,11,12	10,13,14**,15	13,14	17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32**,32.2, 33,33.2	11,12,14,15	X,Y	10,11,12
	20,21,23,23.2,24			15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
9MBR8G	Q - 5901 Powe	erPlex® 21- STRMix	™ 2.8.0			
	11,14,16,17.3,18.3	17,18,20,25		15,16,18	9,10,11,12,13	11,12,13,18
3e	9,10,11,12	10,13,15		16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33		X,Y	10,11,12
	20,21,23,23.2,24	9,10,11,12,13	5,8,10,11,12		6,7,9,9.3	8,10,11,14
	15,16,17					
9WLTTR	- 5902 Powe	erPlex® 21- STRMix	™ 2.10.0			
	11,14,16,17.3,18.3	17,18,20,25		15,16,18	9,10,11,12,13	11,12,13,18
3e	9,10,11,12	10,13,14,15		16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 .2		X,Y	10,11,12
	20,21,23,23.2,24	9,10,11,12,13	5,8,10,11,12		6,7,9,9.3	8,10,11,14
	15,16,17					
B7W38V	' - 5901 Glob	oalFiler™- STRMix™				
	11,13,14,15,16,16.3 ,17.3,18.3	16,17,18,20,24,25	10,11,13,14	15,16,17,18	9,10,11,12,13	
3e	9,10,11,12	10,12,13,14,15	12,13,14	16,17,18,19,21	8,10,11,12	8,9,12,13,14
	14,15,16,17,18,19	11,12,13,15.2	30,31.2,32.2,33,33	11,12,13,14,15	X,Y	9,10,11,12
	19,20,21,23,23.2,2 <i>A</i>			15,16,17,18,24.2,2	6,7,9,9.3	8,10,11,14
	15,16,17	9,10			2	
BDR7Z7	- 5901 Powe	erPlex® Fusion 6C-	STRMix™ 2.6.2			
	(11),14,(16),17.3,(18. 3)	17,(18),20,25	11,14	(15),16,(18)	9,(10),11,(12),13	
3e	9,(10),11,(12)	(10),13,(15)	13,(14)	(16),(17),18,19,(20),(21)	(8),(11),12	8,9,(12),13,14
	14,15,17,18,19	12,(13),(15.2)	30,31.2,(32.2),33,3 3.2	(11),(12),(14),15	X,Y	10,(11),(12)
	20,(21),(23),(23.2),(24)	9,(10),(12),(13)	(5),8,(10),11,12	(15),(17),(18),(24.2), 28.2	6,7,(9),(9.3)	8,10,(11),14
	(15),16,17	10	(18),20	17,19		

			TABLE	3		
WebCo	de - Test An	plification Kits -	Probablistic Gene	otyping Software		
	D1\$1656	D2S1338	D2S441	D3S1358	D5S818	D6S1043
ltem	D7\$820	D8S1179	D10S1248	D12S391	D13S317	D16S539
	D18551	D195433	D21S11	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	5E33		IPOX
	VWA	013371	ltom 20 STP	Populto	i indei	
вкннви	l - 5901 Glob	alFiler™- STRMix™	liem Se - STR	Kesulis		
	11,13,14,15,16,16.3	16,17,18,19,20,24, 25	10,11,13,14	14,15,16,17,18	8,9,10,11,12,13	
3e	8,9,10,11,12	9,10,12,13,14,15	12,13,14	16,17,18,19,21	8,10,11,12	8,9,12,13,14
	14,15,16,17,18,19	11,12,13,14.2,15.2	29,30,30.2,31.2,32 ,32.2,33,33.2	11,12,13,14,15,16	X,Y	9,10,11,12
	19,20,21,22,23,23. 2,24			14,15,16,17,17.2,1 8,23.2,24.2,27.2,2 8,2	6,7,9,9.3	8,10,11,14
	14,15,16,17	9,10			2	
BNGT8U	J - 5902 Glob	alFiler™- STRMix™	2.8			
	11,13,14,15,16,16.3 ,17.3,18.3	16,17,18,19,20,24, 25	10,11,13,14,15	14,15,16,17,18	8,9,10,11,12,13	
3e	8,9,10,11,12	9,10,12,13,14,15	12,13,14	16,17,18,19,20,21	8,10,11,12,13	8,9,11,12,13,14
	14,15,16,17,18,19	11,12,13,14.2,15.2	29,30,30.2,31.2,32 ,32.2,33,33.2	11,12,13,14,15,16	X,Y	9,10,11,12
	19,20,21,22,22.2,2 3,23.2,24			14,14.2,15,16,16.2 ,17,17.2,18,23.2,2 4.2,27.2,28,28.2,2 9.2	5,6,7,8,9,9.3	8,10,11,14
	14,15,16,17	9,10			2	
CDTG9N	M - 5901 Glob	alFiler™				
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,10,11,12,13	-
3e	9,10,11,12	10,13,15	13,14	16,17,18,19	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	31.2,32.2,33,33.2	11,12,14,15	X,Y	10,11,12
	20,21,23,23.2,24	-	-	15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10	-	-	2	
CKECAG	Q - 5902 Glob	oalFiler™- TrueAllele	® VUler Release 202	2b		
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,10,11,12,13	
3e	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 .2	11,12,14,15	X,Y	10,11,12
	20,21,23,23.2,24			15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
CRBAWL	J - 5901 Powe	erPlex® ESI-17 Fast	- LiRa v3.0			
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18		
3e		10,13,15	13,14	16,17,18,19,21		8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 .2	11,12,14,15	X,Y	
	20,21,23,23.2,24			15,17,18,24.2,28.2	6,7,9,9.3	
	15,16,17					

			TABLE	3		
WebCo	de - Test 🛛 🗛 An	nplification Kits -	Probablistic Gene	otyping Software		
	D1S1656	D2S1338	D2\$441	D3S1358	D5S818	D6S1043
ltem	D7\$820	D8S1179	D10S1248	D12S391	D13S317	D16S539
	D18S51	D195433	D21511	D22S1045	Amelogenin	CSF1PO
		DYS391	DYS570	5E33 DYS576	Yindel	IPOX
		010071	Item 3e STR	Results	T Maci	
CRVQCV	V - 5901 Glol	oalFiler™- STRMix™		Kesolis		
	11,14,16,17.3,18.3	16,17,18,19,20,25	10,11,13,14	15,16,17,18	9,10,11,12,13	
3e	9,10,11,12	10,12,13,14,15,16	12,13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	11,12,13,14.2,15.2	30,31.2,32.2,33,33	11,12,14,15	X,Y	10,11,12
	19,20,21,22,23,23. 2,24			15,17,18,24.2,27.2 ,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
CXYRCQ	- 5902 Glol	oalFiler™- TrueAllele	® VUIer Release 202	2b		
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,10,11,12,13	
3e	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 .2	11,12,14,15	Х,Ү	10,11,12
	20,21,23,23.2,24			15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
CY6DP6	- 5901 Glol	oalFiler™- STRMix™				
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,10,11,12,13	
3e	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 .2	11,12,14,15	X,Y	10,11,12
	20,21,23,23.2,24			15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10,11**			2	
CZ6EPV	- 5901 Glol	oalFiler™- STRMix™				
	11,13,14,15,16,16.3 ,17.3,18.3	16,17,18,19,20,24, 25	10,11,13,14	15,16,17,18	9,10,11,12,13	
3e	9,10,11,12	10,12,13,14,15	12,13,14	16,17,18,19,21	8,10,11,12	8,9,12,13,14
	14,15,16,17,18,19	11,12,13,14.2,15.2	30,30.2,31.2,32.2, 33,33.2	11,12,13,14,15,16	X,Y	9,10,11,12
	19,20,21,23,23.2,2 4			14,15,17,18,24.2,2 7.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
E34ANN	- 5902 Glol	oalFiler™- TrueAllele	® VUIer Release 202	2b		
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,11,13	
3e	9,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 .2	11,12,14,15	X,Y	10,11,12
	20,21,23,23.2,24			15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
EHXZJK -	5902 Pow	erPlex® 21- STRMix ⁺	™ 2.10			
-	11,14,16,17.3	17,18,20,25		16,18	9,10,11,12,13	11,12,13
3e	9,10,11,12	10,13,15		16,17,18,19	8,12	9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33.2, 33		Х,Ү	10,11,12
	20,21,23,23.2,24	9,11,12	5,8,10,11		6,7,9,9.3	8,10,11,14
	15,16,17					

			TABLE	3		
WebCo	de - Test 🛛 An	plification Kits	- Probablistic Gene	otyping Software		
	D151656	D251338	D25441	D351358	D55818	D651043
Item	D75820	D8S1179	D10S1248	D12S391	D13S317	D16S539
	D18S51	D19S433	D21S11	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	TPOX
	vWA	DYS391	DYS570	DYS576	Y Indel	
	5000 CI-I		Item 3e - SIK			
ELJJPP -		17 19 20 25		15 14 19	0 11 10 12	
30	0 11 12	17,18,20,25	13.14	16,10,10	9,11,12,13	Q Q 1 Q 1 Q 1 A
56	1/ 15 17 18 10	10,13,15	30 31 2 32 2 33 33	11 12 14 15	0,11,12 X V	10 11 12
	14,13,17,10,17	12,13,13.2	.2	11,12,14,13	7,1	10,11,12
	20,21,23,23.2,24			15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
EY6YTN	- 5902 Glob	palFiler™- TrueAllel	e® VUler Release 202	2b		
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,11,12,13	
3e	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 2	11,12,14,15	X,Y	10,11,12
	20,21,23,23.2,24			15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
F7H8UP	- 5902 Glob	oalFiler™- STRMix™	v2.9.1			
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,10,11,12,13	
3e	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 .2	11,12,14,15	X,Y	10,11,12
	20,21,23,23.2,24			15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
F8XQDN	1 - 5902 Gloł	oalFiler™- TrueAllel	e® VUIer Release 202	2b		
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,11,12,13	
3e	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33	11,12,14,15	X,Y	10,11,12
	20,21,23,23.2,24		.2	15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
FPXE6L -	5902 Iden	tifiler® Plus- STRMi	X TM			
		17,18,20,25		15,16,18	9,10,11,12,13	
3e	9,10,11,12	10,13,15			8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33		X,Y	10,11,12
	20,21,23,23.2,24		٤.		6,7,9,9.3	8,10,11,14
	15,16,17					
GCH7H	K - 5902 Gloł	oalFiler™- TrueAllel	e® VUIer Release 202	2b		
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,10,11,12,13	
3e	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33	11,12,14,15	X,Y	10,11,12
	20,21,23,23.2,24		.2	15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	

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			TABLE	3		
WebCo	de - Test Am	plification Kits	- Probablistic Gene	otyping Software		
	D1S1656	D2S1338	D2S441	D3S1358	D5\$818	D6S1043
Item	D7\$820	D8S1179	D10S1248	D125391	D13S317	D16S539
	D18S51	D195433	D21S11	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	VWA	D12391			r Indel	
H484W/I	L. 5902 Glob	alFiler™- TrueAllel	ITEM JE - JIK	Kesuits		
11-0-113	11.14.16.17 3.18 3	17.18.20.25	11.14	15.16.18	9.11.12.13	
3e	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8.9.12.13.14
	14,15,17,18,19	12,13,15.2	, 30,31.2,32.2,33,33	11,12,14,15	X,Y	10,11,12
	00.01.02.02.0.04		.2	15 17 10 04 0 00 0	(7 0 0 0	0 10 11 14
	20,21,23,23.2,24	10		15,17,18,24.2,28.2	0,7,9,9.3	8,10,11,14
	5000				2	
heQhil	3902 GIOD	alFiler - TrueAllel	e® VUIEr Kelease 202	15 14 19	0 10 11 10 12	
30	0 10 11 12	17,18,20,25	13.14	16,10,10	9,10,11,12,13	80121314
56	14 15 17 18 10	10,13,15 2	30 31 2 33 33 2	11 12 14 15	0,11,12 X V	10 11 12
	20 21 23 23 2 24	12,13,13.2	00,01.2,00,00.2	15 17 18 24 2 28 2	67993	8 10 11 14
	15.16.17	10		10,17,10,24.2,20.2	2	0,10,11,14
	- 5901 Glob	nlFiler™ IQC				
	(11).14.16.17 3.18 3	17.18.20.25	11.14	15,16,18	(9.10).11.(12.13)	
3e	9.(10).11.(12)	(10),13,15	13.(14)	(16,17),18,19,21	(8,11),12	8,9,(12),13,14
	(14),15,17,18,19	12,13,(15.2)	30,31.2,(32.2),33,3	(11,12,14),15	X,Y	10,11,(12)
			3.2			0.10.11.14
	20,(21,23),23.2,(24			15,(17),18,(24.2),2 8.2	6,7,9,(9.3)	8,10,11,14
	(15),16,17	10			2	
J37QNH	1 - 5902 Powe	rPlex® 21- STRMi	∢™ 2.10.0			
	11,14,16,17.3,18.3	17,18,20,25		15,16,18	9,10,11,12,13	11,12,13,18
3e	9,10,11,12	10,13,15		16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33		X,Y	10,11,12
	20,21,23,23.2,24	9,10,11,12,13	5,8,10,11,12		6,7,9,9.3	8,10,11,13,14
	15,16,17					
J772JG	- 5902 Powe	rPlex® 21- STRMix	<™ 2.10			
	11,14,16,17.3,18.3	17,18,20,25		15,16,18	9,10,11,12,13	11,12,13,18
3e	9,10,11,12	10,13,15		18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,15.2	30,31.2,32.2,33,33		X,Y	10,11,12
	20.21.23.23.2.24	9.10.11.12.13	.2		6.7.9.9.3	8.10.11.14
	15.16.17	,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	0,0,10,12		0,,,,,	0,10,11,11
JHLQHM	1 - 5902 Glob	alFiler™ - DNA×s				
5.120,110	11,14,16,17,3,18,3	17,18.20.25	11,14	15.16.18	9,10.11.12.13	
3e	9,10,11,12	10,13,14,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33	11,12,14,15	X,Y	10,11,12
	20,21,23,23.2,24		.∠	15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	

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			TABLE	3					
WebCo	de - Test 🛛 Am	plification Kits -	Probablistic Geno	otyping Software					
	D1S1656	D2S1338	D2S441	D3S1358	D5S818	D6S1043			
Item	D7S820	D8S1179	D10S1248	D125391	D13S317	D16S539			
	D18551	D195433	D21511	D22S1045	Amelogenin	CSF1PO			
	FGA vWA	DYS391	Penta E DYS570	5E33 DYS576	Yindel	ΤΡΟΧ			
	VIIA	013071	Item 3e - STR	Results	T Maci				
JPGUBX	- 5901 Glob	alFiler™- STRMix™	nem de - ork	110113					
	11,14,16,17.3,18.3	17,18,20,24,25	11,14	16,18	9,11,12,13				
3e	9,10,11,12	10,13,15	13	16,18,19,21	8,11,12	9,12,13,14			
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33	11,12,14,15	X,Y	10,11,12			
	20,23,23.2,24		.2	15,18,24.2,28.2	6,7,9,9.3	8,10,11,14			
	15,16,17	10,11			2				
K9ECCE	- 5902 Powe	erPlex® 21- STRMix	™ 2.10						
	11,14,16,17.3,18.3	17,18,20,25		15,16,18	9,10,11,12,13	11,12,13,18			
Зe	9,10,11,12	10,13,15		16,17,18,19,21	8,11,12	8,9,12,13,14			
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 .2		X,Y	10,11,12			
	20,21,23,23.2,24	9,10,11,12,13	5,8,10,11,12		6,7,9,9.3	8,10,11,14			
	15,16,17								
KPDQAK	KPDQAK - 5902 GlobalFiler™ - STRMix™ v2.9.1								
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,10,11,12,13				
Зe	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14			
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 .2	11,12,14,15	Х,Ү	10,11,12			
	20,21,23,23.2,24			15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14			
	15,16,17	10			2				
L67JVL -	5901 Glob	alFiler™- STRMix™							
	11,13,14,16,16.3,17 .3,18.3	16,17,18,20,24,25	10,11,13,14	15,16,17,18	9,10,11,12,13				
3e	9,10,11,12	10,12,13,14,15	12,13,14	16,17,18,19,21	8,11,12	8,9,12,13,14			
	14,15,16,17,18,19	11,12,13,15.2	30,31.2,32.2,33,33 .2	11,12,14,15	X,Y	9,10,11,12			
	19,20,21,22,22.2,2 3,23.2,24			15,17,18,24.2,27.2 ,28.2	6,7,9,9.3	8,10,11,14			
	15,16,17	10			2				
L8XEJE -	5901 Powe	erPlex® 21- STRMix	™ v2.8.0						
	11,14,16,17.3,18.3	17,18,20,25		15,16,18	9,10,11,12,13	11,12,13,18			
3e	9,10,11,12	10,13,15		16,17,18,19,21	8,11,12	8,9,12,13,14			
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 .2		X,Y	10,11,12			
	20,21,23,23.2,24	9,10,11,12,13	5,8,10,11,12		6,7,9,9.3	8,10,11,14			
	15,16,17								
LBAKVF -	- 5902 Ident	ifiler® Plus- STRMix	™ 2.7.0						
		17,18,20,25		15,16,18	9,10,11,12,13				
3e	9,10,11,12	10,13,15			8,11,12	8,9,12,13,14			
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 .2		X,Y	10,11,12			
	20,21,23,23.2,24				6,7,9,9.3	8,10,11,14			
	15,16,17								

			TABLE	3		
WebCo	de - Test 🛛 Am	plification Kits -	Probablistic Gene	otyping Software		
	D1S1656	D2S1338	D2S441	D3S1358	D5S818	D6S1043
Item	D7S820	D8S1179	D10S1248	D12S391	D13S317	D16S539
	D18551	D195433	D21S11	D22S1045	Amelogenin	CSF1PO
	FGA vWA	DYS391	DYS570	5E33 DYS576	Yindel	ΤΡΟΧ
	VIIA		Item 3e - STR	Results	i maci	
LHUN3H	l - 5902 Glob	oalFiler™- TrueAllele	® VUler Release 202	2b		
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,11,12,13	
3e	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33	11,12,14,15	X,Y	10,11,12
	20,21,23,23.2,24			15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
LUPUWF	- 5901 Powe	erPlex® Fusion 6C-	STRMix™ STRmix V2.	5.11		
	11,16,17.3,18.3	17,18,20,25	11,14	15,16,18	10,11,12,13	
3e	9,10,11,12	10,13,15	13	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 .2	11,14,15	X,Y	10,11,12
	20,21,23,23.2,24	9,10,11,12,13	5,8,10,11	15,17,18,24.2,28.2	6,7,9,9.3	8,11,14
	15,16,17	10	18,20	17,19		
LYKQJL -	5901 Glob	oalFiler™- STRMix™				
	11,13,14,15,16,16.3 ,17.3,18.3	17,18,19,20,24,25	10,11,13,14	14,15,16,17,18	9,10,11,12,13	
3e	8,9,10,11,12	10,12,13,14,15	12,13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,16,17,18,19	11,12,13,15.2	30,31.2,32.2,33,33 .2	11,12,14,15,16	X,Y	10,11,12
	19,20,21,23,23.2,2 4			14,15,17,18,24.2,2 7.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
MW99ZF	- 5902 Glob	oalFiler™- TrueAllele	® VUler Release 202	2b		
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,11,12,13	
3e	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 .2	11,12,14,15	X,Y	10,11,12
	20,21,23,23.2,24			15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
NWNRKI	D - 5901 Powe	erPlex® Fusion 6C-	STRMix™ V2.5.11			
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,10,11,13	
3e	9,11	10,13,15	13,14	16,17,18,19,21	11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 .2	14,15	X,Y	10,11,12
	20,21,23,23.2,24	9,10,11,12,13	5,8,10,12	15,17,18,28.2	6,7,9	8,10,11,14
	15,16,17	10	18,20	17,19		
P3QH3A	- 5901 Powe	erPlex® 21- STRMix	™ 2.8			
	11,14,16,17.3,18.3	17,18,20,25		15,16,18	9,10,11,12,13	11,12,13,18
3e	9,10,11,12	10,13,15		16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 .2		X,Y	10,11,12
	20,21,23,23.2,24	9,10,11,12,13	5,8,10,11,12		6,7,9,9.3	8,10,11,14
	15,16,17					

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			TABLE	3		
WebCo	de - Test 🛛 An	plification Kits -	Probablistic Gene	otyping Software		
	D1\$1656	D251338	D2S441	D3S1358	D5\$818	D6\$1043
ltem	D7\$820	D8S1179	D1051248	D12S391	D13S317	D16S539
	D18551	D195433	D21511	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	VWA	DYS391		DYS576	Y Indel	
	5002 Clak	alF:lor™ CTDM/iv™	Item 3e - SIR	Kesults		
FORLET -			11.14	15 16 18	0 10 11 12 13	
30	9 10 11 12	10 13 15	13.14	16 17 18 19 21	8 11 12	89121314
00	14 15 17 18 19	12 13 15 2	30 31 2 32 2 33 33	11 12 14 15	X Y	10 11 12
	11,10,17,10,17	12,10,10.2	.2	11,12,11,10	~~~	10,11,12
	20,21,23,23.2,24			15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
P8TP7Q	- 5901 Glok	oalFiler™- STRMix™				
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,10,11,12,13	
3e	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33	11,12,14,15	X,Y	10,11,12
	20,21,23,23.2,24		.2	15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
PD88ZF	- 5902 Glob	alFiler™- STRMix™	v2.8			
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,10,11,12,13	
3e	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33	11,12,14,15	X,Y	10,11,12
	00.01.02.02.0.04		.2	15 17 10 04 0 00 0	(7 0 0 0	0 10 11 14
	20,21,23,23.2,24	10		15,17,10,24.2,20.2	0,7,9,9.3	8,10,11,14
	15,10,17 5 5001 Clab				Z	
Q99DQE			11X Z.O.U	15 14 19	0 10 11 10 12	
30	0 10 11 12	10 13 15	13.14	16 17 18 10 21	9,10,11,12,13	80121314
56	14 15 17 18 19	12 13 15 2	30 31 2 32 2 33 33	11 12 14 15	× Y	10 11 12
		12,10,10.2	.2	11,12,14,13	7,1	10,11,12
	20,21,23,23.2,24			15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
R3MFPB	- 5901 Powe	erPlex® Fusion 6C- S	STRMix™ STRmix v2.5	5.11		
	11,14,16,17.3,18.3	17,18,20,25	11,14	16,18	10,11,12,13	
3e	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	31.2,32.2,33,33.2	11,12,14,15	X,Y	10,11,12
	20,21,23,23.2,24	9,10,11,12	5,8,10,11	15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10	18,20	17		
R9RDYE	- 5901 Glok	oalFiler™- STRMix™	STRmix			
	11,13,14,15,16,16.3 ,17.3,18.3	16,17,18,19,20,24, 25	10,11,13,14	14,15,16,17,18	9,10,11,12,13	
3e	8,9,10,11,12	10,12,13,14,15,16	12,13,14	16,17,18,19,20,21	8,10,11,12	8,9,12,13,14
	14,15,16,17,18,19	11,12,13,14.2,15.2	30,31.2,32,32.2,33 ,33.2	11,12,13,14,15,16	X,Y	9,10,11,12
	19,20,21,22.2,23,2 3.2,24			14,15,17,18,24.2,2 7.2,28.2,29.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	

			TABLE	3		
WebCo	de - Test 🛛 Am	plification Kits	- Probablistic Gena	otyping Software		
	D1\$1656	D2S1338	D2S441	D3S1358	D5S818	D6S1043
ltem	D7\$820	D8S1179	D1051248	D12S391	D135317	D16S539
	D18S51	D195433	D21511	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D DVS391	Penta E DYS570	5E33 DV\$576	I HUI V Indel	ΤΡΟΧ
	VWA	013071	ltem 3e STR	Results	i ilidei	
RCPXZD	- 5901 Inves	tiggtor® 24plex Q	S - EuroForMix	INESUIIS		
	[11],14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,10,11,12,13	
3e	9,[10],11,[12]	10,13,15	13,[14]	16,17,18,19,21	[8],11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33	[11,12,14],15	Х,Ү	10,11,12
	20,[21],23,23.2,[24]		.2	15,17,18,24.2,28.2	6,7,9,[9.3]	8,10,11,14
	[15],16,17	10				
TMDEV4	- 5902 Glob	alFiler™				
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,10,11,12,13	
3e	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 .2	11,12,14,15	X,Y	10,11,12
	20,21,23,23.2,24			15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
TZBZVA -	- 5901 Powe	erPlex® Fusion 6C-	STRMix™ V2.5.11			
	14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,10,11,12,13	
3e	9,11	10,13,15	13,14	16,17,18,19,21	11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 .2	14,15	X,Y	10,11,12
	20,21,23,23.2,24	9,10,12,13	5,8,10,12	15,18,28.2	6,7,9	8,10,11,14
	15,16,17	10	18,20	17,19		
UNVR99	- 5902 Glob	alFiler™- TrueAllel	e® VUIer Release 202	2b		
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,11,12,13	
Зе	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 .2	11,12,14,15	X,Y	10,11,12
	20,21,23,23.2,24			15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
VFHZHA	- 5902 Glob	alFiler™- STRMix™	v2.9.1			
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,10,11,12,13	
3e	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 .2	11,12,14,15	X,Y	10,11,12
	20,21,23,23.2,24			15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
VK9834	- 5902 Powe	erPlex® Fusion 6C-	STRMix™ v2.10			
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,10,11,12,13	
3e	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 .2	11,12,14,15	Х,Ү	10,11,12
	20,21,22.2,23,23.2 ,24	9,10,11,12,13	5,8,10,11,12	15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10	18,20	17,19		

			TABLE	3		
WebCoo	de - Test 🛛 🗛 An	plification Kits -	Probablistic Gene	otyping Software		
	D1\$1656	D251338	D2S441	D3\$1358	D5\$818	D6\$1043
ltem	D7\$820	D8S1179	D1051248	D12S391	D13S317	D16S539
	D18551	D195433	D21511	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	VWA	DYS391		DYS576	t Indel	
	5001 Clal		Item 3e - STR	Kesults		
VVZKLNZ			11.14	15 14 19	0 10 11 12 13	
30	0 10 11 12	17,10,20,23	13.14	16 17 18 10 21	9,10,11,12,13	80121314
Je	14 15 17 18 19	12 13 15 2	30 31 2 32 2 33 33	11 12 14 15	× Y	10 11 12
		12,10,10.2	.2	11,12,14,13	7,1	10,11,12
	20,21,23,23.2,24			15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
WRGKYZ	C - 5901 Glob	oalFiler™- STRMix™				
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,10,11,12,13	
Зe	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33	11,12,14,15	X,Y	10,11,12
	20,21,23,23.2,24			15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
WU8M28	3 - 5901 Gloł	oalFiler™- STRMix™	2.8			
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,10,11,12,13	
3e	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 .2	11,12,14,15	Х,Ү	10,11,12
	20,21,23,23.2,24			15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
X28843 -	- 5901 Pow	erPlex® 21- STRMix	™ V2.8.0			
	11,14,16,17.3,18.3	17,18,20,25		15,16,18	9,10,11,12,13	11,12,13,18
3e	9,10,11,12	10,13,15		16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33		X,Y	10,11,12
	20,21,23,23.2,24	9,10,11,12,13	5,8,10,11,12		6,7,9,9.3	8,10,11,14
	15,16,17					
YGJ6L9 -	- 5901 Glob	oalFiler™- STRMix™				
	11,14,16,17.3,18.3	17,18,19,20,24,25	10,11,13,14	15,16,17,18	9,10,11,12,13	
3e	9,10,11,12	10,12,13,14,15	12,13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,16,17,18,19	11,12,13,15.2	30,31.2,32.2,33,33	11,12,14,15	X,Y	10,11,12
	19,20,21,22,22.2,2 3,23.2,24		.۷	15,17,18,24.2,27.2 ,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10			2	
YJMUQ3	- 5901 Pow	erPlex® Fusion 6C-	STRMix™ V2.5.11			
	16,17.3	17,18,20	11,14	15,16	10,11	
3e	ND	10,13,15	13	18,19	12	9
	15,17,19	12,15.2	ND	ND	Y	10
	ND	12	5,8	17	ND	8
	15,16	ND	ND	ND		

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			TABLE	3					
WebCoo	WebCode - Test Amplification Kits - Probablistic Genotyping Software								
	D1S1656	D2S1338	D2S441	D3S1358	D5S818	D6S1043			
ltem	D75820	D8S1179	D1051248	D125391	D135317	D16S539			
	FGA	Penta D	Penta E	SE33	TH01	ТРОХ			
	vWA	DYS391	DY\$570	DYS576	Y Indel				
			ltem 3e - STR	Results					
Z8LUQ4	- 5902 Glob	alFiler™- TrueAllel	e® VUler Release 202	2b					
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,11,12,13				
3e	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14			
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 .2	11,12,14,15	Х,Ү	10,11,12			
	20,21,23,23.2,24			15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14			
	15,16,17	10			2				

			TABLE	3		
WebCoo	de - Test	Amplification Kits -	Probablistic Gene	otyping Software		
	D151656	D251338	D25441	D351358	D55818	D6510/3
ltem	D75820	D251338	D1051248	D125391	D135317	D16S539
	D18\$51	D195433	D21S11	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	vWA	DYS391	DY\$570	DYS576	Y Indel	
			Item 3sp - STR	Results		
2QGMC	2 - 5902	GlobalFiler™- TrueAllele	® "VUIer Release 20:	22b		
	14,18.3	17,20,24,25	11,14	15,16	9,11	
Зsр	9,11	13,15	13,14	19,21	11,12	8,13
	17,18	12,13	30,31.2	15,15	X,Y	10,11
	20,23.2			15,27.2,28.2	6,9	8,10
	16,17	10			2	
34G3B3	- 5902	GlobalFiler™- STRMix™ 2	2.8			
	13,13.2,14,17.3, 1,18.3	,18. 19,20,21,24,25,26	10,11,12,13,14,15	14,15,16,17,18	8,9,10,11,12	
Зsр	8,9,10,11,12	12,13,14,15,16	12,13,14	18,19,20,21	10,11,12,13	7,8,12,13,14
	16,17,18,19	11,12,13	29,30,30.2,31,31.2	13,14,15,16	X,Y	9,10,11,12
	19,20,21,21.2,22, 23.2,24.2	2.2		14,14.2,15,16,26.2	5,6,8,9	7,8,9,10
	15,16,17,18	9,10			2	
37J2CU	- 5901	GlobalFiler™- STRMix™				
	14,18.3	20,25	11,14	15,18	9,11	
Зsр	9,11	13,15	13,14	19,21	11,12	8,13
	17,18	12,13	30,31.2	15,15	X,Y	10,11
	20,23.2			15,28.2	6,9	8,10
	16,17	10			2	
3MQZUY	(- 5902	Identifiler® Plus- STRMix	м			
		20,25		15,18	9,11	
3sp	9,11	13,15			11,12	8,13
	17,18	12,13	30,31.2		X,Y	10,11
	20,23.2				6,9	8,10
	16,17					
3WHFJ3	- 5902	GlobalFiler™- STRMix™ \	/2.9.1			
	14,18.3	20,25	11,14	15,18	9,11	
3sp	9,11	13,15	13,14	19,21	11,12	8,13
	17,18	12,13	30,31.2	15	Х,Ү	10,11
	20,23.2			15,28.2	6,9	8,10
	16,17	10			2	
44PDRD	- 5901	GlobalFiler™- STRMix™				
	14,18.3	20,25	11,14	15,18	9,11	
3sp	9,11	13,15	13,14	19,21	11,12	8,13
	17,18	12,13	30,31.2	15	Х,Ү	10,11
	20,23.2,24.2**	*		15,28.2	6,9	8,10
	16,17	10			2	
4BEUJW	- 5902	PowerPlex® 21- STRMix™	2.10.0			
	14,18.3	20,25		15,18	9,11	12,18
Зsр	9,11	13,15		19,21	11,12	8,13
	17,18	12,13	30,31.2		X,Y	10,11
	20,23.2	10,13	8,12		6,9	8,10
	16,17					

			TABLE	3		
WebCod	le - Test 🛛 🗛	mplification Kits -	Probablistic Gen	otyping Software		
	D1S1656	D2S1338	D2S441	D3S1358	D5S818	D6S1043
ltem	D7\$820	D8S1179	D10S1248	D12S391	D13S317	D16S539
	D18S51	D195433	D21511	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	vWA	DYS391	DYS570	DYS576	Y Indel	
	5000 D		Item 3sp - STI	R Results		
4FBJY4 -	5902 Pov	werPlex® Fusion 6C -	DINAXS	15.10	0.11	
0	14,18.3	20,25	11,14	15,18	9,11	0.10
3sp	9,11	13,15	13,14	19,21	11,12	8,13
	17,18	12,13	30,21.2	15	X, Y	10,11
	20,23.2	10,13	8,12	15,28.2	6,9	8,10
	16,17	10	18	19		
4UQG64	- 5901 Glo	balFiler™- STRMix™				
	13,14,17.3,18.3	19,20,24,25	10,11,13,14	14,15,17,18	8,9,10,11	
Зsр	8,9,10,11	12,13,14,15	12,13,14	18,19,20,21	10,11,12	8,12,13
	16,17,18	12,13	29,30,30.2,31.2	14,15,16	X,Y	9,10,11
	19,20,22.2,23.2			14,15,27.2,28.2	6,9	8,10
	15,16,17	10			2	
4XQRZ4 -	- 5901 Glo	balFiler™- STRMix™				
	13,14,17.3,18.3	19,20,24,25	10,11,13,14	14,15,17,18	8,9,10,11	
Зsр	8,9,10,11	12,13,14,15	12,13,14	18,19,20,21	10,11,12	8,12,13
	16,17,18	12,13	29,30,30.2,31.2	14,15,16	X,Y	9,10,11
	19,20,22.2,23.2			14,15,27.2,28.2	6,9	8,10
	15,16,17	10			2	
62K9Q4	- 5901 Glo	balFiler™- STRMix™				
	13,14,17.3,18.3	19,20,24,25	10,11,13,14	14,15,17,18	8,9,10,11	
Зsр	8,9,10,11	12,13,14,15	12,13,14	18,19,20,21	10,11,12	8,12,13
	16,17,18	12,13	29,30,30.2,31.2	14,15,16	X,Y	9,10,11
	19,20,22.2,23.2,24			14,15,27.2,28.2,29	6,9	8,10
	.2	10		.2	0	
	15,10,17				2	
63VBC2 -	- 5901 Glo	obalFiler ^m - STRMix ^m				
	13,14,15,17.3,18.3	19,20,24,25	10,11,13,14	14,15,17,18	8,9,10,11	
Зsр	8,9,10,11	12,13,14,15	12,13,14	18,19,20,21	10,11,12	8,12,13
	16,17,18	11,12,13	29,30,30.2,31.2	14,15,16	X,Y	9,10,11,12
	19,20,22.2,23.2,24			14,15,27.2,28.2,29	6,9	8,10
	15,16,17	10		.2	2	
6PTH6Y -	5902 Glo	balFiler™- TrueAllele	® VUIer Release 202	22b		
	14,18.3	20,25	11,14	15,18	9,11	
3sp	9,11	_13,15	13,14	19,21	11,12	8,13
1	17,18	12,13	30,31.2	15	X,Y	10,11
	20,23.2	·		15,28.2	6,9	8,10
	16,17	10			2	

			TABLE	3		
WebCoo	de - Test 👘	Amplification Kits -	Probablistic Gene	otyping Software		
	D1S1656	D2S1338	D2S441	D3S1358	D5S818	D6S1043
ltem	D7S820	D8S1179	D10S1248	D12S391	D13S317	D16S539
	D18S51	D195433	D21S11	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	VWA	D122A1		D13576	r Indei	
7746RV	5901 G	SlobalFiler™ STRMix™	Item 3sp - 51k	Kesuits		
7770001	13 14 17 3 18 3	19 20 24 25	10 11 13 14	14 15 17 18	891011	
3sp	8.9.10.11	12.13.14.15	12.13.14	18.19.20.21	10.11.12	7.8.12.13
1-	16,17,18	12,13	29.30.30.2.31.2	14,15,16	X.Y	9,10,11
	19,20,22.2,23.2	,	., , . , .	14,14.2,15,27.2,28	6,9	8,10
				,28.2		
	15,16,17	9,10			2	
786NUZ	- 5901 lo	lentifiler® Plus - EuroF	orMix			
		17,18,20,25		15,16,18	9,10,11,12,13	
Зsр	9,10,11,12	10,13,15			8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 .2		X,Y	10,11,12
	20,21,23,23.2,24	4			6,7,9,9.3	8,10,11,14
	15,16,17					
7H4JRV -	- 5902 G	GlobalFiler™- TrueAllele	e® VUIer Release 202	2b		
	14,18.3	20,25	11,14	15,18	9,11	
Зsр	9,11	13,15	13,14	19,21	8,11,12	8,13
	17,18	12,13	30,31.2	15,15	X,Y	10,11
	20,23.2			15,28.2	6,9	8,10
	16,17	10			2	
8DDAHX	- 5901 G	GlobalFiler™- STRMix™	2.8			
	14,18.3	20,25	11,14	15,18	9,11	
Зsр	9,11	13,15	13,14	19,21	11,12	8,13
	17,18	12,13	30,31.2	15	X,Y	10,11
	20,23.2			15,28.2	6,9	8,10
	16,17	10			2	
8Y8DYV	- 5902 G	GlobalFiler™- TrueAllele	e® VUIer Release 202	2b		
Зsр						
	5001					
YCLYKY -	- 5901 G	lobalFiler™-SIKMix™	11.14	15.10	0.13	
2	14,18.3	20,25	11,14	15,18	9,11	0.10
Зsр	9,11	13,15	13,14	19,21	11,12	8,13
	1/,18	12,13	30,31.2	15 29 2	Λ, Ϊ	9 10
	16.17	10		13,28.2	رب رب ر	0,10
	10,17	ĨŬ			۷.	

			TABLE	3		
WebCo	de - Test 🛛 🗛	mplification Kits -	Probablistic Gene	otyping Software		
	D1\$1656	D2S1338	D2S441	D3S1358	D5\$818	D6S1043
ltem	D7\$820	D8S1179	D1051248	D12S391	D13S317	D16S539
	D18S51	D195433	D21511	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	VWA	DYS391	DYS570	DYS576	Y Indel	
	5001 Pou	UnerDlay® 01 STRAG	Item 3sp - SIN	R Results		
71010100	11 14 16 17 3 18 3	17 18 20 25	2.0.0	15 16 18	9 10 11 12 13	11 12 13 18
3sn	9 10 11 12	10 13 15		17 18 19 21	8 11 12	8 9 12 13 14
035	14 15 17 18 19	12 13 15 2	30 31 2 32 2 33 33	17,10,17,21	X Y	10 11 12
		12,10,10.2	.2		7,1	10,11,12
	20,23,23.2,24	9,10,12,13	5,8,10,11,12		6,7,9,9.3	8,10,11,14
	15,16,17					
9WLTTR	- 5902 Pov	verPlex® 21- STRMix*	™ 2.10.0			
	11,14,16,17.3,18.3	17,18,20,25		15,16,18	9,11,12,13	12,18
Зsр	9,10,11,12	10,13,15		16,17,19,21	11,12	8,9,13
	14,15,17,18,19	12,13,15.2	30,31.2,32.2		X,Y	10,11
	20,23,23.2	9,10,13	8,10,11,12		6,7,9	8,10,14
	16,17					
B7W38V	- 5901 Glo	balFiler™- STRMix™				
	13,14,17.3,18.3	19,20,24,25	10,11,13,14	14,15,17,18	8,9,10,11	
Зsр	8,9,10,11	12,13,14,15	12,13,14	18,19,20,21	10,11,12	8,12,13
	16,17,18	12,13	29,30,30.2,31.2	14,15,16	X,Y	9,10,11
	19,20,22.2,23.2			14,14.2,15,27.2,28	6,9	8,10
	15,16,17	9,10		,20.2	2	
BDR7Z7	- 5901 Pov	verPlex® Fusion 6C- 3	STRMix™ 2.6.2			
	14,18.3	(18),20,25	11,14	15,18	9,11	
Зsp	9,11	13,15	13,14	(16),19,21	11,12	8,13
	17,18	12,13	30,31.2,(33.2)	15	X,Y	10,11
	20,23.2	10,13	8,(10),12	15,(17),28.2	6,9	8,10
	16,17	10	18	19		
BKHHBU	- 5901 Glo	balFiler™- STRMix™				
	13,13.2,14,17.3,18. 1,18.3	19,20,24,25	10,11,13,14	14,15,17,18	8,9,10,11	
Зsр	8,9,10,11	12,13,14,15,16	12,13,14	18,19,20,21	10,11,12	7,8,12,13,14
	16,17,18	11,12,13	29,30,30.2,31.2	14,15,16	X,Y	9,10,11
	19,20,22.2,23.2,24 .2			14,14.2,15,16,27.2 ,28,28.2	6,9	8,10
	15,16,17	9,10			2	
BNGT8U	I - 5902 Glo	balFiler™- STRMix™	2.8			
	13,13.2,14,17.3,18. 1,18.3	19,20,24,25	10,11,12,13,14,15	14,15,17,18	8,9,10,11	
Зsр	8,9,10,11	12,13,14,15	12,13,14	18,19,20,21	10,11,12,13	7,8,12,13
	16,17,18	11,12,13	29,30,30.2,31.2	13,14,15,16	Х,Ү	9,10,11
	19,20,22.2,23.2,24 .2			14,14.2,15,16,27.2 ,28,28.2,29.2	5,6,8,9	7,8,9,10
	15,16,17	9,10			2	

WebCode Test Amplification Kits - Probabilistic Genotyping Softwore Dissist DoS1043 Item D151656 D25133 D25411 D251315 D153213 D153213 D153213 D153213 D153213 D153231 D153313 D153333 D133313 D153333 D133313 D153333 D133313 D133313 D133313 D133313 D133313 D133313 D133313 D133313				TABLE	3		
Item D151656 D251338 D25411 D351358 D353136 D551143 D16553 Item D15531 D195433 D213179 D16553 D16553 D16553 FGA Penta D Penta E E33 TH01 TPCX VM DYS370 DYS370 DYS370 V Indel CDTG9M - 5901 GlobalFiler* Item 3sp - STR Results 0.11 - 20232 - - 15.28.2 6.9 8.10 16,17 10 - - 2 CKCCAQ 5002 GlobalFiler* 10.11 20.23 - 15.28.2 6.9 8.10 16,17 10 - - 2 CKCCAQ 5002 GlobalFiler*< 10.11 20.32 6.9 8.13 20,17.18 12,13 30,31.2 15,15 X,Y 10.11 20,23.2 11,14 15,18 9,11 3.14 12,21 8.13 301 17,18 12,13 30	WebCo	de - Test	Amplification Kits -				
Hem D75820 D6351179 D1051248 D125301 D135317 D165530 CA Penico Derritol S333 THO TPOX VMA D13531 D195433 D21511 D2251045 Amelogenin C5F1P0 VMA D13570 D75570 Y Indel TPOX TPOX CDTG9W -5901 GlobalFiler" THEm 35p - STR Results State State CDTG9W -5901 GlobalFiler" 11,14 15,18 9,11 - - 20232 - - 15,282 6,9 8,10 16,17 10 - - 2 CKCCQ0 5902 GlobalFiler"- TrueAllele® VUle Releose 2022b - 2 1,14 15,18 9,11 2,023 10,11 2,023 10,11 2,023 10,11 2,023 10,11 2,023 10,11 2,023 10,11 2,023 10,11 2,023 10,11 2,023 10,11 2,023 10,11 1,13,13 13		D151656	D251338	D25441	D351358	D55818	D651043
D1853 D195433 D21511 D2251043 Amelogenin CSFIPO road Diss70 DYS370 DYS370 DYS376 Y Indel road Item 35p - STR Results Diss70 DYS370 DYS370 DYS370 Y Indel CDIO9M - 5901 GlobalFiler** 14,18,3 20,25 11,14 15,18 9,11 - - 30 9,11 13,15 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15 XY 10,11 20,23.2 - - 2 - 2 - - 2 - - 2 - - 2 - - 2 - - 2 - - 2 - - 2 - - - - 2 - - - 2 - - - - - - - - - - - - - - -	ltem	D75820	D8S1179	D1051248	D125391	D13S317	D165539
FGA Penta B SE33 TH01 TPOX VWA DYS391 DYS370 DYS370 <t< th=""><th></th><th>D18S51</th><th>D195433</th><th>D21S11</th><th>D22S1045</th><th>Amelogenin</th><th>CSF1PO</th></t<>		D18S51	D195433	D21S11	D22S1045	Amelogenin	CSF1PO
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Hem 3sp - STR Results 3up 61,1 61,0bol/Filer** 3up 9,11 13,15 13,14 19,21 11,12 8,13 20,23.2 - - 15,28.2 6,9 8,10 16,17 10 - - 2 2 CKECAQ - 5902 Globol/Filer**< TrueAllele® VUler Release 2022b		vWA	DYS391	DYS570	DYS576	Y Indel	
CDI GWA - 3901 GlobalPiler ¹⁸ 34p 9,11 13,15 13,14 19,21 11,12 8,13 34p 9,11 13,15 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15 X,Y 10,11 20,23.2 - - 15,28.2 6,9 8,10 16,17 10 - - 2 2 14,18.3 20,25 11,14 15,18 9,11 31,3 31 13,15 13,14 19,27 11,12 8,13 17,18 12,13 30,31.2 15,15 X,Y 10,11 20,23.2 . 15,28.2 6,9 8,10 16,17 10 2 2 15,15 X,Y 20,23.2 . 15,15 X,Y 10,11 12,13 30,31.2 15,15 X,Y 13,14 19,21 13,15 13,14 19,20 10,11,12 8,12,13,14				ltem 3sp - ST	R Results		
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$\begin{array}{ c c c c c c c c c c c c c c c c c c c$	Зsр	9,11	13,15	13,14	19,21	11,12	8,13
20,23 2 . . 15,28 2 6,9 8,10 16,17 10 . . 2 CKECAQ - 5902 GlobalFiler [™] - TrueAllele [®] VUler Release 2022b . 2 14,18.3 20,25 11,14 15,18 9,11 3ap 9,11 13,15 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15,15 X,Y 10,11 2023.2 15,15 X,Y 10,11 2 CRBAWU - 5901 PowerNew® ESI-17 Fast - LiRa v3.0 14,18.3 20,25 11,14 15,15 X,Y 2023.2 13,14 19,21 8,13 17,18 12,13 30,31.2 15,15 X,Y 2023.2 16,17 10 2 16,17 8,9,10,11 11,14,17.3,18.3 19,20,24,25 10,11,13,14 14,15,17,18 8,9,10,11 3sp 8,9,10,11 12,13,14,15 12,13,14 18,19,20,21 10,11,12 8,13 15,16,17 10 2		17,18	12,13	30,31.2	15	X,Y	10,11
16,17 10 . 2 CKECAQ 5902 GlobalFiler**. TrueAllele® VUler Release 2022b 14,18.3 20,25 11,14 15,18 9,11 3sp 9,11 13,15 13,14 19,21 11,12 8,13 20,232 30,31.2 15,15 X,Y 10,11 20,232 30,31.2 15,15 X,Y 10,11 20,232 16,17 10 2 2 CRBAWU<-5901		20,23.2	-	-	15,28.2	6,9	8,10
CKECAQ 5902 GlobalFiler ** TrueAllele® VUler Release 2022b 14,18.3 20,25 11,14 15,18 9,11 3sp 9,11 13,15 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15,15 X,Y 10,11 20,232 15,28.2 6,9 8,10 16,17 10 2 2 CRAWU 5901 PowerPlex® E51.17 Fost - LiRa v3.0 8,13 17,18 12,13 30,31.2 15,15 XY 20,25.2 16,17 8,13 17,18 12,13 8,13 17,18 12,13 30,32,12 15,15 XY 9,11 13,14,17,3,18.3 19,20,24,25 10,11,13,14 14,15,17,18 8,9,10,11 19,00,22,23,2 12,13 29,30,30,2,31.2 14,15,16 XY 9,10,11 19,00,22,23,2 11,14 15,18,920,21 10,11,12 8,12,13,14 19,20,22,23,2 GlobalFiler** - TrueAllele® VUler Release 20222b 2 2 <td></td> <td>16,17</td> <td>10</td> <td>-</td> <td>-</td> <td>2</td> <td></td>		16,17	10	-	-	2	
$\begin{array}{ c c c c c } & 14,18.3 & 20,25 & 11,14 & 15,18 & 9,11 \\ \hline & 17,18 & 12,13 & 30,31.2 & 15,15 & XY & 10,11 \\ \hline & 20,23.2 & 15,28.2 & 6,9 & 8,10 \\ \hline & 16,17 & 10 & 2 \\ \hline & 20,23.2 & 15,28.2 & 6,9 & 8,10 \\ \hline & 16,17 & 10 & 2 \\ \hline & 16,17 & 10 & 2 \\ \hline & 16,18 & 12,13 & 30,31.2 & 15,15 & XY & 10,11 \\ \hline & 12,13 & 13,14 & 19,21 & 8,13 \\ \hline & 17,18 & 12,13 & 30,31.2 & 15,15 & XY & 10,11 \\ \hline & 12,13 & 12,13 & 30,31.2 & 15,15 & XY & 10,11 \\ \hline & 12,13 & 12,13 & 30,31.2 & 15,15 & XY & 10,11 \\ \hline & 12,13 & 12,13 & 12,20,24,25 & 10,11,13,14 & 14,15,17,18 & 8,9,10,11 \\ \hline & 12,0222,23.2 & 14,15,17,18 & 8,9,10,11 & 12,13,14,15 & 12,13,14 & 18,19,20,21 & 10,11,2 & 8,12,13,14 \\ \hline & 16,17,18 & 12,13 & 29,30,30,2,31.2 & 14,15,16 & XY & 9,10,11 \\ \hline & 12,0222,23.2 & 10,11,13 & 13,14 & 15,18 & 9,11 \\ \hline & 12,0222,23.2 & 11,14 & 15,18 & 9,11 \\ \hline & 12,0222,23.2 & 11,14 & 15,18 & 9,11 \\ \hline & 12,0222,23.2 & 11,14 & 15,18 & 9,11 \\ \hline & 12,0222,23.2 & 11,14 & 15,18 & 9,11 \\ \hline & 12,0222,23.2 & 11,14 & 15,18 & 9,11 \\ \hline & 12,0222,23.2 & 11,14 & 15,18 & 9,11 \\ \hline & 12,0223 & 10,11,12 & 8,13 \\ \hline & 16,17 & 10 & 2 \\ \hline & 16,1$	CKECAG	2 - 5902	GlobalFiler™- TrueAllele®	VUler Release 202	22b		
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$ \begin{array}{ c c c c c c } \hline 17,18 & 12,13 & 30,31.2 & 15,15 & X,Y & 10,11 \\ \hline 20,23.2 & 15,28.2 & 6,9 & 8,10 \\ \hline 16,17 & 10 & 2 \\ \hline \hline$	Зsр	9,11	13,15	13,14	19,21	11,12	8,13
20,23.2 15,28.2 6,9 8,10 16,17 10 2 CRBAWU<-5901		17,18	12,13	30,31.2	15,15	X,Y	10,11
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CRBAWU - 5901 PowerPlex® ESI-17 Fast - LiRa v3.0 14,18.3 20,25 11,14 15,18 3sp 17,18 12,13 30,31.2 15,15 X,Y 20,23.2 15,28.2 6,9 16,17 13,14 14,15,17,18 8,9,10,11 3sp 8,9,10,11 12,13,14,15 12,13,14 14,15,17,18 8,9,10,11 3sp 8,9,10,11 12,13,14,15 12,13,14 14,15,17,18 8,9,10,11 19,20,22,2,3.2 10,11,13,14 14,15,17,18 8,9,10,11 19,20,24,25 10,11,13,14 14,15,17,18 8,9,10,11 3sp 8,9,10,11 12,13 29,30,30,2,31.2 14,15,17,18 8,9,10,11 15,16,17 10 2 CXYRCQ - 5902 GlobalFiler ^w TrueAllele® VUIer Release 2022b 4,18.3 20,25 11,14 15,18 9,11 3sp 9,11 13,15 13,14 19,21 11,12 8,13 16,17 10 2 2 11,14 15,18,19** 9,11		16,17	10			2	
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$\begin{tabular}{ c c c c c c c } \hline $15,16,17$ 10 & 2 \\ \hline $CYRCQ - $5902 & $GlobalFiler''' - $TrueAllele® VUler Release 2022b \\ $14,18.3$ $20,25$ $11,14$ $15,18$ $9,11$ \\ \hline $14,18.3$ $20,25$ $11,14$ $15,18$ $9,11$ \\ \hline $20,23.2$ $15,15$ X,Y $10,11$ \\ \hline $20,23.2$ $15,28.2$ $6,9$ $8,10$ \\ \hline $16,17$ 10 2 \\ \hline $CY6DP6 - 5901 $GlobalFiler''' - $TRMix'''$ $$$ $11,14$ $15,18,19**$ $9,11$ \\ \hline $14,18.3$ $20,21**,25$ $11,14$ $15,18,19**$ $9,11$ \\ \hline $14,18.3$ $20,21**,25$ $11,14$ $15,18,19**$ $9,11$ \\ \hline $14,18.3$ $20,21**,25$ $11,14$ $15,18,19**$ $9,11$ \\ \hline $20,23.2,24.2^{**}$ $12,13$ $30,31.2$ 15 X,Y $10,11$ \\ \hline $20,23.2,24.2^{**}$ 15 X,Y $10,11$ \\ \hline $20,23.2,24.2^{**}$ $15,28.2$ $6,9$ $8,10$ \\ \hline $16,17$ 10 2 \\ \hline $CZ6EPV - 5901 $GlobalFiler''' - $TRMix'''$ $15,28.2$ $6,9$ $8,10$ \\ \hline $16,17$ 10 2 \\ \hline $CZ6EPV - 5901 $GlobalFiler''' - $TRMix'''$ $13,14$ $14,15,17,18$ $8,9,10,11$ \\ \hline $3sp$ $8,9,10,11$ $12,13,14,15$ $12,13,14$ $18,19,20,21$ $10,11,12$ $8,12,13$ \\ \hline $16,17,18$ $12,13$ $29,30,30,2,31.2$ $14,15,16$ X,Y $9,10,11$ \\ \hline $19,20,22,2,23.2$ $10,11,13,14$ $18,19,20,21$ $10,11,12$ $8,12,13$ \\ \hline $16,17,18$ $12,13$ $29,30,30,2,31.2$ $14,15,16$ X,Y $9,10,11$ \\ \hline $19,20,22,2,23.2$ $10,11,13,14$ $18,19,20,21$ $10,11,12$ $8,12,13$ $16,17,18$ $12,13$ $29,30,30,2,31.2$ $14,15,16$ X,Y $9,10,11$ \\ \hline $19,20,22,2,23.2$ $10,11,13,14$ $18,19,20,21$ $10,11,12$ $8,12,13$ $16,17,18$ $12,13$ $29,30,30,2,31.2$ $14,15,16$ X,Y $9,10,11$ \\ \hline $19,20,22,2,23.2$ $10,11,13,14$ $18,19,20,21$ $10,11,12$ $8,12,13$ $16,17,18$ $12,13$ $29,30,30,2,31.2$ $14,15,16$ X,Y $9,10,11$ $12,13,14$ $12,13$ $12,13$ $12,13$ $12,13$ $12,13$ $12,13$ $12,13$ $12,13$ $14,15,16$ X,Y $9,10,11$ $12,13,14$ $13,15$ $13,14$ $13,15$ $13,14$ $14,15,16$ $12,2,2,28$ $14,15,16$ $12,2,2,28$ $15,16$ 17 10 15 $14,15,16$ $12,2,28$ $14,15,16$ $12,2,$		19,20,22.2,23	3.2		14,15,27.2,28.2	6,9	8,10
CXYRCQ - 5902 GlobalFiler™ - TrueAllele® VUler Release 2022b 3sp 14,18.3 20,25 11,14 15,18 9,11 3sp 9,11 13,15 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15,15 X,Y 10,11 20,23.2 15,28.2 6,9 8,10 16,17 10 2 CY6DP6 - 5901 GlobalFiler™ - STRMix™ 14,18.3 20,21**,25 11,14 15,18,19** 9,11 3sp 9,11 13,15 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15 X,Y 10,11 20,23,24.2** 15,28.2 6,9 8,10 2 CZ6EPV - 5901 GlobalFiler™ - STRMix™ 2 2 2 CZ6EPV - 5901 GlobalFiler™ - STRMix™ 13,14,17,3,18,3,19. 19,20,24,25 10,11,13,14 14,15,17,18 8,9,10,11 3sp 8,9,10,11 12,13,14,15 12,13,14 18,19,20,21 10,11,12 <t< td=""><td></td><td>15,16,17</td><td>10</td><td></td><td></td><td>2</td><td></td></t<>		15,16,17	10			2	
$\begin{array}{c c c c c c c c c c c c c c c c c c c $	CXYRCQ	2 - 5902	GlobalFiler™- TrueAllele@	NUler Release 202	22b		
3sp 9,11 13,15 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15,15 X,Y 10,11 20,23.2 15,28.2 6,9 8,10 16,17 10 2 CY6DP6 - 5901 GlobalFiler TM - STRMix TM 1 1,12 8,13 3sp 9,11 13,15 11,14 15,18,19** 9,11 3sp 9,11 13,15 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15 X,Y 10,11 20,23.2,24.2** 10 2 2 10,11 10,11 20,23.2,24.2** 10 2 10,11,13,14 14,15,17,18 8,9,10,11 16,17 10 2 2 10,11,13,14 14,15,17,18 8,9,10,11 3sp 8,9,10,11 12,13,14,15 12,13,14 18,19,20,21 10,11,12 8,12,13 3sp 8,9,10,11 12,13,14,15 12,13,14 18,19,20,21 1		14,18.3	20,25	11,14	15,18	9,11	
$ \begin{array}{ c c c c c c c } \hline 17,18 & 12,13 & 30,31.2 & 15,15 & X,Y & 10,11 \\ \hline 20,23.2 & 15,28.2 & 6,9 & 8,10 \\ \hline 16,17 & 10 & 2 \\ \hline CY6DP6 - 5901 & GlobalFiler^{TM} - STRMix^{TM} & \\ \hline 14,18.3 & 20,21^{**},25 & 11,14 & 15,18,19^{**} & 9,11 \\ \hline 3sp & 9,11 & 13,15 & 13,14 & 19,21 & 11,12 & 8,13 \\ \hline 17,18 & 12,13 & 30,31.2 & 15 & X,Y & 10,11 \\ \hline 20,23.2,24.2^{**} & 15,28.2 & 6,9 & 8,10 \\ \hline 16,17 & 10 & 2 \\ \hline CZ6EPV - 5901 & GlobalFiler^{TM} - STRMix^{TM} & \\ \hline 13,14,17,3,18,3,19 & 19,20,24,25 & 10,11,13,14 & 14,15,17,18 & 8,9,10,11 \\ \hline 3sp & 8,9,10,11 & 12,13,14,15 & 12,13,14 & 18,19,20,21 & 10,11,12 & 8,12,13 \\ \hline 3sp & 8,9,10,11 & 12,13,14,15 & 12,13,14 & 18,19,20,21 & 10,11,12 & 8,12,13 \\ \hline 16,17,18 & 12,13 & 29,30,30,2,31.2 & 14,15,16 & X,Y & 9,10,11 \\ \hline 19,20,22,2,23.2 & \hline 10,11 & 12,13,14,15 & 12,13,14 & 14,15,17,18 & 8,9,10,11 \\ \hline 19,20,22,2,23.2 & \hline 10,11 & 12,13,14,15 & 12,13,14 & 18,19,20,21 & 10,11,12 & 8,12,13 \\ \hline 15,16,17 & 10 & 2 \\ \hline 15,16,17 & 10 & 2 \\ \hline \end{array}$	Зsр	9,11	13,15	13,14	19,21	11,12	8,13
$\begin{array}{c c c c c c c c c c c c c c c c c c c $		17,18	12,13	30,31.2	15,15	X,Y	10,11
$ \begin{array}{ c c c c c c } \hline 16,17 & 10 & 2 \\ \hline CY6DP6 - 5901 & GlobalFiler & STRMix & \\ \hline 14,18.3 & 20,21 & 2,25 & 11,14 & 15,18,19 & 9,11 \\ \hline 3sp & 9,11 & 13,15 & 13,14 & 19,21 & 11,12 & 8,13 \\ \hline 17,18 & 12,13 & 30,31.2 & 15 & X,Y & 10,11 \\ \hline 20,23.2,24.2 & & 15,28.2 & 6,9 & 8,10 \\ \hline 16,17 & 10 & 2 \\ \hline CZ6EPV - 5901 & GlobalFiler & STRMix & \\ \hline 13,14,17.3,18.3,19 & 19,20,24,25 & 10,11,13,14 & 14,15,17,18 & 8,9,10,11 \\ \hline 3sp & 8,9,10,11 & 12,13,14,15 & 12,13,14 & 18,19,20,21 & 10,11,12 & 8,12,13 \\ \hline 16,17,18 & 12,13 & 29,30,30.2,31.2 & 14,15,16 & X,Y & 9,10,11 \\ \hline 19,20,22.2,23.2 & 10,11 & 12,13,14 & 18,19,20,22 & 6,9 & 8,10 \\ \hline 15,16,17 & 10 & 2 & \\ \hline 15,16,17 & 10 & \\ \hline 15,16,17 & 10 & 2 & \\ \hline 15,16,17 & 10 &$		20,23.2			15,28.2	6,9	8,10
GlobalFiler™ - STRMix™ 14,18.3 20,21**,25 11,14 15,18,19** 9,11 3sp 9,11 13,15 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15 X,Y 10,11 20,23.2,24.2** 15,28.2 6,9 8,10 16,17 10 2 CZ6EPV - 5901 GlobalFiler™ - STRMix™ 13,14,17.3,18.3,19. 19,20,24,25 10,11,13,14 14,15,17,18 8,9,10,11 3sp 8,9,10,11 12,13,14,15 12,13,14 18,19,20,21 10,11,12 8,12,13 3sp 8,9,10,11 12,13,14,15 12,13,14 18,19,20,21 10,11,12 8,12,13 16,17,18 12,13 29,30,30.2,31.2 14,15,16,27.2,28.2 6,9 8,10 19,20,22.2,23.2 10 14,15,16,27.2,28.2 6,9 8,10 15,16,17 10 2 2 2		16,17	10			2	
$\begin{array}{c c c c c c c c c c c c c c c c c c c $	CY6DP6	- 5901	GlobalFiler™- STRMix™				
3sp 9,11 13,15 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15 X,Y 10,11 20,23.2,24.2** 15,28.2 6,9 8,10 16,17 10 2 CZ6EPV - 5901 GlobalFiler™ - STRMix™ 3sp 8,9,10,11 12,13,14,15 12,13,14 14,15,17,18 8,9,10,11 3sp 8,9,10,11 12,13,14,15 12,13,14 18,19,20,21 10,11,12 8,12,13 3sp 8,9,10,11 12,13 29,30,30.2,31.2 14,15,16 X,Y 9,10,11 19,20,22.2,23.2 10 14,15,16,27.2,28.2 6,9 8,10 15 16 17	-	14,18.3	20,21**.25	11,14	15,18,19**	9,11	
17,18 12,13 30,31.2 15 X,Y 10,11 20,23.2,24.2** 15,28.2 6,9 8,10 16,17 10 2 CZ6EPV - 5901 GlobalFiler™- STRMix™ 13,14,17.3,18.3,19. 19,20,24,25 10,11,13,14 14,15,17,18 8,9,10,11 3sp 8,9,10,11 12,13,14,15 12,13,14 18,19,20,21 10,11,12 8,12,13 16,17,18 12,13 29,30,30.2,31.2 14,15,16 X,Y 9,10,11 19,20,22.2,23.2 10 14,15,16,27.2,28.2 6,9 8,10 15,16,17 10 2 2 2	3sp	9,11	13,15	13,14	19,21	11,12	8,13
20,23,2,24.2** 15,28.2 6,9 8,10 16,17 10 2 CZ6EPV - 5901 GlobalFiler™- STRMix™ 13,14,17.3,18.3,19. 19,20,24,25 10,11,13,14 14,15,17,18 8,9,10,11 3sp 8,9,10,11 12,13,14,15 12,13,14 18,19,20,21 10,11,12 8,12,13 16,17,18 12,13 29,30,30.2,31.2 14,15,16 X,Y 9,10,11 19,20,22.2,23.2 10 14,15,16,27.2,28.2 6,9 8,10 15,16,17 10 2 2	I-	17.18	12,13	30,31.2	15	X,Y	10,11
16,17 10 2 CZ6EPV - 5901 GlobalFiler™ - STRMix™ 13,14,17.3,18.3,19. 19,20,24,25 10,11,13,14 14,15,17,18 8,9,10,11 3sp 8,9,10,11 12,13,14,15 12,13,14 18,19,20,21 10,11,12 8,12,13 16,17,18 12,13 29,30,30.2,31.2 14,15,16 X,Y 9,10,11 19,20,22.2,23.2 10 14,15,16,27.2,28.2 6,9 8,10 15,16,17 10 2 2		20,23.2,24.2	**		15,28.2	6,9	8,10
CZ6EPV - 5901 GlobalFiler™- STRMix™ 13,14,17.3,18.3,19. 19,20,24,25 10,11,13,14 14,15,17,18 8,9,10,11 3sp 8,9,10,11 12,13,14,15 12,13,14 18,19,20,21 10,11,12 8,12,13 16,17,18 12,13 29,30,30.2,31.2 14,15,16 X,Y 9,10,11 19,20,22.2,23.2 10 14,15,16,27.2,28.2 6,9 8,10 15,16,17 10 2 2		16.17	10		, _	2	
13,14,17.3,18.3,19. 19,20,24,25 10,11,13,14 14,15,17,18 8,9,10,11 3sp 8,9,10,11 12,13,14,15 12,13,14 18,19,20,21 10,11,12 8,12,13 16,17,18 12,13 29,30,30.2,31.2 14,15,16 X,Y 9,10,11 19,20,22.2,23.2 10 14,15,16,27.2,28.2 6,9 8,10 15,16,17 10 2 2	C76FP\/	- 5901	GlobalFiler™- STRMiv™				
3sp 8,9,10,11 12,13,14,15 12,13,14 18,19,20,21 10,11,12 8,12,13 16,17,18 12,13 29,30,30.2,31.2 14,15,16 X,Y 9,10,11 19,20,22.2,23.2 10 14,15,16,27.2,28.2 6,9 8,10 15,16,17 10 2 2		131/17310		10111314	1/151718	8 9 10 11	
3sp 8,9,10,11 12,13,14,15 12,13,14 18,19,20,21 10,11,12 8,12,13 16,17,18 12,13 29,30,30.2,31.2 14,15,16 X,Y 9,10,11 19,20,22.2,23.2 10 14,15,16,27.2,28.2 6,9 8,10 15,16,17 10 2 2 2 2		3	D, 17, 17, ZU, Z4, ZU	10,11,10,14	14,13,17,10	0,7,10,11	
16,17,18 12,13 29,30,30.2,31.2 14,15,16 X,Y 9,10,11 19,20,22.2,23.2 14,15,16,27.2,28.2 6,9 8,10 15,16,17 10 2	Зsр	8,9,10,11	12,13,14,15	12,13,14	18,19,20,21	10,11,12	8,12,13
19,20,22.2,23.2 14,15,16,27.2,28.2 6,9 8,10 ,29.2 2		16,17,18	12,13	29,30,30.2,31.2	14,15,16	X,Y	9,10,11
,29.2		19,20,22.2,23	3.2		14,15,16,27.2,28.2	6,9	8,10
		15 16 17	10		,29.2	2	

			TABLE	3		
WebCoo	de - Test	Amplification Kits - P	robablistic Geno	typing Software		
	D151656	- D251338	D25441	D3\$1358	D55818	D6510/3
ltem	D75820	D851179	D1051248	D125391	D135317	D165539
	D18S51	D19S433	D21S11	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	vWA	DYS391	DYS570	DYS576	Y Indel	
			Item 3sp - STR	Results		
E34ANN	- 5902	GlobalFiler [™] - IrueAllele®	VUIer Release 202	2b		
	14,18.3	20,25	11,14	15,16,18	9,11	
Зsр	9,11	13,15	13,14	19,21	11,12	8,13
	17,18	12,13	30,31.2	15	X,Y	10,11
	20,23.2	10		15,28.2	6,9	8,10
	5, 6, /	10			2	
EHXZJK -	5902	PowerPlex® 21- STRMix™	2.10			
	14,18.3	20,25		15,18	9,11	12,18
Зsр	9,11	13,15		19,21	11,12	8,13
	17,18	12,13	30,31.2		X,Y	10,11
	20,23.2	10,13	8,12		6,9	8,10
	16,17					
ELJJPP - 5	5902	GlobalFiler™- TrueAllele®	VUler Release 202	2b		
	11,14,18.3	20,25	11,14	15,16,18	9,11	
Зsр	9,11	13,15	13,14	19,21	11,12	8,9,12,13,14
	14,17,18	12,13,15.2	30,31.2,33.2	11,15	X,Y	10,11,12
	20,23.2			15,17,28.2	6,9	8,10,14
	16,17	10			2	
EY6YTN -	- 5902	GlobalFiler™ - TrueAllele®	VUler Release 202	2b		
	11,14,18.3	17,18,20,25	11,14	15,16,18	9,11,12,13	
Зsр	9,10,11	13,15	13,14	16,17,19,21	8,11,12	8,9,13
	14,17,18,19	12,13,15.2	30,31.2,33,33.2	15	X,Y	10,11
	20,21,23.2			15,28.2	6,7,9	8,10
	16,17	10			2	
F7H8UP	- 5902	GlobalFiler™- STRMix™ v2	2.9.1			
	14,18.3	20,25	11,14	15,18	9,11	
Зsр	9,11	13,15	13,14	19,21	11,12	8,13
	17,18	12,13	30,31.2	15	X,Y	10,11
	20,23.2			15,28.2	6,9	8,10
	16,17	10			2	
F8XQDM	- 5902	GlobalFiler™- TrueAllele®	VUIer Release 202	2b		
	14,18.3	18,20,25	11,14	15,18	9,11	
Зsр	9,11	13,15	13,14	19,21	11,12	8,13
	17,18	12,13	30,31.2	15,15	X,Y	10,11
	20,23.2			15,28.2	6,9	8,10
	16,17	10			2	
FPXE6L -	5902	Identifiler® Plus- STRMix™				
		20,25		15,18	9,11	
Зsр	9,11	13,15			11,12	8,13
	17,18	12,13	30,31.2		X,Y	10,11
	20,23.2				6,9	8,10
	16,17					

			TABLE	3		
WebCod	de - Test	Amplification Kits -	Probablistic Gen	otyping Software		
	D151656	D251338	D25441	D3\$1358	D5\$818	D651043
ltem	D75820	D8S1179	D10S1248	D125391	D135317	D165539
	D18S51	D19S433	D21S11	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	VWA	DYS391	DYS570	DYS576	Y Indel	
	(5000		Item 3sp - SII	R Results		
GCH/HK	5902	GlobalFiler - TrueAllele	® VUIEr Release 202	220	0.11	
2	14,18.3	20,25	11,14	15,18	9,11	0.10
Ssp	9,11	10,10	13,14	19,21	11,1Z	0,13
	20.23.2	12,13	30,31.2	15,15	۸,1	8 10
	16.17	10		13,20.2	0,7	0,10
	5000				Σ	
∏404 VVJ	- 3902		vuler Release 202	15 10	0.11	
2	14,18.3	20,25	12.14	10,10	9,11	0 1 2
Ssp	9,11	10,10	30.31.2	19,21	××	0,13
	20.23.2	12,13	50,51.2	15 28 2	6.7.9	8 10
	16.17	10		10,20.2	2	0,10
	5002	GlabalFilar™ TrueAllala	® VIIIar Palagaa 200	206	Z	
HLQIIIL	1/102			15 19	0 1 1	
300	0 11	13 15	13.14	10.21	11 12	8 1 3
Jsh	17.18	12.13	30 31 2	15.15	× ×	10.11
	20.23.2	12,13	50,51.2	15 28 2	6.9	8 10
	16.17	10		10,20.2	2	0,10
	5901	GlobalEilar™ IQC				
	14 18 3	20.25	11 14	15 18	9.11	
3sn	9 11	13 15	13.14	19.21	11 12	8 1 3
Osp	17.18	12,13	30.31.2	15.15	X.Y	10.11
	20.23.2	.2/.0	00,0112	15.28.2	6.9	8.10
	16,17	10		/	2	- / · -
137QNH	- 5902	PowerPlex® 21- STRMix ⁺	^ 2 10 0			
507 QI 11	14.18.3	20.25	2.10.0	15.18	9.11	12.18
3sp	9,11	13,15		19,21	11.12	8,13
I	17,18	12,13	30,31.2	.,	, Х,Ү	10,11
	20,23.2	10,13	8,12		6,9,9.3	8,10
	16,17					
	5902	PowerPlex® 21- STRMix ⁺	™ 2.10			
	14,18.3	20,25		15,18	9,11	12,18
Зsp	9,11	13,15		19,21	11,12	8,13
	17,18	12,13	30,31.2		X,Y	10,11
	20,23.2	10,13	8,12		6,9	8,10
	16,17					
JHLQHM	- 5902	GlobalFiler™ - DNAxs				
	14,18.3	20,25	11,14	15,18	9,11	
Зsp	9,11	13,15	13,14	19,21	11,12	8,13
	17,18	12,13	30,31.2	15,15	X,Y	10,11
	20,23.2			15,28.2	6,9	8,10
	16,17	10			2	

			TABLE (3		
WebCo	de - Test 🛛 Am	plification Kits	- Probablistic Geno	typing Software		
	D151656	- D251338	D25441	D351358	D55818	D651043
ltem	D75820	D231338	D1051248	D125391	D135317	D165539
	D18551	D195433	D21S11	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH 01	ΤΡΟΧ
	vWA	DYS391	DY\$570	DYS576	Y Indel	
			ltem 3sp - STR	Results		
JPGUBX	- 5901 Glob	balFiler™- STRMix™	И			
	14,16,18.3	17,18,20,25	11,14	15,16,18	9,11	
Зsр	9,11	10,13,15	13,14	19,21	11,12	8,13
	17,18	12,13,15.2	30,31.2	15	X,Y	10,11
	20,23.2			15,18,28.2	6,9	8,10
	16,17	10			2	
K9ECCE	- 5902 Powe	erPlex® 21- STRMi	x™ 2.10			
	14,16,18.3	17,18,20,25		15,16,18	9,11	12,18
Зsр	9,11	13,14,15		17,19,21	11,12	8,12,13
	17,18	12,13	30,31.2		X,Y	10,11
	20,23,23.2	9,10,13	8,12		6,9	8,10
	16,17					
KPDQAK	(- 5902 Glob	balFiler™- STRMix™	″ v2.9.1			
	14,18.3	20,25	11,14	15,18	9,11	
Зsр	9,11	13,15	13,14	19,21	11,12	8,13
	17,18	12,13	30,31.2	15	X,Y	10,11
	20,23.2			15,28.2	6,9	8,10
	16,17	10			2	
L67JVL -	5901 Glob	balFiler™- STRMix™	и			
	13,14,17.3,18.3	19,20,24,25	10,11,13,14	14,15,16,17,18	8,9,10,11	
Зsр	8,9,10,11	12,13,14,15	12,13,14	18,19,20,21	10,11,12	7,8,12,13
	16,17,18	11,12,13	29,30,30.2,31.2	14,15,16	X,Y	9,10,11
	19,20,22.2,23.2			14,15,27.2,28.2	6,8,9	8,10
	15,16,17	10			2	
L8XEJE -	5901 Powe	erPlex® 21- STRMi	x™ v2.8.0			
	14,16,18.3	20,25		15,16,18	9,11,12	12,13,18
Зsр	9,11	13,14,15		16,19,21	8,11,12,13	8,13,14
	17,18,19	12,13	30,31.2,32.2,33,33 2		X,Y	10,11,12
	20,23.2,24.2	9,10,11,13	8,12		6,9	8,10
	16,17,18					
LBAKVF -	- 5902 Iden	tifiler® Plus- STRM	ix™ 2.7.0			
		20,25		15,18	9,11	
3sp	9,11	13,15			11,12	8,13
1	17,18	12,13	30,31.2		X,Y	10,11
	20,23.2				6,9	8,10
	16,17					
LHUN3H	I - 5902 Glob	oalFiler™- TrueAlle	le® VUIer Release 2022	2b		
	14.18.3	20.25	11,14	15.16.18	9.11	
3sp	9,11	10,13,15	13,14	19,21	11,12	8,12,13
1	17,18	12,13	30,31.2	15,15	Х,Ү	10,11
	20,23.2			15,28.2	6,9	8,10
	16,17	10			2	

			TABLE	3		
WebCo	de - Test 🛛 An	nplification Kits -	Probablistic Geno	typing Software		
	D151656	D251338	D25441	D351358	D55818	D651043
ltem	D75820	D251338	D10S1248	D125391	D135317	D16\$539
	D18S51	D195433	D21S11	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	vWA	DYS391	DYS570	DYS576	Y Indel	
			Item 3sp - STR	Results		
LUPUWF	- 5901 Powe	erPlex [®] Fusion 6C-	STRMix™ STRmix V2.5).		
_	14,18.3	17,18,20,25	11,14	15,16,18	9,11,12	
Зsр	9,11	10,13,15	13,14	16,18,19,21	11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2	11,15	X,Y	10,11,12
	20,23.2	9,10,11,12,13	8,12	15,17,18,28.2	6,/,9	8,10
	6, /	10	18,20	17,19		
LYKQJL -	- 5901 Gloł	oalFiler™- STRMix™				
	13,14,17.3,18.3	19,20,24,25	10,11,13,14	14,15,17,18	8,9,10,11	
Зsр	8,9,10,11	12,13,14,15	12,13,14	18,19,20,21	10,11,12,13	7,8,12,13
	16,17,18	12,13	29,30,30.2,31.2	14,15,16	X,Y	9,10,11
	19,20,21,22.2,23.2			14,15,27.2,28.2	6,8,9	8,10
	15,16,17	10			2	
MW99ZF	- 5902 Gloł	oalFiler™- TrueAllele	® VUIer Release 2022	2b		
	11,14,16,18.3	17,18,20,25	11,14	15,18	9,11,13	
Зsр	9,11	13,15	13,14	19,21	11,12	8,13
	17,18	12,13,15.2	30,31.2	15	X,Y	10,11
	20,23.2			15,28.2	6,9	8,10
	6, /	10			2	
NWNRKI	D - 5901 Pow	erPlex® Fusion 6C-	STRMix™ V2.5.11			
	14,18.3	18,20,25	11,14	15,18	9,11	
Зsр	9,11	13,15	13,14	19,21	11,12	8,9,12,13,14
	17,18	12,13	30,31.2	15	X,Y	10,11,12
	20,23.2	9,10,12,13	8,12	15,28.2	6,9	8,10
	6, /	10	18	19		
P3QH3A	- 5901 Pow	erPlex® 21- STRMix	™ 2.8			
	14,16,17.3,18.3	17,18,20,25		15,16,18	9,11,13	11,12,13,18
Зsр	9,11	10,13,15		16,17,18,19,21	8,11,12	8,9,13,14
	4, 5, /, 8	12,13	30,31.2,32.2,33,33		Х,Ү	10,11
	20,21,23,23.2,24	9,10,13	5,8,10,12		6,9	8,10,11,14
	16,17					
P8KEPF -	- 5902 Gloł	oalFiler™- STRMix™				
	14,18.3	20,25	11,14	15,18	9,11	
Зsp	9,11	13,15	13,14	19,21	11,12	8,13
	17,18	12,13	30,31.2	15	X,Y	10,11
	20,23.2			15,28.2	6,9	8,10
	16,17	10			2	
P8TP7Q	- 5901 Gloi	oalFiler™- STRMix™				
	14,18.3	20,25	11,14	15,18	9,11	
Зsp	9,11,12**	13,14**,15	13,14	19,21	11,12	8,13
	17,18	12,13	30,31.2	15	X,Y	10,11
	20,23.2			15,28.2	6,9	8,10
	16,17	10			2	

			TABLE	3		
WebCo	de - Test	Amplification Kits -	Probablistic Gene	otyping Software		
	D151656	D251338	D25441	D351358	D55818	D651043
ltem	D75820	D8S1179	D1051248	D125391	D135317	D16S539
	D18S51	D195433	D21S11	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH 01	ΤΡΟΧ
	vWA	DYS391	DY\$570	DY\$576	Y Indel	
000075	5000		Item 3sp - STR	R Results		
PD882F	- 5902	GlobalFiler''' - STRMix'''	v2.8			
0	14,18.3	17,20,25	11,14	15,18	9,11	0.10
Зsр	9,11	13,15	13,14	19,21	11,12	8,13
	17,18	12,13	30,31.2	15 00 0	X, Y	10,11
	20,21,23,23.2	10		15,28.2	0,7,9	8,10
	10,17		1: TM 0 0 0		Z	
Q99BQE	= - 5901	GlobalFiler'''' IQC- STRA	Aix [™] 2.8.0	15.10	0.11	
0	14,18.3	20,25	11,14	15,18	9,11	0.10
Зsр	9,11	13,15	13,14	19,21	11,12	8,13
	17,18	12,13	30,31.2	15,15	Χ,Υ	10,11
	20,23.2	10		15,28.2	0,9	8,10
	10,17				Z	
R3MFPB	- 5901	PowerPlex® Fusion 6C-	STRMIX [™] STRmix v2.5	D.	0.11.10	
0	14,17.3,18.3	17,18,20,25	11,14	15,16,18	9,11,13	0.0.10.14
Зsр	9,11	13,15	13,14	18,19,21	11,12	8,9,13,14
	15,17,18,19	12,13,15.2	30,31.2	11,15	λ, Ϊ	10,11,12
	20,23.2	9,10,12,13	0,12	17.10	0,/,9	8,10
	5001		T0,20	17,17		
RYRDIE	- 3901			14151710	0 0 10 11	
2	13,14,17.3,18	3 19,20,24,25	10,11,13,14	14,15,17,18	8,9,10,11	0 10 10
Ssp	9,10,11	12,13,14,15	12,13,14	10,19,20,21	10,11,12	0,12,13
	10,17,10	24	29,30,30.2,31.2	14,15,10	۸,1	9,10,11
	.2	,24		14,13,27.2,20.2	0,7	8,10
	15,16,17	10			2	
RCPXZD	- 5901	nvestigator® 24plex QS	- EuroForMix			
	14,18.3	20,25	11,14	15,18	9,11	
Зsр	9,11	13,15	13,14	19,21	11,12	8,13
	17,18	12,13	30,31.2	15,15	X,Y	10,11
	20,23.2			15,28.2	6,9	8,10
	16,17	10				
TMDEV4	- 5902	GlobalFiler™				
	14,18.3	20,25	11,14	15,18	9,11	
Зsр	9,11	13,15	13,14	19,21	11,12	8,13
	17,18	12,13	30,31.2	15,15	X,Y	10,11
	20,23.2			15,28.2	6,9	8,10
	16,17	10			2	
TZBZVA	- 5901	PowerPlex® Fusion 6C-	STRMix™ V2.5.11			
	14,18.3	20,25	11,14	15,18	9,11	
Зsр	9,11	13,15	13,14	19,21	11,12	8,13
	17,18	12,13	30,31.2	15	X,Y	10,11,12
	20,23.2	10,13	8,12	15,28.2	6,9	8,10
	16,17	10	18	19		

WebCode - Test Amplification Kits - Probabilistic Genotyping Software DIS1656 DS1178 DS1178 DS1178 DS12817 DIS5317 DIS5317 DIS5317 DIS5317 DIS5317 DIS5317 DIS5317 DIS5337 DIS5317 DIS5337 DIS5337 DIS5337 DIS5337 DIS5337 DIS5337 DIS5337 DIS5337 DIS5337 DIS537 VINTUPESTIC CEFTER UNVR99 - 5902 GlobalFiler ** - Trou-Allele® VUler Release 2022b 11,14 15,16,18 9,11 14,18,3 17,18 17,18 17,14 15,11,14 14,18,3 0,20,32 11,14 15,12 6,9 8,10 14,18,3 0,20,32 11,14 15,282 6,9 8,10 <t< th=""><th></th><th></th><th></th><th>TABLE</th><th>3</th><th></th><th></th></t<>				TABLE	3			
Item D151636 D251338 D25441 D351350 D35818 D651043 Item D15532 D15537 D15537 D15537 D15537 F6A Pente D Pente E SE33 TH01 TPOX V/A DYS370 V Intel TH01 TPOX UNVR99 GlobalFiler** Turwhleewe D23141 15,16,18 9,11 3ap 9,11 10,1315 13,14 17,19,21 11,12 8,13,14 10,17,18 12,13 30,31,2,33 15,18,18 9,11 30,31,4 15,18,18 9,11 3ap 9,11 13,15 13,14 15,18,18 9,11 30,31,2 15 X,Y 10,11 20,23,23 11,14 15,18,18 9,11 3,14 10,21 11,12 8,13 20,73,2 C10,17 0 2 2 11,14 10,11 2,13 11,14 10,11 2,13 11,14 12,12 8,13 11,14 12,12 11	WebCo	de - Test 🛛 🗛	mplification Kits -	Probablistic Gen	otyping Software			
Item D75820 D1351179 D1351179 D135317 D13117 D13117 <thd111< th=""> <thd111< th=""> <thd1111< th=""></thd1111<></thd111<></thd111<>		D1\$1656	D251338	D25441	D3\$1358	D5\$818	D651043	
D1851 D195433 D21511 D2251045 Amelogenin CSFIPO VWA DYS391 DYS570 DYS576 Y Indel TPOX VWA DYS391 DYS570 DYS576 Y Indel TPOX UNVR99 5902 GlobolFiler*-Trueklie® VUer Releace 2022b 11,14,16,18,3 17,20,25 11,14 15,16,18 9,11 3vp 9,11 10,13,15 15,14 17,18 12,13 30,31,2,33 15,15 X,Y 10,11 2023,232 GlobolFiler** STRMis" v2.9,1 11,42 8,13 17,18 12,13 30,31,2 15 X,Y 10,11 2023,22 GlobolFiler** STRMis" v2.9,1 11,12 8,13 17,18 12,13 30,31,2 15 X,Y 10,11 20,32 GlobolFiler** STRMis" v2.10 14,18,3 20,25 11,14 15,18 9,11 13,15 13,14 19,21 11,12 8,13 3vp 9,11 13,15 15,14 19,21 11,12 8,13 10,11	ltem	D75820	D8S1179	D10S1248	D12S391	D135317	D165539	
FGA Pento B SE33 TH01 TPOX VMA DY5391 DY5597 Y Indel ILINX89 5502 GlobalFiler**. TrueAllele® VUer Release 2022b 11.114.16.18.8 9.11 10.13.15 13.14 17.19.21 11.12 8.13.14 3sp 9.11 10.13.15 13.14 17.19.21 11.12 8.13.14 17.18 12.13 30.31.2.33 15.15 X.Y 10.11 16.17 10 2 2 8.10.11.14 16.17 10 2 11.14 15.18 9.11 3sp 9.11 13.15 13.14 19.27 11.12 8.13 17.18 12.13 30.31.2 15 X.Y 10.11 3sp 9.11 13.15 13.14 19.27 11.12 8.13 16.17 10 18 19.21 15.28.2 6.9 8.10 16.17 10 18 19 11.12 8.13 10.11 20		D18S51	D195433	D21S11	D22S1045	Amelogenin	CSF1PO	
VWA DYS370 DYS376 Y Indel Imam Sap - STR Results UNVR99 - 5902 GlobalFiler™-TrueAllele® VUler Release 2022b 11,14,16,18,3 17,20,2,5 11,14 15,16,18 9,11 10,13,15 13,14 17,19,21 11,12 8,13,14 17,18 12,13 30,31,2,33 15,15 X,Y 10,11 20,23,23,2 6,09 8,10,11,14 15,18 9,11 30,31,2,33 15,15 X,Y 10,11 3op 9,11 13,15 13,14 19,21 11,12 8,13 11 10,17 10 2 VK9834 · 5902 PowerPlex® Fusion 6C- STRMix * v2.10 14,18,3 20,25 11,14 15,18 9,11 31,31 19,21 11,12 8,13 3op 9,11 13,15 15,14 19,21 11,12 8,13 17,18 12,13 30,31,2 15 X,Y 10,11 20,232,24,24*** 15,28,2 6,9 8,100 16,17		FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ	
UNR89 - 502 GlobalFiler* - TrueAlleber VUer Release 2022b 11,14,16,18.3 17,20,25 11,14 15,16 9,11 0,10,11 30,9 9,11 10,20,25 11,14 15,28.2 6,7,9 8,10,11,14 20,233,22 6,10,0,11,14 15,28.2 6,7,9 8,10,11,14 15,28.2 6,7,9 8,10,11,14 1,1,14 15,28.2 6,7,9 8,10,11 30,9 9,11 13,13,1 11,14 15,28.2 6,9 8,10 14,18.3 20,25 11,14 15,18 9,11 3,13,15 13,12 1,12 8,10 14,18.3 20,25 11,14 15,28.2 6,9 8,10 14,18.3 <th col<="" td=""><td></td><td>vWA</td><td>DYS391</td><td>DYS570</td><td>DYS576</td><td>Y Indel</td><td></td></th>	<td></td> <td>vWA</td> <td>DYS391</td> <td>DYS570</td> <td>DYS576</td> <td>Y Indel</td> <td></td>		vWA	DYS391	DYS570	DYS576	Y Indel	
Order Notable 2020 Stabuline - The Ambrie For Notable 2020 Stabuline - The Ambrie For Notable 2020 Stabuline - The Ambrie For Notable 2020 11,14 15,15 X,Y 10,11 Stabuline -		5002	abalE:lar™ TruaAllala@	Item 3sp - SII	K Kesults			
Intervention Intervention Intervention Intervention 3sp 5,11 10,13,15 13,14 17,18,21 11,12 6,13,14 17,18 12,13 30,31,2,33 15,15 XY 10,11 20,23,23,2 15,28,2 6,7,9 8,10,11,14 16,17 10 2 VFHZHA - \$902 GlobalFiler ^w - STRMix ^w V2,9,1 11,12 8,13 17,18 12,13 30,31,2 15 X,Y 10,11 20,23,2 15,28,2 6,9 8,10 11,12 8,13 16,17 10 2 2 11,12 8,13 11,12 8,13 16,17 10 15,28,2 6,9 8,10 11,12 8,13 11,12 8,13 16,17 10 18 19 11,12 8,13 11,12 8,13 16,17 10 18 19 11,12 8,13 11,12 8,13 16,17 10 18 19 11	UNVK77	- 3702 Gi			15 16 19	0 1 1		
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10.1 12.10 0.00.12.00 16.17 0.1 16.17 16.17 10 2 VFHZHA - 5902 GlobalFiler"- STRMix " v2.9.1 2 14.18.3 20.25 11.14 15.18 9.11 3sp 9.11 13.15 13.14 19.21 11.12 8.13 17.18 12.13 30.31.2 15 X.Y 10.11 20.23.2 15.28.2 6.9 8.10 11.14 16.17 10 2 2 15.28.2 6.9 8.10 16.17 10 2 2 15.28.2 6.9 8.10 16.17 10 18 19.21 11.12 8.13 16.17 10 18 19 11.12 8.13 11.18 12.13 30.31.2 15 X.Y 10.11 20.23.2.2.4.2** 15.28.2 6.9 8.10 11.12 8.13 12.18 12.13 30.31.2 15 X.Y 10.	035	17.18	12.13	30 31 2 33	15 15	X Y	10.11	
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VFHZHA - 5902 GlobalFiler"- STRMix" v2.9.1 3sp 9,11 13,15 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15 X,Y 10,11 20,23.2 15,28.2 6,9 8,10 11,12 8,13 16,17 10 2 VK9834 - 5902 PowerPlex® Fusion 6C: STRMix" v2.10 14,18.3 20,25 11,14 15,18 9,11 3sp 9,11 13,15 13,14 19,21 11,12 8,13 17,18 20,23 10,13 8,12 15,28.2 6,9 8,10 16,17 10 18 19 11,12 8,13 17,18 12,13 30,31.2 15 X,Y 10,11 20,23,24,24** 15,16* 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15 X,Y 10,11 20,23,24,24** 15,28.2 6,9 8,10 16,17,18* 10,11 20,23,24,24** 13,14		16.17	10		10,20.2	2	0,10,11,11	
Million of the formation of the fo	VFH7HA	- 5902 GI	obalEiler™- STRMix™ v	291				
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$\begin{array}{ c c c c c c c c c c c c c c c c c c c$		14,18.3	20,25	11,14	15,18	9,11		
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$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$		17,18	12,13	30,31.2	15	X,Y	10,11	
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$		20,23.2	10,13	8,12	15,28.2	6,9	8,10	
W2RLKZ - 5901 Global/Filer** - STRMix** 3sp 14,18.3 20,25 11,14 15,18 9,11 3sp 9,11 13,15,16** 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15 X,Y 10,11 20,32,2,24.2** 15,28.2 6,9 8,10 16,17,18** 10,11** 2 WRGKYZ - 5901 Global/Filer** STRMix** 14,18.3 20,25 11,14 15,18 9,11 3sp 9,11 13,14**,15 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15 X,Y 10,11 20,23.2,24.2** 15,28.2 6,9 8,10 16,17 10 2 10,11 20,23.2,24.2** 15,28.2 6,9 8,10 16,17 10 2 10,11 20,23.2 11,14 15,18 9,11 3sp 9,11 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15 X,Y 10,11		16,17	10	18	19			
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17,18 12,13 30,31.2 15 X,Y 10,11 20,23,2,24.2** 15,28.2 6,9 8,10 16,17,18** 10,11** 2 WRGKYZ - 5901 GlobalFiler [™] - STRMix [™] 2 3sp 9,11 13,14**,15 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15 X,Y 10,11 20,23,2,24.2** 15,28.2 6,9 8,10 16,17 10 2 2 WU8M28 - 5901 GlobalFiler [™] - STRMix [™] 2.8 4,18.3 20,25 11,14 15,18 9,11 3sp 9,11 13,15 13,14 19,21 11,12 8,13 3sp 9,11 13,15 13,14 19,21 11,12 8,13 3sp 9,11 13,15 13,14 19,21 11,12 8,13 16,17 10 2 2 4,9 8,10 2 16,17 10 2 2 2,18 <td>Зsр</td> <td>9,11</td> <td>13,15,16**</td> <td>13,14</td> <td>19,21</td> <td>11,12</td> <td>8,13</td>	Зsр	9,11	13,15,16**	13,14	19,21	11,12	8,13	
20,23,2,24.2** 15,28.2 6,9 8,10 16,17,18** 10,11** 2 WRGKYZ - 5901 GlobalFiler [™] - STRMix [™] 4,18.3 20,25 11,14 15,18 9,11 3sp 9,11 13,14**,15 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15 X,Y 10,11 20,23,2,24,2** 15,28.2 6,9 8,10 16,17 10 2 WU8M28 - 5901 GlobalFiler [™] - STRMix [™] 2.8 14,18.3 20,25 11,14 15,18 9,11 3sp 9,11 13,15 13,14 19,21 11,12 8,13 3sp 9,11 13,15 13,14 19,21 11,12 8,13 16,17 10 2 2 6,9 8,10 2 X28843 - 5901 PowerPlex® 21 - STRMix [™] V2.8.0 14,17,3,18.3 20,25 15,18 9,11,12,13 12,18 3sp 9,11 13,15 19,21 11,12 8,		17,18	12,13	30,31.2	15	X,Y	10,11	
16,17,18** 10,11** 2 WRGKYZ - 5901 GlobalFiler ** - STRMix ** 14,18.3 20,25 11,14 15,18 9,11 3sp 9,11 13,14**,15 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15 X,Y 10,11 20,23.2,24.2** 15,28.2 6,9 8,10 16,17 10 2 WU8M28 - 5901 GlobalFiler ** - STRMix ** 2.8 14,18.3 20,25 11,14 15,18 9,11 3sp 9,11 13,15 13,14 19,21 11,12 8,13 3sp 9,11 13,15 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15 X,Y 10,11 20,23.2 15,28.2 6,9 8,10 16,17 10 2 X28843 - 5901 PowerPlex® 21- STRMix ** V2.8.0 11,14,17.3,18.3 20,25 15,18 9,11,12,13 12,18 3sp 9,11 13,15 <t< td=""><td></td><td>20,23.2,24.2**</td><td></td><td></td><td>15,28.2</td><td>6,9</td><td>8,10</td></t<>		20,23.2,24.2**			15,28.2	6,9	8,10	
WRGKYZ - 5901 GlobalFiler [™] - STRMix [™] 3sp 14,18.3 20,25 11,14 15,18 9,11 3sp 9,11 13,14**,15 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15 X,Y 10,11 20,23,2,24.2** 15,28.2 6,9 8,10 16,17 10 2 WU8M28 - 5901 GlobalFiler [™] - STRMix [™] 2.8 14,18.3 20,25 11,14 15,18 9,11 3sp 9,11 13,15 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15 X,Y 10,11 20,23.2 15,28.2 6,9 8,10 16,17 10 2 2 X28843 - 5901 PowerPlex® 21- STRMix [™] V2.8.0 2 2 X28843 - 5901 PowerPlex® 21- STRMix [™] V2.8.0 15,18 9,11,12,13 12,18 3sp 9,11 13,15 19,21 11,12 8,13,14 3sp 9,11 13,15 19,21 11,12 8,13,14 <t< td=""><td></td><td>16,17,18**</td><td>10,11**</td><td></td><td></td><td>2</td><td></td></t<>		16,17,18**	10,11**			2		
14,18.3 20,25 11,14 15,18 9,11 3sp 9,11 13,14**,15 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15 X,Y 10,11 20,23.2,24.2** 15,28.2 6,9 8,10 16,17 10 2 WU8M28 - 5901 GlobalFiler ^{Tw} - STRMix TM 2.8 14,18.3 20,25 11,14 15,18 9,11 3sp 9,11 13,15 13,14 19,21 11,12 8,13 3sp 9,11 13,15 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15 X,Y 10,11 20,23.2 15 X,Y 10,11 20,23.2 6,9 8,10 16,17 10 2 2 2 2 2 2 X28843 - 5901 PowerPlex® 21 - STRMix TM V2.8.0 2 15,18 9,11,12,13 12,18 3sp 9,11 13,15 19,21 11,12 8,13,14 17,18,19 12,13 30,	WRGKYZ	C - 5901 GI	obalFiler™- STRMix™					
3sp 9,11 13,14**,15 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15 X,Y 10,11 20,23.2,24.2** 15,28.2 6,9 8,10 16,17 10 2 WU8M28 - 5901 GlobalFiler™- STRMix™ 2.8 2 14,18.3 20,25 11,14 15,18 9,11 3sp 9,11 13,15 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15 X,Y 10,11 20,23.2 15,28.2 6,9 8,10 16,17 20,23.2 10 2 11,14 19,21 11,12 8,13 16,17 10 2 2 10 2 12,18 12,18 12,18 12,18 12,18 12,18 12,18 12,18 12,18 12,18 12,18 12,18 13,14 14,17 10,11 10,11 10,11 10,11 10,11 10,11 10,11 <t< td=""><td></td><td>14,18.3</td><td>20,25</td><td>11,14</td><td>15,18</td><td>9,11</td><td></td></t<>		14,18.3	20,25	11,14	15,18	9,11		
17,18 12,13 30,31.2 15 X,Y 10,11 20,23.2,24.2** 15,28.2 6,9 8,10 16,17 10 2 WU8M28 - 5901 GlobalFiler [™] - STRMix [™] 2.8 2 14,18.3 20,25 11,14 15,18 9,11 3sp 9,11 13,15 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15 X,Y 10,11 20,23.2 10 15,28.2 6,9 8,10 16,17 10 2 2 2 2 2 X28843 - 5901 PowerPlex® 21 - STRMix [™] V2.8.0 15,18 9,11,12,13 12,18 3sp 9,11 13,15 19,21 11,12 8,13,14 X28843 - 5901 PowerPlex® 21 - STRMix [™] V2.8.0 2 5,18 9,11,12,13 12,18 3sp 9,11 13,15 19,21 11,12 8,13,14 17,18,19 12,13 30,31.2,33 X,Y 10,11 20,23.2,24.2 9,10,13 5,8,10,12 6,9 8,10	Зsр	9,11	13,14**,15	13,14	19,21	11,12	8,13	
20,23,2,24.2** 15,28.2 6,9 8,10 16,17 10 2 WU8M28 - 5901 GlobalFiler [™] - STRMix [™] 2.8 11,14 15,18 9,11 3sp 9,11 13,15 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15 X,Y 10,11 20,23.2 16,17 10 2 2 X28843 - 5901 PowerPlex® 21- STRMix [™] V2.8.0 2 3 2 15 11,12,13 12,18 3sp 9,11 13,15 19,21 11,12,13 12,18 3sp 9,11 13,15 19,21 11,12 8,13,14 17,18,19 12,13 30,31.2,33 X,Y 10,11 20,23.2,24.2 9,10,13 5,8,10,12 6,9 8,10		17,18	12,13	30,31.2	15	X,Y	10,11	
16,17 10 2 WU8M28 - 5901 GlobalFiler [™] - STRMix [™] 2.8 14,18.3 20,25 11,14 15,18 9,11 3sp 9,11 13,15 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15 X,Y 10,11 20,23.2 15,18 9,11,12,13 8,10 16,17 10 2 X28843 - 5901 PowerPlex® 21- STRMix [™] V2.8.0 11,14,17.3,18.3 20,25 15,18 9,11,12,13 12,18 3sp 9,11 13,15 19,21 11,12 8,13,14 3sp 9,11 13,15 19,21 11,12 8,13,14 20,23.2,24.2 9,10,13 5,8,10,12 6,9 8,10		20,23.2,24.2**			15,28.2	6,9	8,10	
GlobalFiler™- STRMix™ 2.8 14,18.3 20,25 11,14 15,18 9,11 3sp 9,11 13,15 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15 X,Y 10,11 20,23.2 15,18 9,11,12,13 8,10 16,17 10 2 X28843 - 5901 PowerPlex® 21- STRMix™ V2.8.0 15,18 9,11,12,13 12,18 3sp 9,11 13,15 19,21 11,12 8,13,14 17,18,19 12,13 30,31.2,33 X,Y 10,11 20,23.2,24.2 9,10,13 5,8,10,12 6,9 8,10		16,17	10			2		
14,18.3 20,25 11,14 15,18 9,11 3sp 9,11 13,15 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15 X,Y 10,11 20,23.2 15,18 9,11,12,13 8,10 16,17 10 2 X28843 - 5901 PowerPlex® 21- STRMix™ V2.8.0 11,14,17.3,18.3 20,25 15,18 9,11,12,13 12,18 3sp 9,11 13,15 19,21 11,12 8,13,14 17,18,19 12,13 30,31.2,33 X,Y 10,11 20,23.2,24.2 9,10,13 5,8,10,12 6,9 8,10	WU8M28	8 - 5901 GI	obalFiler™- STRMix™ 2	2.8				
3sp 9,11 13,15 13,14 19,21 11,12 8,13 17,18 12,13 30,31.2 15 X,Y 10,11 20,23.2 15,28.2 6,9 8,10 16,17 10 2 X28843 - 5901 PowerPlex® 21- STRMix™ V2.8.0 15,18 9,11,12,13 12,18 3sp 9,11 13,15 19,21 11,12 8,13,14 17,18,19 12,13 30,31.2,33 X,Y 10,11 20,23.2,24.2 9,10,13 5,8,10,12 6,9 8,10		14,18.3	20,25	11,14	15,18	9,11		
17,18 12,13 30,31.2 15 X,Y 10,11 20,23.2 15,28.2 6,9 8,10 16,17 10 2 X28843 - 5901 PowerPlex® 21- STRMix™ V2.8.0 11,14,17.3,18.3 20,25 15,18 9,11,12,13 12,18 3sp 9,11 13,15 19,21 11,12 8,13,14 17,18,19 12,13 30,31.2,33 X,Y 10,11 20,23.2,24.2 9,10,13 5,8,10,12 6,9 8,10	Зsр	9,11	13,15	13,14	19,21	11,12	8,13	
20,23.2 15,28.2 6,9 8,10 16,17 10 2 X28843 - 5901 PowerPlex® 21- STRMix™ V2.8.0 11,14,17.3,18.3 20,25 15,18 9,11,12,13 12,18 3sp 9,11 13,15 19,21 11,12 8,13,14 17,18,19 12,13 30,31.2,33 X,Y 10,11 20,23.2,24.2 9,10,13 5,8,10,12 6,9 8,10		17,18	12,13	30,31.2	15	X,Y	10,11	
IO IO X28843 - 5901 PowerPlex® 21- STRMix™ V2.8.0 11,14,17.3,18.3 20,25 15,18 9,11,12,13 3sp 9,11 17,18,19 12,13 20,23.2,24.2 9,10,13 5,8,10,12 6,9		20,23.2	10		15,28.2	6,9	8,10	
X28843 - 5901 PowerPlex® 21- STRMix** V2.8.0 11,14,17.3,18.3 20,25 15,18 9,11,12,13 12,18 3sp 9,11 13,15 19,21 11,12 8,13,14 17,18,19 12,13 30,31.2,33 X,Y 10,11 20,23.2,24.2 9,10,13 5,8,10,12 6,9 8,10		10,17				Z		
11,14,17.3,18.3 20,25 15,18 9,11,12,13 12,18 3sp 9,11 13,15 19,21 11,12 8,13,14 17,18,19 12,13 30,31.2,33 X,Y 10,11 20,23.2,24.2 9,10,13 5,8,10,12 6,9 8,10	х28843	- 2901 Po	werrlex® 21- 51KMix™	v2.8.0	16.10	0.11.10.10	10.10	
35p 9,11 13,15 19,21 11,12 8,13,14 17,18,19 12,13 30,31.2,33 X,Y 10,11 20,23.2,24.2 9,10,13 5,8,10,12 6,9 8,10	0.	11,14,17.3,18.3	20,25		15,18	9,11,12,13	12,18	
17,10,17 12,13 30,31,2,33 A,1 10,11 20,23.2,24.2 9,10,13 5,8,10,12 6,9 8,10 14.17 14.17 14.17 14.17 14.17	Jsp	7,11	10.12	30 21 0 22	19,21	11,12	0,13,14	
14 17		20 23 2 24 2	0 10 12	581012		A, I	810	
		16.17	7,10,10	0,0,10,12		0,7	0,10	

			TABLE	3		
WebCoo	le - Test An	nplification Kits -	Probablistic Gene	otyping Software		
	D1S1656	D2S1338	D2S441	D3S1358	D5S818	D6S1043
Item	D7S820	D8S1179	D10S1248	D12S391	D13S317	D16S539
	D18S51	D195433	D21S11	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	TPOX
	vWA	DYS391	DY\$570	DYS576	Y Indel	
			ltem 3sp - STR	R Results		
YGJ6L9 -	5901 Glob	oalFiler™- STRMix™				
	13,14,17.3,18.3	19,20,24,25	10,11,13,14	14,15,17,18,19	8,9,10,11	
Зsр	8,9,10,11	12,13,14,15	12,13,14	18,19,20,21	10,11,12	7,8,12,13
	16,17,18	11,12,13	29,30,30.2,31.2	14,15,16	X,Y	9,10,11
	19,20,22.2,23.2			14,15,27.2,28.2	6,8,9	8,10
	15,16,17	10			2	
YJMUQ3	- 5901 Pow	erPlex® Fusion 6C-	STRMix™ V2.5.11			
	11,14,16,17.3,18.3	17,18,20,25	11,14	15,16,18	9,10,11,12,13	
Зsр	9,10,11,12	10,13,15	13,14	16,17,18,19,21	8,11,12	8,9,12,13,14
	14,15,17,18,19	12,13,15.2	30,31.2,32.2,33,33 .2	11,12,14,15	Х,Ү	10,11,12
	20,21,23,23.2,24	9,10,11,12,13	5,8,10,11,12	15,17,18,24.2,28.2	6,7,9,9.3	8,10,11,14
	15,16,17	10	18,20	17,19		
Z8LUQ4	- 5902 Gloł	oalFiler™- TrueAllele	e® VUler Release 202	2b		
	14,18.3	17,18,20,25	11,14	15,18	9,11,13	
Зsр	9,11,12	10,13,15	13,14	19,21	11,12	8,9,12,13
	17,18	12,13	30,31.2	11,15	X,Y	10,11
	20,23.2			15,28.2	6,7,9	8,10
	16,17	10			2	

			TABLE	3		
WebCo	de - Test 👘	Amplification Kits -	Probablistic Gene	otyping Software		
	D1\$1656	D251338	D25441	D3\$1358	D55818	D651043
ltem	D7S820	D8S1179	D10S1248	D12S391	D135317	D16S539
	D18551	D195433	D21511	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	VWA	D12241		Dission	Tindei	
20GMC	·2 - 5902 -	JobalFiler™- TrueAllele	R "VI ller Release 20	22b		
200110	11.16.17.3	17.18.23.24.25	10.14	14.15.16	11.12.13	
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26			14,18,23.2,26.2,28	7,9.3	8,11,14
	15,16,17	10,11			2	
34G3B3	- 5902 G	GlobalFiler™- STRMix™ 2	2.8			
	11,15,15.2,16,16. 17.1,17.3	3, 16,17,18,22,23,24, 25	9,10,13,14,15	13,14,15,16	10,11,12,13	
4	7,8,9,10,11,12	9,10,12,13,14,15,1 6	12,13,14,15	17,18,19,20,21	8,9,10,11,12,13	8,9,10,11,12,13,1 4
	13,14,15,16,17,18 19	8, 11,12,12.2,13,13.2 ,14,14.2,15.2	28,29,30,32,32.2,3 3,33.2	11,13,14,15,16,17	X,Y	9,10,11,12,13
	19,20,21,22,23,23 26,27	5,		13,13.2,14,17,17.2 ,18,22.2,23.2,25.2, 26.2,27.2,28,28.2	6,7,9.3	8,10,11,13,14
	14,15,16,17	9,10,11			2	
37J2CU	- 5901 G	GlobalFiler™- STRMix™				
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26			14,18,23.2,26.2,28	7,9.3	8,11,14
	15,16,17	10,11		.2	2	
3MQZU	Y - 5902 ld	lentifiler® Plus- STRMix ⁺	TM			
		17,18,23,24,25		14,15,16	11,12,13	
4	8,9,11,12	10,13,14,16			8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2		X,Y	10,11,12,13
	20,21,23,26				7,9.3	8,11,14
	15,16,17					
3WHFJ3	- 5902 G	GlobalFiler™- STRMix™ \	/2.9.1			
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26			14,18,23.2,26.2,28	7,9.3	8,11,14
	15,16,17	10,11		.2	2	
44PDRD	- 5901 G	GlobalFiler™- STRMix™				
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26			14,18,23.2,26.2,28 .2	7,9.3	8,11,14
	15,16,17	10,11			2	
			TABLE	3		
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WebCo	de - Test 🛛 An	nplification Kits -	Probablistic Gene	otyping Software		
	D1\$1656	D2\$1338	D2S441	D3\$1358	D5S818	D6S1043
ltem	D7\$820	D8S1179	D1051248	D12\$391	D13S317	D16S539
	D18S51	D195433	D21511	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	VWA	D13391		Dission	rindei	
4BEUJW	- 5902 Pow	rerPlex® 21- STRMix⁺	[™] 2.10.0	Kesuits		
	11,16,17.3	17,18,23,24,25		14,15,16	11,12,13	10,11,12,13,14
4	8,9,11,12	10,13,14,16		18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2		X,Y	10,11,12,13
	20,21,23,26	9,10,11,12,15	5,10,11,12,14		7,9.3	8,11,14
	15,16,17					
4FBJY4 -	5902 Pow	erPlex® Fusion 6C -	DNAxs			
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26	9,10,11,12,15	5,10,11,12,14	14,18,23.2,26.2,28 .2	7,9.3	8,11,14
	15,16,17	10,11	17,20	16,17,21		
4UQG64	4 - 5901 Glo	balFiler™- STRMix™				
	11,15,16,17.3	16,17,18,23,24,25	9,10,13,14	13,14,15,16	10,11,12,13	
4	8,9,11,12	9,10,12,13,14,16	12,13,14,15	17,18,19,20,21	8,10,11,12,13	9,11,12,13,14
	13,14,15,17,18,19	12,12.2,13,13.2,14 ,14.2,15.2	28,29,30,32,32.2,3 3,33.2	11,13,14,15,16	Х,Ү	10,11,12,13
	19,20,21,22,23,25, 26			14,17,18,23.2,25.2 ,26.2.27.2.28.2	6,7,9.3	8,11,14
	15,16,17	10,11			2	
4XQRZ4	- 5901 Glo	balFiler™- STRMix™				
	11,16,17.3	16,17,18,23,24,25	9,10,13,14	14,15,16	11,12,13	
4	8,9,11,12	10,12,13,14,16	12,13,14,15	18,19,21	8,10,11,12,13	9,11,12,13,14
	13,15,17,18,19	12,12.2,13,13.2,14	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	19,20,21,22,23,25,			14,17,18,23.2,26.2	6,7,9.3	8,11,14
	15,16,17	10,11		,20.2	2	
62K9Q4	- 5901 Glo	balFiler™- STRMix™				
	10,11,15,16,16.3,17	16,17,18,23,24,25	9,10,13,14	14,15,16	10,11,12,13	
4	8,9,10,11,12	9,10,12,13,14,15,1	12,13,14,15	17,18,19,21	8,10,11,12,13	9,11,12,13,14
	13,14,15,17,18,19	11,12,13,13.2,14,1	28,29,30,32,32.2,3	11,13,14,15,16	X,Y	9,10,11,12,13
	19,20,21,22,23,25,	4.2,13.2	0,00.2	14,17,18,23.2,26.2	6,7,9.3	8,10,11,14
	14,15,16,17	10,11		,27.2,20.2	2	
63VBC2	- 5901 Glo	balFiler™- STRMix™				
	11,15,16,16.3,17.3	16,17,18,23,24,25	9,10,13,14	14,15,16	10,11,12,13	
4	8,9,11,12	9,10,12,13,14,16	12,13,14,15	17,18,19,20,21	8,10,11,12,13	9,11,12,13,14
	13,14,15,16,17,18, 19	12,12.2,13,13.2,14	28,29,30,32,33,33.	11,13,14,15,16	X,Y	9,10,11,12,13
	20,21,22,23,25,26	,		14,17,18,23.2,26.2	6,7,9.3	8,10,11,14
	15,16,17	10,11		,,	2	

			TABLE	3		
WebCo	de - Test 🛛 🗛	mplification Kits -	Probablistic Gene	otyping Software		
	D1S1656	D2S1338	D2S441	D3\$1358	D5\$818	D6S1043
ltem	D7\$820	D8S1179	D10S1248	D12S391	D135317	D16S539
	D18551	D19S433	D21S11	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	VWA	D12391		DTS570	t Indel	
ζρτμζν	5002 CL	obalEilor™ TruoAllolo(Item 4 - SIK			
0111101	11 16 17 3			17 15 16	11 12 13	
4	891112	10 13 14 16	13 14 15	18 19 21	8 10 12 13	9 11 12 13 14
-	13 15 17 19	12 13 13 2 14 15 2	28 29 30 33 33 2	11 14 15 16	X Y	10 11 12 13
	20.21.23.26	12,10,10.2,14,10.2	20,27,00,00,00.2	14.18.23 2.26 2.28	7.9.3	8.11.14
	20/2:/20/20			.2	.,,	0,,
	15,16,17	10,11			2	
77A6RY	- 5901 Gl	obalFiler™- STRMix™				
	11,15,16,16.3,17.3	3 16,17,18,22,23,24, 25	9,10,13,14	13,14,15,16	10,11,12,13	
4	8,9,10,11,12	9,10,12,13,14,16	12,13,14,15	17,18,19,20,21	8,10,11,12,13	9,10,11,12,13,14
	13,14,15,16,17,18, 19	, 11,12,12.2,13,13.2 ,14,14.2,15.2	28,29,30,32,32.2,3 3,33.2	11,13,14,15,16	X,Y	9,10,11,12,13
	19,20,21,22,23,25, 26	,		14,17,17.2,18,22.2 ,23.2,25.2,26.2,27. 2,28,28.2	6,7,9.3	8,10,11,13,14
	14,15,16,17	9,10,11			2	
786NUZ	- 5901 Ide	entifiler® Plus - EruoFo	rMix			
		17,18,23,24,25		14,15,16	11,12,13	
4	8,9,11,12	10,13,14,16			8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2		X,Y	10,11,12,13
	20,21,23,26				7,9.3	8,11,14
	15,16,17					
7H4JRV -	- 5902 Gl	obalFiler™- TrueAllele	® VUIer Release 202	2b		
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26			14,18,23.2,26.2,28	7,9.3	8,11,14
	15,16,17	10,11		.2	2	
8Y8DYV	- 5902 Gl	obalFiler™- TrueAllele	® VUIer Release 202	2b		
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26			14,18,23.2,26.2,28	7,9.3	8,11,14
	151(17	10.11		.2		
	5, 6, /	10,11			2	
8YUL9Z	- 5901 Po	werPlex® Fusion 6C, N	NGM Detect - LRmix			
4	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	0 11 10 10 14
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	0 10 11 10 15	28,29,30,33,33.2	11,14,15,16	Χ,Υ	10,11,12,13
	20,21,23,20	9,10,11,12,15	5,10,11,12,14	.2	7,7.3	0,11,14
	15,16,17	10,11	17,20	16,17,21	2	

TABLE 3							
WebCoo	de - Test 🛛 🗛 An	nplification Kits -	Probablistic Gen	otyping Software			
	D151656	D251338	D25441	D3\$1358	D55818	D651043	
ltem	D75820	D8S1179	D10S1248	D12S391	D135317	D165539	
	D18551	D195433	D21S11	D22S1045	Amelogenin	CSF1PO	
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ	
	vWA	DYS391	DYS570	DYS576	Y Indel		
	5001 011		Item 4 - STR	Results			
9CLYK9 -	- 5901 Gloi	balFiler''' - STRMix'''					
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13		
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14	
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13	
	20,21,23,26			14,18,23.2,26.2,28	7,9.3	8,11,14	
	15,16,17	10,11			2		
9MBR8Q	- 5901 Pow	erPlex® 21- STRMix⁺	™ 2.8.0				
	11,16,17.3	17,18,23,24,25		14,15,16	11,12,13	10,11,12,13,14	
4	8,9,11,12	10,13,14,16		18,19,21	8,10,12,13	9,11,12,13,14	
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2		X,Y	10,11,12,13	
	20,21,23,26	9,10,11,12,15	5,10,11,12,14		7,9.3	8,11,14	
	15,16,17						
9WLTTR -	- 5902 Pow	erPlex® 21- STRMix⁺	[™] 2.10.0				
	11,16,17.3	17,18,23,24,25		14,15,16	11,12,13	10,11,12,13,14	
4	8,9,11,12	10,13,14,16		18,19,21	8,10,12,13	9,11,12,13,14	
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2		X,Y	10,11,12,13	
	20,21,23,26	9,10,11,12,15	5,10,11,12,14		7,9.3	8,11,14	
	15,16,17						
B7W38V	- 5901 Glol	balFiler™- STRMix™					
	11,15,16,16.3,17.3	16,17,18,23,24,25	10,13,14	14,15,16	10,11,12,13		
4	8.9.11.12	9.10.12.13.14.16	12,13,14,15	17,18,19,20,21	8.10.11.12.13	9,11,12,13,14	
	13,14,15,17,18,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,13,14,15,16	X,Y	10,11,12,13	
	20,21,23,25,26			14,17,18,23.2,26.2	7,9.3	8,11,14	
				,27.2,28.2			
	15,16,17	9,10,11			2		
BDR7Z7	- 5901 Pow	erPlex® Fusion 6C- S	STRMix™ 2.6.2				
	(11),16,17.3	17,18,(24),(25)	(10),14	(14),(15),16	11,(12),13		
4	(8),9,11,(12)	10,13,(14),(16)	13,(14),(15)	18,19,(21)	(8),(10),12,(13)	9,(11),(12),(13),14	
	(13),15,(17),19	12,(13),(13.2),(14),1	(28),(29),(30),33,33.	(11),14,15,(16)	X,Y	10,(11),12,(13)	
	20,(21),23,(26)	9,(10),(11),12,(15)	5,10,(11),(12),(14)	(14),18,(23.2),(26.2) ,28.2	7,(9.3)	(8),11,(13),14	
	15,(16),17	10,(11)	(17),20	(16),17,(21)			
BKHHBU	- 5901 Glol	balFiler™- STRMix™					
	11,15,16,16.3,17.3	16,17,18,22,23,24, 25	9,10,13,14	13,14,15,16	10,11,12,13		
4	7,8,9,10,11,12	9,10,12,13,14,16	12,13,14,15	17,18,19,20,21	8,10,11,12,13	9,10,11,12,13,14	
	13,14,15,16,17,18, 19	11,12,12.2,13,13.2 ,14,14.2,15.2	28,29,30,32,32.2,3 3,33.2	11,13,14,15,16	X,Y	9,10,11,12,13	
	19,20,21,22,23,25, 26		·	13,13.2,14,17,18,2 3.2,25.2,26.2,27.2, 28.2	6,7,9.3	8,10,11,14	
	14,15,16,17	9,10,11			2		

			TABLE	3		
WebCod	de - Test 🛛 🗛 An	nplification Kits -	Probablistic Gene	otyping Software		
	D1\$1656	D251338	D25441	D3\$1358	D5\$818	D651043
ltem	D7\$820	D851179	D1051248	D125391	D135317	D165539
	D18551	D195433	D21S11	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	vWA	DY\$391	DY\$570	DYS576	Y Indel	
DUCTOU	5000 011		Item 4 - STR	Results		
BNG180	- 5902 Glol	palFiler ** - STRMix ** 2	2.8			
	11,15,15.2,16,16.3, 17.1,17.3	16,17,18,22,23,24, 25	9,10,13,14	13,14,15,16	10,11,12,13,14	
4	7,8,9,10,11,12	9,10,12,13,14,15,1 6	12,13,14,15	17,18,19,20,21	8,9,10,11,12,13	8,9,10,11,12,13,1 4
	13,14,15,16,17,18, 19	11,12,12.2,13,13.2 ,14,14.2,15.2	27,28,29,30,32,32. 2,33,33.2	11,13,14,15,16,17	X,Y	9,10,11,12,13
	19,20,21,22,23,24, 25,26			13,14,17,17.2,18,2 2.2,23.2,25.2,26.2, 27.2,28,28.2	6,7,9.3	8,10,11,13,14
	14,15,16,17	9,10,11			2	
CDTG9N	1 - 5901 Glol	oalFiler™				
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	-
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26	-	-	14,18,23.2,26.2,28	7,9.3	8,11,14
	15,16,17	10,11	-	-	2	
CKECAQ	- 5902 Glol	oalFiler™- TrueAllele	® VUIer Release 202	22b		
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26			14,18,23.2,26.2,28	7,9.3	8,11,14
	15 16 17	10 11		.2	2	
	L- 5901 Pow	erPlex® FSI-17 Fast.	LiRa v3 0			
	11 16 17 3	17 18 23 24 25	10.14	14 15 16		
4	11,10,17.0	10 13 14 16	13 14 15	18 19 21		9 11 12 13 14
-	13 15 17 19	12 13 13 2 14 15	28 29 30 33 33 2	11 14 15 16	XY	7,11,12,10,14
	20,21,23,26	12,10,10.2,14,10	20,27,00,00,00.2	14.18.23.2.26.2.28	7,9.3	
	, , ,			.2		
	15,16,17					
CRVQCV	V - 5901 Glol	oalFiler™- STRMix™				
	11,15,16,16.3,17.3	17,18,23,24,25	9,10,13,14	13,14,15,16	10,11,12,13	
4	8,9,10,11,12	9,10,12,13,14,16	12,13,14,15	18,19,21	8,10,11,12,13	9,11,12,13,14
	13,14,15,17,18,19	12,12.2,13,13.2,14 ,14.2,15.2	28,29,30,32.2,33,3 3.2	11,14,15,16	X,Y	10,11,12,13
	19,20,21,22,23,25, 26			14,17,18,23.2,26.2 ,27.2,28.2	6,7,9.3	8,11,14
	15,16,17	10,11			2	
CXYRCQ	- 5902 Glo	oalFiler™- TrueAllele	® VUIer Release 202	22b		
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26			14,18,23.2,26.2,28 .2	7,9.3	8,11,14
	15,16,17	10,11			2	

			TABLE	3		
WebCo	de - Test 🛛 🖉	Amplification Kits -	Probablistic Gen	otyping Software		
	D1S1656	D2S1338	D2S441	D3S1358	D5\$818	D6S1043
ltem	D75820	D851179	D1051248	D125391	D135317	D16S539
	D18S51	D195433	D21S11	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	VWA	D123A1			t Indei	
СУКПРК	- 5901	ilobalEiler™- STRMix™	ITEM 4 - SIK	Kesuits		
CIODIO	11 16 17 3	17 18 23 24 25	10 14	14 15 16	11 12 13	
4	8.9.11.12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9.11.12.13.14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26			14,18,23.2,26.2,28	7,9.3	8,11,14
				.2		
	15,16,17	10,11			2	
CZ6EPV	- 5901 G	ilobalFiler™- STRMix™				
	11,15,16,16.3,17	.3 16,17,18,22,23,24, 25	9,10,13,14,15	13,14,15,16	10,11,12,13	
4	8,9,10,11,12	9,10,12,13,14,16	12,13,14,15	17,18,19,21	8,9,10,11,12,13	9,10,11,12,13,14
	13,14,15,17,18,1	9 11,12,13,13.2,14,1 4.2,15.2	28,29,30,32,32.2,3 3,33.2	11,13,14,15,16	X,Y	9,10,11,12,13
	19,20,21,22,23,25 26	5,	ŕ	13,14,17,18,22.2,2 3.2,25.2,26.2,27.2, 28.2	6,7,9.3	8,10,11,14
	14,15,16,17	10,11			2	
E34ANN	- 5902 G	iobalFiler™- TrueAllele	® VUIer Release 202	22b		
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26			14,18,23.2,26.2,28 .2	7,9.3	8,11,14
	15,16,17	10,11			2	
EHXZJK -	- 5902 Po	owerPlex® 21- STRMix⁺	₫ 2.10			
	11,16,17.3	17,18,23,24,25		14,15,16	11,12,13	10,11,12,13,14
4	8,9,11,12	10,13,14,16		18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2		X,Y	10,11,12,13
	20,23,26	9,10,11,12,15	5,10,11,12,14		7,9.3	8,11,14
	15,16,17					
ELJJPP - 🗄	5902 G	lobalFiler™- TrueAllele	® VUIer Release 202	22b		
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26			14,18,23.2,26.2,28	7,9.3	8,11,14
	15,16,17	10,11		.2	2	
EY6YTN	- 5902 G	ilobalFiler™- TrueAllele	® VUler Release 202	22b		
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26			14,18,23.2,26.2,28 .2	7,9.3	8,11,14
	15,16,17	10,11			2	

			TABLE	3		
WebCod	e - Test	Amplification Kits -	Probablistic Gen	otyping Software		
	D1\$1656	D251338	D25441	D3\$1358	D5\$818	D651043
Item	D7S820	D8S1179	D1051248	D125391	D135317	D16S539
	D18S51	D195433	D21S11	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	TPOX
	vWA	DYS391	DYS570	DYS576	Y Indel	
	5000		Item 4 - STR	Results		
F/H8UP -	3902		2.9.1		11 10 10	
4	0.011.10	17,18,23,24,25	10,14	14,15,10	11,12,13	0 11 10 12 14
4	0,9,11,12	10,13,14,10	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	20 21 23 24	12,13,13.2,14,15.2	26,29,30,33,33.2	14,19,03,0,06,0,09	7.0.2	9 11 14
	20,21,23,20			.2	7,9.3	0,11,14
	15,16,17	10,11			2	
F8XQDM ·	- 5902	GlobalFiler™- TrueAllele	® VUIer Release 202	22b		
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26			14,18,23.2,26.2,28	7,9.3	8,11,14
	15 16 17	10.11		.2	2	
EPXEAL 5	3002	Identifiler® Plus STRMix	м		L	
TTALOL - J	702	17 18 23 24 25		14 15 16	11 12 13	
4	8.9.11.12	10.13.14.16		14,10,10	8.10.12.13	9.11.12.13.14
	13,15,17,19	12,13,13,2,14,15,2	28.29.30.33.33.2		X,Y	10,11,12,13
	20,21,23,26	· · · · · · · · · · · ·			7,9.3	8,11,14
	15,16,17				,	
GCH7HK	- 5902	GlobalFiler™- TrueAllele	® VUIer Release 202	22b		
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26			14,18,23.2,26.2,28	7,9.3	8,11,14
				.2		
	15,16,17	10,11			2	
H484WJ -	5902	GlobalFiler [™] - TrueAllele	® VUIer Release 202	22b		
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26			14,18,23.2,26.2,28	7,9.3	8,11,14
	15,16,17	10,11			2	
HEQHYL -	5902	GlobalFiler™- TrueAllele	® VUIer Release 202	22b		
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26			14,18,23.2,26.2,28	7,9.3	8,11,14
	15 16 17	10.11		.2	2	
	13,10,17	10,11			2	

			TABLE	3		
WebCoo	de - Test 🛛 🖌	Amplification Kits -	Probablistic Gen	otyping Software		
	D1\$1656	D2S1338	D2S441	D3S1358	D5\$818	D6S1043
ltem	D75820	D8S1179	D10S1248	D12S391	D135317	D165539
	D18551	D195433	D21511	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	VWA	D¥5391	DYS570	DY5576	¥ Indel	
	5901	lobalFilor™ IOC	Item 4 - SIR	Results		
	(11) 14 17 3	17 18 (23 24 25)	(10) 14	(14) 15 16	11/10/13	
1	8 9 11 (12)	10 13 (14 16)	13 (14 15)	18 (19 21)	(8 10) 12 (13)	011/1013\14
-	(13) 15 (17) 19	12 13 13 2 (14) 15	(28) 29 (30 32) 33	(11) 14 15 (16)	(0,10),12,(10) X Y	10 11 12 (13)
	(10),10,(17),17	2	33.2	(11),14,13,(10)	Λ, Ι	10,11,12,(10)
	20,(21),23,26			(14),18,(23.2,26.2), 28.2	7,(9.3)	(8),11,14
	15,16,17	10,(11)			2	
J37QNH	- 5902 Po	owerPlex® 21- STRMix⊺	™ 2.10.0			
	11,16,17.3	17,18,23,24,25		14,15,16	11,12,13	10,11,12,13,14
4	8,9,11,12	10,13,14,16		18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2		X,Y	10,11,12,13
	20,21,23,26	9,10,11,12,15	5,10,11,12,14		7,9.3	8,11,13,14
	15,16,17					
J772JG -	5902 Po	owerPlex® 21- STRMix⁺	™ 2.10			
	11,16,17.3	17,18,23,24,25		14,15,16	11,12,13	10,11,12,13,14
4	8,9,11,12	10,13,14,16		18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2		X,Y	10,11,12,13
	20,21,23,26	9,10,11,12,15	5,10,11,12,14		7,9.3	8,11,14
	15,16,17					
JHLQHM	- 5902 G	lobalFiler™ - DNAxs				
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26			14,18,23.2,26.2,28	7,9.3	8,11,14
	15,16,17	10,11		۷.	2	
JPGUBX -	- 5901 G	lobalFiler™- STRMix™				
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26			14,18,23.2,26.2,27	7,9.3	8,11,14
	15,16,17	10,11		.∠,∠0.∠	2	
K9ECCE	- 5902 Pa	owerPlex® 21- STRMix™	[™] 2.10			
	11,16,17.3	17,18,23,24,25		14,15,16	11,12,13	10,11,12,13,14
4	8,9,11,12	10,13,14,16		18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2		X,Y	10,11,12,13
	20,21,23,26	9,10,11,12,15	5,10,11,12,14		7,9.3	8,11,14
	15,16,17					

			TABLE	3		
WebCo	de - Test 🛛 🗛	mplification Kits -	Probablistic Gen	otyping Software		
	D151656	· •	D26441	D261250	D50010	D461042
Item	D75820	D231338	D1051248	D125391	D135317	D165539
	D18S51	D19S433	D21S11	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	vWA	DYS391	DYS570	DYS576	Y Indel	
			ltem 4 - STR	Results		
KPDQAK	(- 5902 Glo	balFiler™- STRMix™ v	/2.9.1			
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26			14,18,23.2,26.2,28 .2	7,9.3	8,11,14
	15,16,17	10,11			2	
L67JVL -	5901 Glo	balFiler™- STRMix™				
	11,16,16.3,17.3	16,17,18,23,24,25	9,10,13,14	14,15,16	11,12,13	
4	8,9,11,12	9,10,12,13,14,16	12,13,14,15	18,19,21	8,10,11,12,13	9,11,12,13,14
	13,14,15,17,18,19	11,12,13,13.2,14,1 4 2 15 2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	19,20,21,22,23,24, 25,26	7.2,10.2		14,17,18,23.2,26.2 ,27.2,28.2	7,9.3	8,11,14
	15,16,17	10,11			2	
L8XEJE -	5901 Pow	verPlex® 21- STRMix™	″ v 2.8.0			
	11,16,17.3	17,18,23,24,25		14,15,16	11,12,13	10,11,12,13,14
4	8,9,11,12	10,13,14,16		18,19,21	8,10,12,13	9,11,12,13,14,15
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2		X,Y	10,11,12,13
	20,21,23,26	9,10,11,12,15	5,10,11,12,14		7,9.3	8,11,14
	15,16,17					
LBAKVF	- 5902 Ider	ntifiler® Plus- STRMix ⁺	™ 2.7.0			
		17,18,23,24,25		14,15,16	11,12,13	
4	8,9,11,12	10,13,14,16			8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2		X,Y	10,11,12,13
	20,21,23,26				7,9.3	8,11,14
	15,16,17					
lg3pnj	- 5901 Pow	verPlex® ESX17 Fast S	öystem- STRMix™ V2	.5.11		
	11,16,17.3	17,18,23,24,25	10,14	14,15,16		
4		10,13,14,16	13,14,15	18,19,21		9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	
	20,21,23,26			14,18,23.2,26.2,28	7,9.3	
	15,16,17			.2		
LHUN3H	1 - 5902 Glo	balFiler™- TrueAllele	® VUIer Release 202	22b		
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26			14,18,23.2,26.2,28	7,9.3	8,11,14
	15,16,17	10,11		.2	2	

			TABLE	3				
WebCo	de - Test 🛛 🗛	mplification Kits -	Probablistic Gen	otyping Software				
	D1\$1656	D251338	D25441	D3\$1358	D55818	D651043		
ltem	D7S820	D8S1179	D1051248	D12S391	D13S317	D16S539		
	D18551	D195433	D21S11	D22S1045	Amelogenin	CSF1PO		
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ		
	vWA	DYS391	DYS570	DYS576	Y Indel			
	5001 D							
LUPUVVF	- 3901 POV		$\frac{10.14}{10.14}$	J. I I	11 10 10			
1	901112	10 13 14 14	10,14	14,15,10	9 10 12 13	0 11 12 13 14		
4	0,7,11,12	10,13,14,10	13,14,13	10,19,21	0,10,12,13	9,11,12,13,14		
	13,13,17,17	12,13,13.2,14,13.2	28,27,30,32,33,33.	11,14,13,10	Λ, Ι	10,11,12,13		
	20,21,23,26	9,10,11,12,14,15	5,10,11,12,14	14,18,23.2,26.2,28 .2	7,9.3	8,11,14		
	15,16,17	10,11	17,20	16,17,21				
LYKQJL -	5901 Glo	obalFiler™- STRMix™						
	11,15,16,16.3,17.3	8 16,17,18,23,24,25	9,10,13,14	13,14,15,16	10,11,12,13			
4	8,9,10,11,12	9,10,12,13,14,16	12,13,14,15	17,18,19,20,21	8,10,11,12,13	9,11,12,13,14		
	13,14,15,17,18,19	11,12,12.2,13,13.2 ,14,14.2,15.2	28,29,30,32,32.2,3 3,33.2	11,13,14,15,16	X,Y	9,10,11,12,13		
	19,20,21,22,23,25, 26			13,14,17,18,23.2,2 5.2,26.2,27.2,28.2	6,7,9.3	8,10,11,14		
	15,16,17	10,11			2			
MW99ZF	MW99ZF - 5902 GlobalFiler™- TrueAllele® VUIer Release 2022b							
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13			
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14		
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13		
	20,21,23,26			14,18,23.2,26.2,28 .2	7,9.3	8,11,14		
	15,16,17	10,11			2			
NWNRK	D - 5901 Pov	werPlex® Fusion 6C- 3	STRMix™ V2.5.11					
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13			
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14		
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13		
	20,21,23,26	9,10,11,12,15	5,10,11,12,14	14,18,23.2,26.2,28 .2	7,9.3	8,11,14		
	15,16,17	10,11	17,20	16,17,21				
P3QH3A	- 5901 Pov	werPlex® 21- STRMix⊺	™ 2.8					
	11,16,17.3	17,18,23,24,25		14,15,16	11,12,13	10,11,12,13,14		
4	8,9,11,12	10,13,14,16		18,19,21	8,10,12,13	9,11,12,13,14		
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2		X,Y	10,11,12,13		
	20,21,23,26	9,10,11,12,15	5,10,11,12,14		7,9.3	8,11,14		
	15,16,17							
P8KEPF -	5902 Glo	obalFiler™- STRMix™						
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13			
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14		
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13		
	20,21,23,26			14,18,23.2,26.2,28 .2	7,9.3	8,11,14		
	15,16,17	10,11			2			

			TABLE	3		
WebCoo	de - Test 🛛 🗛	mplification Kits -	Probablistic Gene	otyping Software		
	D1\$1656	D2S1338	D2S441	D3S1358	D5\$818	D6S1043
ltem	D75820	D8S1179	D1051248	D125391	D13S317	D16S539
	D18S51	D195433	D21511	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	VWA	D12241		Dission	rindei	
	- 5901 Gla	balFiler™- STRMix™	ITEM 4 - 31K	Kesuits		
10117Q	11 16 17 3	17 18 23 24 25	10.14	14 15 16	11 12 13	
4	8 9 11 12	10 13 14 16	13 14 15	18 19 21	8 10 12 13	9 11 12 13 14
	13.15.17.19	12.13.13 2.14.15 2	28.29.30.33.33.2	11.14.15.16	X.Y	10.11.12.13
	20,21,23,26	/ . 0 / . 0 / / . 0	20/2//00/00/00/2012	14,18,23.2,26.2,28	7,9.3	8,11,14
	, , ,			.2		, ,
	15,16,17	10,11			2	
Q99BQE	- 5901 Glo	balFiler™ IQC- STRM	1ix™ 2.8.0			
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26			14,18,23.2,26.2,28	7,9.3	8,11,14
	15,16,17	10,11		.2	2	
R3MFPB	- 5901 Pov	werPlex® Fusion 6C- 3	STRMix™ STRmix v2.5	5.11		
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26	9,10,11,12,15	5,10,11,12,14	14,18,23.2,26.2,28	7,9.3	8,11,14
	15 14 17	10.11	17 10 20	.2		
	5001		17,17,20 CTD :	10,17,21		
KYKDIE -	- 5901 Gio			141514	10 11 10 12	
4	11,15,16,16.3,17.3	0 10 10 12 14 14	9,10,13,14	14,15,16	10,11,12,13	0 11 10 10 14
4	0,9,10,11,12	9,10,12,13,14,10	12,13,14,15	11,10,19,20,21	0,10,11,12,13	9,11,12,13,14
	13,14,13,17,10,17	,15.2	3,33.2	11,14,13,10	Λ, Ι	10,11,12,13
	20,21,22,23,25,26			14,17,18,23.2,26.2	7,9.3	8,10,11,14
	15 16 17	10.11		,27.2,28.2	2	
	5901 lpv	octigator® 24 play OS	EuroEarMix		L	
KCI AZD	[11] 16 17 3	17 [18 23 24 25]	10.14	14 15 16	11 12 13	
1	8 9 11 12	10 13 [14 16]	13 [14 15]	18 [19] 21	8 10 12 13	011121314
-	[13] 15 [17] 19	12 13 13 2 14 15 2	28 29 30 33 33 2	11 14 15 16	X Y	10 11 12 13
	20.[21].23.26	12,10,10.2,11,10.2	20,27,00,00,00.2	14.18.23.2.26.2.28	7.9.3	[8].11.[14]
	20/[2:]/20/20			.2	.,,	[0]//[]
	15,16,17	10,[11]				
TMDEV4	- 5902 Glo	obalFiler™				
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	Х,Ү	10,11,12,13
	20,21,23,26			14,18,23.2,26.2,28	7,9.3	8,11,14
	15,16,17	10,11		.2	2	

			TABLE	3		
WebCod	e - Test	Amplification Kits -	Probablistic Gen	otyping Software		
	D1S1656	D2S1338	D2S441	D3S1358	D5S818	D6S1043
ltem	D7S820	D8\$1179	D10S1248	D12S391	D13S317	D16S539
	D18551	D195433	D21511	D22S1045	Amelogenin	CSF1PO
	vWA	DYS391	DYS570	DYS576	Yindel	IFOX
			ltem 4 - STR	Results		
TZBZVA -	5901	PowerPlex® Fusion 6C- S	5TRMix™ V2.5.11	Kesens		
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26	9,10,11,12,15	5,10,11,12,14	14,18,23.2,26.2,28 .2	7,9.3	8,11,14
	15,16,17	10,11	17,20	16,17,21		
UNVR99	- 5902	GlobalFiler™- TrueAllele	® VUIer Release 202	22b		
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26			14,18,23.2,26.2,28 .2	7,9.3	8,11,14
	15,16,17	10,11			2	
VFHZHA	- 5902	GlobalFiler™- STRMix™ \	/2.9.1			
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26			14,18,23.2,26.2,28	7,9.3	8,11,14
	15,16,17	10,11			2	
VK9834 -	5902	PowerPlex® Fusion 6C- S	STRMix™ v2.10			
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4	8,9,11,12	10,12,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,12.2,13,13.2,14 ,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26	9,10,11,12,15	5,10,11,12,14	14,18,23.2,26.2,28 .2	7,9.3	8,11,14
	15,16,17	10,11	17,20	16,17,21		
W2RLKZ -	- 5901	GlobalFiler™- STRMix™				
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,32,33,33. 2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26			14,18,23.2,26.2,28 .2	7,9.3	8,11,14
	15,16,17	10,11			2	
WRGKYZ	- 5901	GlobalFiler™- STRMix™				
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26			14,18,23.2,26.2,28 .2	7,9.3	8,11,14
	15,16,17	10,11			2	

	TABLE 3							
WebCoo	le - Test	Amplification Kits -	Probablistic Gen	otyping Software				
	D1S1656	D2S1338	D2S441	D3S1358	D5S818	D6S1043		
ltem	D7S820	D8S1179	D10S1248	D12S391	D13S317	D16S539		
	D18551	D195433	D21S11	D22S1045	Amelogenin	CSF1PO		
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ		
	vWA	DYS391	DYS570	DYS576	Y Indel			
			ltem 4 - STR	Results				
X28843 -	5901	PowerPlex® 21- STRMix™	⁴ V2.8.0					
	11,16,17.3	3 17,18,23,24,25		14,15,16	11,12,13	10,11,12,13,14		
4	8,9,11,12	10,13,14,16		18,19,21	8,10,12,13	9,11,12,13,14		
	13,15,17,1	9 12,13,13.2,14,15.2	28,29,30,33,33.2		X,Y	10,11,12,13		
	20,21,23,2	6 9,10,11,12,15	5,10,11,12,14		7,9.3	8,11,14		
	15,16,17							
YGJ6L9 -	5901	GlobalFiler™- STRMix™						
	11,15,16,17	7.3 17,18,23,24,25	9,10,13,14	14,15,16	10,11,12,13			
4	8,9,11,12	10,12,13,14,16	12,13,14,15	18,19,21	8,10,11,12,13	9,11,12,13,14		
	13,15,17,18,	19 11,12,12.2,13,13.2 ,14,14.2,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13		
	20,21,22,23,2	5,26		14,17,18,23.2,26.2 ,27.2,28.2	6,7,9.3	8,11,14		
	15,16,17	10,11			2			
YJMUQ3	- 5901	PowerPlex® Fusion 6C- S	5TRMix™ V2.5.11					
	11,16,17.3	3 17,18,23,24,25	10,14	14,15,16	11,12,13			
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14		
	13,15,17,1	9 12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13		
	20,21,23,2	6 9,10,11,12,15	5,10,11,12,14	14,18,23.2,26.2,28 .2	7,9.3	8,11,14		
	15,16,17	10,11	17,20	16,17,21				
Z8LUQ4	- 5902	GlobalFiler™- TrueAllele	® VUler Release 202	22b				
	11,16,17.3	3 17,18,23,24,25	10,14	14,15,16	11,12,13			
4	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14		
	13,15,17,1	9 12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13		
	20,21,23,2	6		14,18,23.2,26.2,28 .2	7,9.3	8,11,14		
	15,16,17	10,11			2			

			TABLE	3		
WebCod	e - Test	Amplification Kits -	Probablistic Gen	otyping Software		
	D1S1656	D2S1338	D2S441	D3S1358	D5S818	D6S1043
Item	D7S820	D8S1179	D10S1248	D12S391	D13\$317	D16S539
	D18S51	D195433	D21511	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	vWA	DYS391	DY\$570	DYS576	Y Indel	
			ltem 4e - STR	R Results		
8DDAHX	- 5901	GlobalFiler™- STRMix™ 2	2.8			
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4e	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26			14,18,23.2,26.2,28 .2	7,9.3	8,11,14
	15,16,17	10,11			2	
PD88ZF -	5902	GlobalFiler™- STRMix™ v	2.8			
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4e	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26			14,18,23.2,26.2,28 .2	7,9.3	8,11,14
	15,16,17	10,11			2	
WU8M28	- 5901	GlobalFiler™- STRMix™ 2	2.8			
	11,16,17.3	17,18,23,24,25	10,14	14,15,16	11,12,13	
4e	8,9,11,12	10,13,14,16	13,14,15	18,19,21	8,10,12,13	9,11,12,13,14
	13,15,17,19	12,13,13.2,14,15.2	28,29,30,33,33.2	11,14,15,16	X,Y	10,11,12,13
	20,21,23,26			14,18,23.2,26.2,28 .2	7,9.3	8,11,14
	15,16,17	10,11			2	

			TABLE	3		
WebCoc	le - Test	Amplification Kits - F	Probablistic Gen	otyping Software		
	D1S1656	D2S1338	D2S441	D3S1358	D5S818	D6S1043
ltem	D7\$820	D8S1179	D10S1248	D12S391	D13S317	D16\$539
	D18S51	D195433	D21S11	D22S1045	Amelogenin	CSF1PO
	FGA	Penta D	Penta E	SE33	TH01	ΤΡΟΧ
	vWA	DY\$391	DYS570	DY\$576	Y Indel	
			Item 4sp - ST	R Results		
8DDAHX	- 5901 0	GlobalFiler™- STRMix™ 2	.8			
4sp						
PD88ZF -	5902 (GlobalFiler™- STRMix™ v2	2.8			
	11,16,17.3	17,18,19,23,24,25	10,14	15,16	11,12,13	
4sp	8,9,11	10,13,14,16	13,14,15	18,19,21	10,12,13	9,11,14
	13,15,17,19	12,13.2,15.2	28,29,33,33.2	11,14,15,16	X,Y	10,11
	20,23,26			14,18,23.2	7,9.3	8,11,14
	15,16,17	10			2	
WU8M28	3 - 5901 (GlobalFiler™- STRMix™ 2	.8			
	11,16,17.3	17,18,24,25	10,14	14,15,16	11,13	
4sp	8,9,11,12	10,13,14	13,15	18,19,21	10,12,13	9,11,14
	17,19	12,13,13.2,15.2	30,33,33.2	11,14,15,16	X,Y	10,11,12
	20,23,26			14,18,26.2,28.2	7,9.3	11
	15,16,17	10			2	

Test 25-5901/2

YSTR Amplification Kit(s) & Results

					TABLE 4				
WebCo	ode - Test	Amplifice	ation Kit						
ltem	DYF38751	DYS19	DYS385	DYS389-I	DYS389-II	DYS390	DYS391	DYS392	DYS393
	DYS437	DYS438	DYS439	DYS448	DYS449	DYS456	DYS458	DYS460	DYS481
	D12218	D12233	D15549	D15570	D15576	D15027	D12032	D12043	IGAIAH4
				Item 2	2 - YSTR Re	esults			
3MQZ	UY - 5902	PowerPlex	® Y 23						
2		13	16,17	13	30	24	10	11	13
	14	10	11	19		17	15	10	22
		12	13	20	17		21	12	11
786NL	JZ - 5901	Yfiler®							
2		13	16,17	13	30	24	10	11	13
	14	10	11	19		17	15		
							21		11
8DDAF	IX - 5901	Yfiler® Plu	S						
2	35,37	13	16,17	13	30	24	10	11	13
	14	10	11	19	32	17	15	9	22
	40	12		20	17	22	21		11
8YUL92	Z - 5901	Yfiler® Plu	S						
2	35,37	13	16,17	13	30	24	10	11	13
	14	10	11	19	32	17	15	9	22
	40	12		20	17	22	21		11
9WLTT	R - 5902	Yfiler® Plu	S						
2	35,37	13	16,17	13	30	24	10	11	13
	14	10	11	19	32	17	15	9	22
	40	12		20	17	22	21		11
EHXZJK	(- 5902	Yfiler® Plu	S						
2	35,37	13	16,17	13	30	24	10	11	13
	14	10	11	19	32	17	15	9	22
	40	12		20	17	22	21		11
FPXE6L	- 5902	PowerPlex	® Y 23						
2		13	16,17	13	30	24	10	11	13
	14	10	11	19		17	15		22
		12	13	20	17		21	12	11
HJX7C	N - 5901	Yfiler® plu	IS						
2	35,37	13	16,17	13	30	24	10	11	13
	14	10	11	19	32	17	15	9	22
	40	12		20	17	22	21		11
J772JC	9 - 5902	Yfiler® Plu	S						
2	35,37	13	16,17	13	30	24	10	11	13
	14	10	11	19	32	17	15	9	22
	40	12		20	17	22	21		11
JHLQH	M - 5902	Yfiler® plu	IS						
2	35,37	13	16,17	13	30	24	10	11	13
	14	10	11	19	32	17	15	9	22
	40	12		20	17	22	21		11
К9ЕСС	E - 5902	Ytiler® Plu	S						
2	35,37	13	16,17	13	30	24	10	11	13
	14	10	11	19	32	17	15	9	22
	40	12		20	17	22	21		11

WebCo	ode - Test	Amplific	ation Kit						
ltem	DYF38751	DYS19	DYS385	DYS389-I	DYS389-11	DYS390	DYS391	DYS392	DYS393
	DYS437	DYS438	DYS439	DYS448	DYS449	DYS456	DYS458	DYS460	DYS481
	DYS518	DYS533	DYS549	DYS570	DYS576	DYS627	DYS635	DYS643	YGATAH4
				Item 2	2 - YSTR R	esults			
LBAKVF	- 5902	PowerPlex	® Y 23						
2		13	16,17	13	30	24	10	11	13
	14	10	11	19		17	15		22
		12	13	20	17		21	12	11
IG3PN	l - 5901	PowerPlex	R Y 23						
2	,	13	16 17	13	30	24	10	11	13
2	14	10	10,17	19	00	17	15		22
	14	12	13	20	17	17	21	12	11
	/E 5001	Vfilor®	10	20			21	12	
201.010	- 5701	12	14 17	10	20	0.4	10	11	10
Z	1.4	10	10,17	10	30	17	10	11	13
	14	10	11	14		17	15		11
		V(·I @					21		11
NWNR	KD - 5901	Yfiler®							
2		13	16,17	13	30	24	10	11	13
	14	10	11	19		17	15		
							21		11
PD88Z	F - 5902	Yfiler® Plu	JS						
2	35,37	13	16,17	13	30	24	10	11	13
	14	10	11	19	32	17	15	9	22
	40	12		20	17	22	21		11
Q99BC	QE - 5901	Yfiler® Plu	JS						
2	35,37	13	16,17	13	30	24	10	11	13
	14	10	11	19	32	17	15	9	22
	40	12		20	17	22	21		11
R3MFP	B - 5901	Yfiler®							
2		13	16,17	13	30	24	10	11	13
	14	10	11	19		17	15		
							21		11
) - 5901	Yfiler® Plu	IS						
2	35.37	13	16 17	13	30	24	10	11	13
2	14	10	11	10	32	17	15	0	22
	40	12		20	17	22	21	/	11
T787\//	<u> </u>	Vfilor®		20		22	21		
120207	- 5701	12	14 17	12	20	24	10	11	12
Z	1.4	10	10,17	10	30	17	10	11	13
	14	10	11	14		/	15		11
<u> </u>							21		11
VK9834	4 - 5902	Yfiler® Plu	JS						
2	35,37	13	16,17	13	30	24	10	11	13
	14	10	11	19	32	17	15	9	22
	40	12		20	17	22	21		11
WU8M	28 - 5901	Yfiler® Plu	JS						
2	35,37	13	16,17	13	30	24	10	11	13
	14	10	11	19	32	17	15	9	22
	40	12		20	17	22	21		11
XCU6U	19 - 5902	Yfiler® Plu	JS						
2	35,37	13	16,17	13	30	24	10	11	13
	14	10	11	19	32	17	15	9	22
	40	12		20	17	22	21		11

					TABLE 4				
WebC	ode - Test	Amplific	ation Kit						
ltem	DYF38751 DYS437 DYS518	DYS19 DYS438 DYS533	DYS385 DYS439 DYS549	DYS389-I DYS448 DYS570	DYS389-II DYS449 DYS576	DYS390 DYS456 DYS627	DYS391 DYS458 DYS635	DYS392 DYS460 DYS643	DYS393 DYS481 YGATAH4
				Item 3	3 - YSTR Re	esults			
8YUL9	Z - 5901	Yfiler® Plu	JS						
3	35,36,37	13,14	11,13,16, 17	13,14	30	24	10	11,13	13
	14,15	10,12	11,12	19	29,32	14,17	15,17	9,11	22
	38,40	12		18,20	17,19	20,22	21,24		11,12
LG3PN	JJ - 5901	PowerPlex	® Y 23						
3		13,14	11,13,16, 17	13,14	30	24	10	11,13	13
	14,15	10,12	11,12	19		14,17	15,17		22
		12	13	18,20	17,19		21,24	10,12	11,12

					TABLE 4				
WebCo	de - Test	Amplific	ation Kit						
ltom	DVE29761		DVC295	DVC280 1		DV6300	DV\$201	DV5202	DV\$202
Item	DYS437	DYS438	DYS439	DYS448	DYS449	DYS456	DYS458	DYS460	DYS481
	DYS518	DY\$533	DYS549	DYS570	DYS576	DYS627	DYS635	DYS643	YGATAH4
				Item 3	e - YSTR R	esults			
3MQZI	JY - 5902	PowerPlex	® Y 23						
Зe		13,14	11,13,16,	13,14	30	24	10	11,13	13
			17						
	14,15	10,12	11,12	19		14,17	15,17		22
		12	13	18,20	17,19		21,24	10,12	11,12
786NU	Z - 5901	Ytiler®							
3e		13,14	11,13,16, 17	13,14	30	24	10	11,13	13
	14,15	10,12	11,12	19		14,17	15,17		
							21,24		11,12
8DDAH	IX - 5901	Yfiler® Plu	JS						
3e	35,36,37	13,14	11,13,16, 17	13,14	30	24	10	11,13	13
	14,15	10,12	11,12	19	29,32	14,17	15,17	9,11	22
	38,40	12		18,20	17,19	20,22	21,24		11,12
9WLTTF	R - 5902	Yfiler® Plu	JS						
3e	35,36,37	13,14	11,13,16, 17	13,14	30,30	24	10	11,12,13	13
	14,15	10,12	11,12	19	29,32	14,17	15,17	9,11	22
	38,40	12		18,20	17,19	20,22	21,24		11,12
EHXZJK	- 5902	Yfiler® Plu	JS						
3e	35,37	13,14	11,13,16, 17	13,14	30	24	10	11,13	13
	14,15	10	11,12	19	29,32	14,17	15,17	9	22
	40	12		18,20	17,19	20,22	21,24		11
FPXE6L	- 5902	PowerPlex	® Y 23						
Зе		13,14	11,13,16, 17	13,14	30	24	10	11,13	13
	14,15	10,12	11,12	19		14,17	15,17		22
		12	13	18,20	17,19		21,24	10,12	11,12
HJX7CI	V - 5901	Yfiler® plเ	JS						
3e	35,36,37	13,14	11,13,16, 17	13,14	30	24,24	10	11,13	13
	14,15	10,12	11,12	19	29,32	14,17	15,17	9,11	22
	38,40	12		18,20	17,19	20,22	21,24		11,12
J772JG	5 - 5902	Yfiler® Plu	JS						
Зe	35	13,14	17	13,14		24	10		13
	14	10			29,32	14	15	9	22
		12		18,20	17	22	21,24		11
JHLQH	M - 5902	Ytiler® plu	JS						
3e	35,36,37	13,14	11,13,16, 17	13,14	30	24	10	11,13	13
	14,15	10,12	11,12	19	29,32	14,17	15,17	9,11	22
	38,40	12		18,20	17,19	20,22	21,24		11,12
К9ЕСС	E - 5902	Yfiler® Plu	JS						
3e	35,36,37	13,14	11,13,16, 17	13,14	30	24	10	11,13	13
	14,15	10,12	11,12	19	29,32	14,17	15,17	9,11	22
	38,40	12		18,20	17,19	20,22	21,24		11,12

WebCo	ode - Test	Amplific	ation Kit						
ltem	DYF387S1	DYS19	DYS385	DYS389-I	DYS389-II	DYS390	DYS391	DYS392	DYS393
	DYS437	DYS438	DYS439	DYS448	DYS449	DYS456	DYS458	DYS460	DYS481
	DY\$518	DYS533	DYS549	DYS570	DYS576	DYS627	DYS635	DYS643	YGATAH4
				Item 3	e - YSTR R	esults			
LBAKVF	- 5902	PowerPlex	® Y 23						
3e		13,14	11,13,16, 17	13,14	30	24	10	11,13	13
	14,15	10,12	11,12	19		14,17	15,17		22
		12	13	18,20	17,19		21,24	10,12	11,12
PD88ZI	- 5902	Yfiler® Plu	JS						
Зе	35,36,37	13,14	11,13,16, 17	13,14	30	24	10	11,13	13
	14,15	10,12	11,12	19	29,32	14,17	15,17	9,11	22
	38,40	12		18,20	17,19	20,22	21,24		11,12
Q99BC	QE - 5901	Yfiler® Plu	JS						
3e	35,36,37	13,14	11,13,16, 17	13,14	30	24	10	11,13	13
	14,15	10,12	11,12	19	29,32	14,17	15,17	9,11	22
	38,40	12		18,20	17,19	20,22	21,24		11,12
VK9834	4 - 5902	Yfiler® Plu	JS						
3e	35,36,37	13,14	11,13,16, 17	13,14	30	24	10	11,13	13
	14,15	10,12	11,12	19	29,32	14,17	15,17	9,11	22
	38,40	12		18,20	17,19	20,22	21,24		11,12
WU8M	28 - 5901	Yfiler® Plu	JS						
Зе	35,36,37	13,14	11,13,16, 17	13,14	30	24	10	11,13	13
	14,15	10,12	11,12	19	29,32	14,17	15,17	9,11	22
	38,40	12		18,20	17,19	20,22	21,24		11,12
XCU6U	9 - 5902	Yfiler® Plu	JS						
Зе	35,36,37	13,14	11,13,16, 17	13,14	30	24	10	11,13	13
	14,15	10,12	11,12	19	29,32	14,17	15,17	9,11	22
	38,40	12		18,20	17,19	20,22	21,24		11,12

WebCo	de - Test	Amplifica	ation Kit						
Item	DYF387S1	DYS19	DYS385	DYS389-I	DYS389-II	DYS390	DYS391	DYS392	DYS393
	DYS437	DYS438	DYS439	DYS448	DYS449	DYS456	DY\$458	DYS460	DYS481
	DYS518	DYS533	DYS549	DYS570	DYS576	DYS627	DYS635	DYS643	YGATAH4
				Item 3s	sp - YSTR F	Results			
3MQZL	JY - 5902	PowerPlex®	€ Y 23		•				
Зsр		14	11,13	14	30	24	10	13	13
	15	12	12	19		14	17		22
		12	13	18	19		24	10	12
786NU	Z - 5901	Yfiler®							
Зsр		13,14	11,13,16, 17	13,14	30	24	10	11,13	13
	14,15	10,12	11,12	19		14,17	15,17		
							21,24		11,12
8DDAHX - 5901		Yfiler® Plus	S						
Зsр	35,36	14	11,13	14	30	24	10	13	13
	15	12	12	19	29	14	17	11	22
	38	12		18	19	20	24		12
9WLTTR	2 - 5902	Yfiler® Plus	S						
Зsр	35,36	13,14	11,13	13,14	30	24	10	13	13
	15	12	12	19	29	14	17	11	22
	38	12		18	19	20	24		11,12
EHXZJK	- 5902	Yfiler® Plu	S						
Зsр	35,36	14	11,13	14	30	24	10	13	13
	15	12	12	19	29	14	17	11	22
	38	12		18	19	20	24		12
FPXE6L	- 5902	PowerPlex®	€ Y 23						
Зsр		14	11,13	14	30	24	10	13	13
	15	12	12	19		14	17		22
		12	13	18	19		24	10	12
HJX7CN	V - 5901	Yfiler® plu:	S						
Зsp	35,36	14	11,13	14	30	24	10	13	13
	15	12	12	19	29	14	17	11	22
	38	12		18	19	20	24		12
J772JG	- 5902	Yfiler® Plus	s						
Зsр	35,36	14	11,13	14	30	24	10	13	13
	15	12	12	19	29	14	17	11	22
	38	12		18	19	20	24		12
JHLQHI	M - 5902	Yfiler® plu	s						
Зsp	35,36	14	11,13	14	30	24	10	13	13
	15	12	12	19	29	14	17	11	22
	38	12		18	19	20	24		12
K9ECCI	E - 5902	Yfiler® Plus	S						
3sp	35,36	14	11,13	14	30	24	10	13	13
	15	12	12	19	29	14	17	11	22
	38	12		18	19	20	24		12
LBAKVF	- 5902	PowerPlex®	9 Y 23						
3sp		14	11,13	14	30	24	10	13	13
,	15	12	12	19		14	17		22
		12	13	18	19		24	10	12
PD88ZF	- 5902	Yfiler® Plus	S						
3sp	35,36	14	11,13	14	30	24	10	13	13
,	15	12	12	19	29	14	15,17	11	22
	38	12		18	19	20,22	24		12

WebC	ode - Test	Amplific	ation Kit							
ltem	DYF387S1	DYS19	DYS385	DYS389-I	DYS389-II	DYS390	DYS391	DYS392	DYS393	
	DYS437	DYS438	DYS439	DYS448	DYS449	DYS456	DYS458	DYS460	DYS481	
	DYS518	DYS533	DYS549	DYS570	DYS576	DYS627	DYS635	DYS643	YGATAH4	
				Item 3	sp - YSTR F	Results				
Q99B0	QE - 5901	Yfiler® Plu	US							
Зsp	35,36	14	11,13	14	30	24	10	13	13	
	15	12	12	19	29	14	17	11	22	
	38	12		18	19	20	24		12	
VK983	4 - 5902	Yfiler® Plu	US							
Зsp	35,36	14	11,13	14	30	24	10	13	13	
	15	12	12	19	29	14	17	11	22	
	38	12		18	19	20	24		12	
WU8M	28 - 5901	Yfiler® Plu	US							
Зsp	35,36	14	11,13	14	30	24	10	13	13	
	15	12	12	19	29	14	17	11	22	
	38	12		18	19	20	24		12	
XCU6L	J9 - 5902	Yfiler® Plu	US							
Зsр	35,36,37	13,14	11,13,16, 17	13,14	30	24	10	11,13	13	
	14,15	10,12	11,12	19	29,32	14,17	15,17	9,11	22	
	38,40	12		18,20	17,19	20,22	21,24		11,12	

WebCode - Test Amplification Kit Item DYF38751 DYS19 DYS385 DYS389-I DYS389-II DYS390 DYS391 DYS392 DYS3	393 181
Item DYF38751 DYS19 DYS385 DYS389-I DYS389-II DYS390 DYS391 DYS392 DYS3	393 181
ITEM DTF36/51 DT519 DT5365 DT5369-1 DT5369-11 DT5390 DT5391 DT5392 DT53	181
015457 015458 015459 015448 015449 015456 015458 015460 0154	
DYS518 DYS533 DYS549 DYS570 DYS576 DYS627 DYS635 DYS643 YGAT	AH4
Itom A VSTP Pocults	
3MQ7UY 5902 PowerPlex® Y 23	
A 13 14 15 11 15 16 12 13 28 20 30 22 24 10 11 10 11 13 1	14
17	14
14,15,16 10,12 11,12 19,21 14,16,17 15,18,19 20,22	2,23
10,12 12,13 17,20 16,17,21 21,23 10,11,12 11,7	13
786NUZ - 5901 Yfiler®	
4 13,14,15 11,15,16, 12,13 28,29,30 22,24 10,11 10,11,14 13,1	4
17	
14,15,16 10,12 11,12 19,21 14,16,17 15,18,19	
21,23 11,	13
8YUL9Z - 59UT Yfiler® Plus	
4 35,36,37,3 13,14,15 11,15,16, 12,13 28,29,30 22,24 10,11 10,11,14 13,1 17	14
14.15.16 10.12 11.12 19.21 28.30.32 14.16.17 15.18.19 9.10.11 20.22	2.23
37,38,40 10,12 17,20 16,17,21 20,22 21,23 11,1	, 13
9WLTTR - 5902 Yfiler® Plus	
4 35,36,37,3 13,14,15 11,15,16, 12,13 28,29,30 22,24 10,11 10,11,14 13,1	4
17	
14,15,16 10,12 11,12 19,21 28,30,32 14,16,17 15,18,19 9,10,11 20,22	2,23
37,38,40 10,12 17,20 16,17,21 20,22 21,23 11,1	13
EHXZJK - 5902 Yfiler® Plus	
4 35,36,37,3 13,14,15 11,15,16, 12,13 28,29,30 22,24 10 10,11,14 13,1	4
	0.03
37 38 40 10 12 17 20 16 17 21 20 22 21 23 11	13
FPXF61 - 5902 PowerPlex® Y 23	
4 13 14 15 11 15 16 12 13 28 29 30 22 24 10 11 10 11 14 13	4
17	1 7
14,15,16 10,12 11,12 19,21 14,16,17 15,18,19 20,22	2,23
10,12 12,13 17,20 16,17,21 21,23 10,11,12 11,1	13
HJX7CN - 5901 Yfiler® plus	
4 35,36,37,3 13,14,15 11,15,16, 12,13 28,29,30 22,24 10,11 10,11,14 13,1	4
14,15,16 10,12 11,12 19,21 28,30,32 14,16,17 15,18,19 9,10,11 20,22 37,38,40 10,12 17,20 16,17,21 20,22 21,23 11	2,23
1772 IC 5002 Villar® Plue	15
J/7ZJG - 590Z THIER® Flos)
4 55,56,57,5 15,14,15 11,15,16, 12,15 26,29,50 22,24 10 10,11,14 13 17)
14,15,16 10,12 11,12 19,21 28,30,32 14,16,17 15,18,19 9,10,11 20,22	2,23
38,40 10,12 17,20 16,17,21 20,22 21,23 11,1	13
JHLQHM - 5902 Yfiler® plus	
4 35,36,37,3 13,14,15 11,15,16, 12,13 28,29,30 22,24 10,11 10,11,14 13,1	4
17	
14,15,16 10,12 11,12 19,21 28,30,32 14,16,17 15,18,19 9,10,11 20,22	2,23
37,38,40 10,12 17,20 16,17,21 20,22 21,23 11,1	13
K9ECCE - 5902 Yfiler® Plus	
4 35,36,37,3 13,14,15 11,15,16, 12,13 28,29,30 22,24 10,11 10,11,14 13,1	4
	23
37,38,40 10,12 17,20 16,17,21 20,22 21,23 11,1	13

					TABLE 4				
WebC	ode - Test	Amplific	ation Kit						
ltem	DYF387S1 DYS437 DYS518	DYS19 DYS438 DYS533	DYS385 DYS439 DYS549	DYS389-I DYS448 DYS570	DYS389-II DYS449 DYS576	DYS390 DYS456 DYS627	DYS391 DYS458 DYS635	DYS392 DYS460 DYS643	DYS393 DYS481 YGATAH4
				Item 4	4 - YSTR Re	esults			
LBAKV	F - 5902	PowerPlex	® Y 23						
4		13,14,15	11,15,16, 17	12,13	28,29,30	22,24	10,11	10,11,14	13,14
	14,15,16	10,12	11,12	19,21		14,16,17	15,18,19		20,22,23
		10,12	12,13	17,20	16,17,21		21,23	10,11,12	11,13
LG3PN	JJ - 5901	PowerPlex	® Y 23						
4		13,14,15	11,15,16, 17	12,13	28,29,30	22,24	10,11	10,11,14	13,14
	14,15,16	10,12	11,12	19,21		14,16,17	15,18,19		20,22,23
		10,12	12,13	17,20	16,17,21		21,23	10,11,12	11,13
Q99B	QE - 5901	Yfiler® Plu	JS						
4	35,36,37,3	13,14,15	11,15,16, 17	12,13	28,29,30	22,24	10,11	10,11,14	13,14
	14,15,16	10,12	11,12	19,21	28,30,32	14,16,17	15,18,19	9,10,11	20,22,23
	37,38,40	10,12		17,20	16,17,21	-,-	21,23		11,13
VK983	4 - 5902	Yfiler® Plu	JS						
4	35,37,38	13,14,15	11,15,16, 17	12,13	28,29,30	22,24	10,11	10,11,14	13,14
	14,15,16	10,12	11,12	19,21	28,30,32	14,16,17	15,18,19	9,10,11	20,22,23
	37,38,40	10,12		17,20	16,17,21	20,22	21,23		11,13
XCU6l	J9 - 5902	Yfiler® Plu	JS						
4	35,36,37,3	13,14,15	11,15,16, 17	12,13	28,29,30	22,24	10,11	10,11,14	13,14
	14,15,16	10,12	11,12	19,21	28,29,30,3	14,16,17	15,18,19	9,10,11	20,22,23
	37,38,40	10,12		17,20	16,17,21	20,22	21,23		11,13

					TABLE 4				
WebCo	ode - Test	Amplific	ation Kit						
ltem	DYF38751 DYS437 DYS518	DYS19 DYS438 DYS533	DYS385 DYS439 DYS549	DYS389-I DYS448 DYS570	DYS389-II DYS449 DYS576	DYS390 DYS456 DYS627	DYS391 DYS458 DYS635	DYS392 DYS460 DYS643	DYS393 DYS481 YGATAH4
				Item 4	e - YSTR R	esults			
8DDAH	HX - 5901	Yfiler® Plu	JS						
4e	35,36,37,3	13,14,15	11,15,16, 17	12,13	28,29,30	22,24	10,11	10,11,14	13,14
	14,15,16	10,12	11,12	19,21	28,30,32	14,16,17	15,18,19	9,10,11	20,22,23
	37,38,40	10,12		17,20	16,17,21	20,22	21,23		11,13
PD88Z	F - 5902	Yfiler® Plu	JS						
4e	35,36,37,3	13,14,15	11,15,16, 17	12,13	28,29,30	22,24	10,11	10,11,14	13,14
	14,15,16	10,12	11,12	19,21	28,30,32	14,16,17	15,18,19	9,10,11	20,22,23
	37,38,40	10,12		17,20	16,17,21	20,22	21,23		11,13
WU8M	28 - 5901	Yfiler® Plu	JS						
4e	35,36,37,3	13,14,15	11,15,16, 17	12,13	28,29,30	22,24	10,11	10,11,14	13,14
	14,15,16	10,12	11,12	19,21	28,30,32	14,16,17	15,18,19	9,10,11	20,22,23
	37,38,40	10,12		17,20	16,17,21	20,22	21,23		11,13

WebCo	ode - Test	Amplifica	tion Kit							
ltem	DYF38751	DYS19	DYS385	DYS389-I	DYS389-II	DYS390	DYS391	DYS392	DYS393	
	DYS437	DYS438	DYS439	DYS448	DYS449	DYS456	DYS458	DYS460	DYS481	
	DYS518	DYS533	DYS549	DYS570	DYS576	DYS627	DYS635	DYS643	YGATAH4	
				Item 4	sp - YSTR F	Results				
8DDAH	HX - 5901	Yfiler® Plus								
4sp										
						22				
PD88Z	F - 5902	Yfiler® Plus								
4sp	35,37	13				24		10,11	13	
				19			19	9,11		
		10		20		22	21,23		11	
WU8M	28 - 5901	Yfiler® Plus								
4sp	35,36,37,3	13	15,16,17	13			10		13	
	14	10	11,12	19	28,32	17	15,18,19	9	23	
	40	12		17.20	16.17	20.22	21		11	

Additional DNA Results

TABLE 5

Additional DNA results found to be concordant at a pre-existing locus are retained solely within the applicable tables. Non-concordant results and results for loci not found elsewhere will remain in this table.

Locus	WebCode-	ltem 1	Item 2	Item 3	Item 3e	ltem 3sp	Item 4	ltem 4e	ltem 4sp
	Test								

No additional DNA results were reported.

DNA Mixture Concentrations and Proportions

		TABLE 6								
	Item 3 Results									
WebCode-Test	Number of Contributors	Contributor(s)	DNA Concentration (ng/uL)	DNA Proportion (%)						
LG3PNJ - 5901	3	Suspect Unknown Victim	208 235 91.0	39.00 44.00 17.00						
		Response Summary	y: Item 3							
	Estimated Numbe	<u>er of Contributors</u>	Percent Reported							
		3	1 (100%)							

Item 3e Results							
WebCode-Test	Number of Contributors	Contributor(s)	DNA Concentration (ng/uL)	DNA Proportion (%)			
2QGMC2 - 5902	3	Suspect Unknown Victim					
34G3B3 - 5902	3	Suspect Unknown Victim		34.73 46.61 18.67			
37J2CU - 5901	3	Suspect Unknown Victim	0.3410 0.1840 0.1350	51.66 27.89 20.46			
3WHFJ3 - 5902	3	Suspect Unknown Victim		35.93 40.95 23.12			
44PDRD - 5901	3	Suspect Unknown Victim		33.00 50.00 17.00			
4BEUJW - 5902	3	Suspect Unknown Victim	0.00210 0.00130 0.00160	42.00 26.00 32.00			
4FBJY4 - 5902	3	Suspect Unknown Victim	46.0 27.0 25.0	471.00 282.00 258.00			
4UQG64 - 5901	3	Suspect Unknown Victim		41.00 36.00 23.00			
4XQRZ4 - 5901	3	Suspect Unknown Victim		39.00 45.00 16.00			
62K9Q4 - 5901	3	Suspect Unknown Victim		36.00 42.00 22.00			
63VBC2 - 5901	3	Suspect Unknown Victim		35.00 43.00 21.00			
6PTH6Y - 5902	3	Suspect Unknown Victim					
77A6RY - 5901	3						
786NUZ - 5901	3	Suspect Unknown Victim	0.09490 0.04450 0.03440	54.56 25.60 19.80			

Item 3e Results						
WebCode-Test	Number of Contributors	Contributor(s)	DNA Concentration (ng/uL)	DNA Proportion (%)		
7H4JRV - 5902	3	Suspect Unknown Victim				
8Y8DYV - 5902	3	Suspect Unknown Victim				
9CLYK9 - 5901	3	Suspect Unknown Victim		73.00 18.00 9.00		
9MBR8Q - 5901	3	Suspect Unknown Victim	0.07000 0.09000 0.04000	37.38 43.24 19.38		
9WLTTR - 5902	3	Suspect Unknown Victim	4.70 3.59 2.27	44.54 33.97 21.49		
B7W38V - 5901	3					
BDR7Z7 - 5901	3	Suspect Unknown Victim		34.00 47.00 19.00		
BKHHBU - 5901	3					
BNGT8U - 5902	3	Suspect Unknown Victim		37.00 42.00 21.00		
CDTG9M - 5901	3	Suspect Unknown Victim	0.2983 0.1120 0.1397	54.23 20.36 25.40		
CKECAQ - 5902	3	Suspect Unknown Victim				
CRBAWU - 5901	3	Suspect Unknown Victim		43.30 33.30 23.30		
CRVQCW - 5901	3	Suspect Unknown Victim		45.00 36.00 19.00		
CXYRCQ - 5902	3	Suspect Unknown Victim				

Item 3e Results							
WebCode-Test	Number of Contributors	Contributor(s)	DNA Concentration (ng/uL)	DNA Proportion (%)			
CY6DP6 - 5901	3	Suspect Unknown Victim		38.00 43.00 19.00			
CZ6EPV - 5901	3	Suspect Unknown Victim		33.00 47.00 20.00			
E34ANN - 5902	3	Suspect Unknown Victim					
EHXZJK - 5902	3	Suspect Unknown Victim	0.2260 0.01290 0.08400	70.00 4.00 26.00			
ELJJPP - 5902	3	Suspect Unknown Victim					
EY6YTN - 5902	3	Suspect Unknown Victim					
F7H8UP - 5902	3	Suspect Unknown Victim		49.16 33.13 17.71			
F8XQDM - 5902	3	Suspect Unknown Victim					
GCH7HK - 5902	3	Suspect Unknown Victim					
H484WJ - 5902	3	Suspect Unknown Victim					
HEQHYL - 5902	3	Suspect Unknown Unknown					
HJX7CN - 5901	3	Suspect Unknown Victim	0.3600 0.3600 0.3600	39.00 45.00 16.00			
J37QNH - 5902	3	Suspect Unknown Victim	0.2210 0.08300 0.05800	61.00 23.00 16.00			
J772JG - 5902	3	Suspect Unknown Victim	0.01504 0.00896 0.00800	47.00 28.00 25.00			

Item 3e Results								
WebCode-Test	Number of Contributors	Contributor(s)	DNA Concentration (ng/uL)	DNA Proportion (%)				
JHLQHM - 5902	3	Suspect Unknown Victim	236 4.00 122	31.00 53.00 16.00				
JPGUBX - 5901	3	Suspect Unknown Victim		93.00 4.00 3.00				
K9ECCE - 5902	3	Suspect Unknown Victim	6.65 5.14 3.37	43.88 33.90 22.22				
KPDQAK - 5902	3	Suspect Unknown Victim		41.54 35.82 22.64				
L67JVL - 5901	3	Suspect Unknown Victim		37.00 43.00 20.00				
L8XEJE - 5901	3	Unknown Unknown Victim		32.78 48.58 18.64				
LHUN3H - 5902	3	Suspect Unknown Victim						
LYKQJL - 5901	3	Suspect Unknown Victim		33.00 40.00 26.00				
MW99ZF - 5902	3	Suspect Unknown Victim						
P3QH3A - 5901	3	Suspect Unknown Victim		41.09 36.09 22.82				
P8KEPF - 5902	3	Suspect Unknown Victim		42.11 36.32 21.57				
P8TP7Q - 5901	3	Suspect Unknown Victim		33.00 43.00 24.00				
Q99BQE - 5901	3	Suspect Unknown Victim	0.1400 0.1140 0.04800	46.45 37.76 15.79				
R9RDYE - 5901	3	Suspect Unknown Victim		34.00 40.00 26.00				

		Item 3e Resul	ts	
WebCode-Test	Number of Contributors	Contributor(s)	DNA Concentration (ng/uL)	DNA Proportion (%)
RCPXZD - 5901	3	Suspect Unknown Victim	0.1600 0.1640 0.07600	40.00 41.00 19.00
TMDEV4 - 5902	3	Suspect Unknown Victim	1.28 1.12 0.6030	42.64 37.26 20.10
UNVR99 - 5902	3	Suspect Unknown Victim		
VFHZHA - 5902	3	Suspect Unknown Victim		35.98 42.13 21.89
VK9834 - 5902	3	Suspect Unknown Victim	1.89 1.43 1.07	43.08 32.57 24.35
W2RLKZ - 5901	3	Suspect Unknown Victim		34.00 48.00 18.00
WRGKYZ - 5901	3	Suspect Unknown Victim		37.00 44.00 18.00
X28843 - 5901	3	Suspect Unknown Victim	0.4683 0.5301 0.2486	37.54 42.53 19.93
YGJ6L9 - 5901	3	Suspect Unknown Victim		37.00 45.00 18.00
Z8LUQ4 - 5902	3	Suspect Unknown Victim		
	R	Response Summary:	Item 3e	
	Estimated Numbe	er of Contributors	Percent Reported	
		3	66 (100%)	

Item 3sp Results							
WebCode-Test	Number of Contributors	Contributor(s)	DNA Concentration (ng/uL)	DNA Proportion (%)			
2QGMC2 - 5902	2	Unknown Unknown					
34G3B3 - 5902	1	Unknown		100.00			
37J2CU - 5901	1	Unknown	0.4412	100.00			
3WHFJ3 - 5902	1	Unknown		100.00			
44PDRD - 5901	1	Unknown		100.00			
4BEUJW - 5902	1	Unknown	0.02800	100.00			
4FBJY4 - 5902	1	Unknown	257	100.00			
4UQG64 - 5901	1	Unknown		100.00			
4XQRZ4 - 5901	1	Unknown		100.00			
62K9Q4 - 5901	1	Unknown		100.00			
63VBC2 - 5901	1	Unknown		100.00			
6PTH6Y - 5902	1	Unknown					
77A6RY - 5901	1						
786NUZ - 5901	3	Suspect Unknown Victim	0.9440 1.24 0.4210	36.20 47.64 16.14			
7H4JRV - 5902	2	Unknown Unknown					
9CLYK9 - 5901	1	Unknown		100.00			
9MBR8Q - 5901	3	Suspect Unknown Victim	0.03000 0.3400 0.02000	8.38 85.69 5.93			
9WLTTR - 5902	3	Suspect Unknown Victim	0.5800 7.08 0.3500	7.28 88.34 4.37			
B7W38V - 5901	1						
BDR7Z7 - 5901	2	Unknown Unknown		1.00 99.00			
BKHHBU - 5901	1						

Item 3sp Results								
WebCode-Test	Number of Contributors	Contributor(s)	DNA Concentration (ng/uL)	DNA Proportion (%)				
BNGT8U - 5902	1	Unknown		100.00				
CDTG9M - 5901	1	Unknown	0.09200	100.00				
CKECAQ - 5902	1	Unknown						
CRBAWU - 5901	1	Unknown		100.00				
CRVQCW - 5901	1	Unknown		100.00				
CXYRCQ - 5902	1	Unknown						
CY6DP6 - 5901	1	Unknown		100.00				
CZ6EPV - 5901	1	Unknown		100.00				
E34ANN - 5902	2	Suspect Unknown						
EHXZJK - 5902]	Unknown	0.3180	100.00				
ELJJPP - 5902	2	Unknown Unknown						
EY6YTN - 5902	2	Unknown Unknown						
F7H8UP - 5902	1	Suspect		100.00				
F8XQDM - 5902	2	Unknown Unknown						
GCH7HK - 5902	1	Unknown						
H484WJ - 5902	2	Unknown Unknown						
HEQHYL - 5902	1	Unknown						
HJX7CN - 5901	1	Unknown	0.7100	100.00				
J37QNH - 5902	2	Unknown Victim	0.7460 0.01500	98.00 2.00				
J772JG - 5902	1	Unknown	0.06700	100.00				
JHLQHM - 5902	1	Unknown	417	100.00				
JPGUBX - 5901	2	Suspect Unknown		1.00 99.00				

Item 3sp Results							
WebCode-Test	Number of Contributors	Contributor(s)	DNA Concentration (ng/uL)	DNA Proportion (%)			
K9ECCE - 5902	2	Suspect Unknown	0.2000 9.19	2.16 97.84			
KPDQAK - 5902	1	Unknown		100.00			
L67JVL - 5901	1	Unknown		100.00			
L8XEJE - 5901	3	Unknown Unknown Victim		97.84 1.43 0.73			
LHUN3H - 5902	2	Unknown Unknown					
LYKQJL - 5901	1	Unknown		100.00			
MW99ZF - 5902	2	Unknown Unknown					
P3QH3A - 5901	3	Suspect Unknown Victim		2.64 95.24 2.12			
P8KEPF - 5902	1	Unknown		100.00			
P8TP7Q - 5901	1	Unknown		100.00			
Q99BQE - 5901	1	Unknown	0.9950	100.00			
R9RDYE - 5901	1	Unknown		100.00			
RCPXZD - 5901	1	Unknown	0.1000	100.00			
TMDEV4 - 5902	1	Unknown	2.65	100.00			
UNVR99 - 5902	3	Unknown Unknown Unknown					
VFHZHA - 5902	1	Unknown		100.00			
VK9834 - 5902	1	Unknown	0.6970	100.00			
W2RLKZ - 5901]	Unknown		100.00			
WRGKYZ - 5901]	Unknown		100.00			
X28843 - 5901	2	Unknown Unknown	0.01240 0.7702	1.59 98.41			

Item 3sp Results							
WebCode-Test	Number of Contributors	Contributor(s)	Conce	DNA entration (ng/uL)	DNA Proportion (%)		
YGJ6L9 - 5901	1	Unknown			100.00		
Z8LUQ4 - 5902	2	Unknown Unknown					
	R	esponse Summary:	: Item 3sp				
	Estimated Numbe	<u>r of Contributors</u>	<u>Percent</u>	Reported			
		1 2	44 15	(68%) (23%)			

6

(9%)

3
Item 4 Results				
WebCode-Test	Number of Contributors	Contributor(s)	DNA Concentration (ng/uL)	DNA Proportion (%)
2QGMC2 - 5902	3	Suspect Unknown Unknown		
34G3B3 - 5902	3	Suspect Unknown Unknown		51.72 31.81 16.47
37J2CU - 5901	3	Suspect Unknown Victim	2.13 1.10 0.7110	54.12 27.82 18.05
3WHFJ3 - 5902	3	Suspect Unknown Unknown		37.85 49.16 12.99
44PDRD - 5901	3	Suspect Unknown Unknown		51.00 22.00 28.00
4BEUJW - 5902	3	Suspect Unknown Unknown	1.09 0.6698 0.3349	52.00 32.00 16.00
4FBJY4 - 5902	3	Suspect Unknown Unknown	166 184 65.0	40.00 444.00 156.00
4UQG64 - 5901	3	Suspect Unknown Unknown		50.00 30.00 20.00
4XQRZ4 - 5901	3	Suspect Unknown Unknown		42.00 33.00 25.00
62K9Q4 - 5901	3	Suspect Unknown Unknown		52.00 27.00 20.00
63VBC2 - 5901	3	Suspect Unknown Unknown		40.00 46.00 14.00
6PTH6Y - 5902	3	Suspect Unknown Unknown		
77A6RY - 5901	3			
786NUZ - 5901	3	Suspect Unknown Unknown	0.1010 0.03300 0.03300	60.33 19.83 19.83

Item 4 Results				
WebCode-Test	Number of Contributors	Contributor(s)	DNA Concentration (ng/uL)	DNA Proportion (%)
7H4JRV - 5902	3	Suspect Unknown Unknown		
8Y8DYV - 5902	3	Suspect Unknown Unknown		
9CLYK9 - 5901	3	Suspect Unknown Unknown		51.00 34.00 16.00
9MBR8Q - 5901	3	Suspect Unknown Unknown	10.4 7.27 4.37	47.27 32.93 19.80
9WLTTR - 5902	3	Suspect Unknown Unknown	4.48 1.84 2.36	51.61 21.22 27.16
B7W38V - 5901	3			
BDR7Z7 - 5901	3	Suspect Unknown Unknown		80.00 8.00 11.00
BKHHBU - 5901	3			
BNGT8U - 5902	3	Suspect Unknown Unknown		55.00 17.00 28.00
CDTG9M - 5901	3	Suspect Unknown Unknown	0.8702 0.3478 0.4959	50.77 20.29 28.93
CKECAQ - 5902	3	Suspect Unknown Unknown		
CRBAWU - 5901	3	Suspect Unknown Unknown		48.30 18.30 33.30
CRVQCW - 5901	3	Suspect Unknown Unknown		46.00 30.00 24.00
CXYRCQ - 5902	3	Suspect Unknown Unknown		

Item 4 Results				
WebCode-Test	Number of Contributors	Contributor(s)	DNA Concentration (ng/uL)	DNA Proportion (%)
CY6DP6 - 5901	3	Suspect Unknown Unknown		40.00 26.00 34.00
CZ6EPV - 5901	3	Suspect Unknown Unknown		46.00 32.00 23.00
E34ANN - 5902	3	Suspect Unknown Unknown		
EHXZJK - 5902	3	Suspect Unknown Unknown	1.35 0.4060 0.9200	50.00 15.00 34.00
ELJJPP - 5902	3	Suspect Unknown Unknown		
EY6YTN - 5902	3	Suspect Unknown Unknown		
F7H8UP - 5902	3	Suspect Unknown Unknown		52.16 31.53 16.31
F8XQDM - 5902	3	Suspect Unknown Unknown		
GCH7HK - 5902	3	Suspect Unknown Unknown		
H484WJ - 5902	3	Suspect Unknown Unknown		
HEQHYL - 5902	3	Suspect Unknown Unknown		
HJX7CN - 5901	4	Suspect Unknown Unknown Unknown	0.7700 0.7700 0.7700 0.7700	
J37QNH - 5902	3	Suspect Unknown Unknown	4.31 2.73 1.76	49.00 31.00 20.00
J772JG - 5902	3	Suspect Unknown Unknown	1.37 0.4323 0.7375	54.00 17.00 29.00

Item 4 Results				
WebCode-Test	Number of Contributors	Contributor(s)	DNA Concentration (ng/uL)	DNA Proportion (%)
JHLQHM - 5902	3	Suspect Unknown Unknown	964 364 364	57.00 215.00 215.00
JPGUBX - 5901	3	Suspect Unknown Unknown		47.00 13.00 40.00
K9ECCE - 5902	3	Suspect Unknown Unknown	7.31 3.65 2.79	53.16 26.52 20.32
KPDQAK - 5902	3	Suspect Unknown Unknown		53.13 26.73 20.14
L67JVL - 5901	3	Suspect Unknown Unknown		57.00 27.00 16.00
L8XEJE - 5901	3	Unknown Unknown Unknown		15.18 25.73 59.09
LG3PNJ - 5901	3	Suspect Unknown Unknown	95.0 29.0 48.0	55.00 17.00 28.00
LHUN3H - 5902	3	Suspect Unknown Unknown		
LYKQJL - 5901	3	Suspect Unknown Victim		49.00 35.00 16.00
MW99ZF - 5902	3	Suspect Unknown Unknown		
P3QH3A - 5901	3	Suspect Unknown Unknown		53.67 17.37 28.96
P8KEPF - 5902	3	Suspect Unknown Unknown		51.50 18.24 30.27
P8TP7Q - 5901	3	Suspect Unknown Unknown		45.00 37.00 18.00
Q99BQE - 5901	3	Suspect Unknown Unknown	1.01 0.3460 0.6550	5,021.00 17.20 32.59

Item 4 Results				
WebCode-Test	Number of Contributors	Contributor(s)	DNA Concentration (ng/uL)	DNA Proportion (%)
R9RDYE - 5901	3	Suspect Unknown Unknown		51.00 18.00 31.00
RCPXZD - 5901	3	Suspect Unknown Unknown	1.23 0.8700 0.9000	41.00 29.00 30.00
TMDEV4 - 5902	3	Suspect Unknown Unknown	1.35 0.6265 0.4264	56.16 26.08 17.75
UNVR99 - 5902	3	Suspect Unknown Unknown		
VFHZHA - 5902	3	Suspect Unknown Unknown		49.39 22.24 28.38
VK9834 - 5902	3	Suspect Unknown Unknown	8.96 3.14 4.76	53.14 18.63 28.22
W2RLKZ - 5901	3	Suspect Unknown Unknown		46.00 16.00 39.00
WRGKYZ - 5901	3	Suspect Unknown Unknown		55.00 16.00 29.00
X28843 - 5901	3	Suspect Unknown Unknown	2.03 1.34 0.5646	51.59 34.08 14.33
YGJ6L9 - 5901	3	Suspect Unknown Unknown		44.00 33.00 22.00
Z8LUQ4 - 5902	3	Suspect Unknown Unknown		

Response Summary	y: Item 4	
Estimated Number of Contributors	Percent	Reported
3	66	(99%)
4	1	(1%)

Statistical Analysis for Item 3

TABLE 7			
WebCode- Test	Item 3 Methods & Results		
2QGMC2 - 5902	Method(s): Likelihood Ratio Stats Analysis: A DNA match was identified between the data obtained from the first fraction from the questioned stain from victim's underwear (item 3) and the male suspect. A match between the data obtained from the first fraction from the questioned stain from victim's underwear (item 3) and the male suspect is approximately fifty quadrillion times more probable than a coincidental match to an unrelated person in the population. A DNA match was identified between the data obtained from the first fraction from the questioned stain from victim's underwear (item 3) and the female victim. A match between the data obtained from the first fraction from the questioned stain from victim's underwear (item 3) and the female victim is approximately nine hundred trillion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the second fraction from the questioned stain from victim's underwear (item 3) and the suspect is approximately three undecillion times less probable than a coincidental match to an unrelated person in the population. The suspect is excluded from the data obtained from the second fraction from the second fraction from the questioned stain from victim's underwear (item 3) and the victim is approximately seven octillion times less probable than a coincidental match to an unrelated person in the population. The suspect is excluded from the data obtained from the second fraction from the second fraction from the questioned stain from victim's underwear (item 3) and the victim is approximately seven octillion times less probable than a coincidental match to an unrelated person in the population. The victim is excluded from the data obtained from the second fraction from the questioned stain from victim's underwear (item 3). DNA match between the data obtained from the questioned stain from victim is excluded from the data obtained from the second fraction from the questioned stain from victim's underwea		
34G3B3 - 5902	 Method(s): Likelihood Ratio Stats Analysis: Assuming three contributors to the non-sperm fraction of item 3, the evidence profile is 1.2E+24 times more likely to be observed if Female Victim and two unknown individuals are the contributors than if three unknown individuals are the contributors. Assuming three contributors to the non-sperm fraction of item 3, the evidence profile is 2.0E+27 times more likely to be observed if Female Victim and two unknown individuals are the contributors. Assuming three contributors to the non-sperm fraction of item 3, the evidence profile is 2.0E+27 times more likely to be observed if Female Victim, Male Suspect, and one unknown individual are the contributors than if Female Victim and two unknown individuals are the contributors. Database(s) Used: NIST Databases - Forensic Sci Int.: Genetics 31 (2017) e36-e40 		
37J2CU - 5901	Method(s): Likelihood Ratio Stats Analysis: [Participant did not return statistical analysis.] Database(s) Used: [Location Identifying Database]		
3MQZUY - 5902	 Method(s): Likelihood Ratio Stats Analysis: 3. The SUSPECT (item 2-1) cannot be excluded as a contributor to Mixture 1 (see Testing Summary) from the VICTIM's underwear (item 3-1). The STR DNA results are estimated to be greater than one trillion times more likely if they originate from the VICTIM, SUSPECT and one unknown person than if they originate from the VICTIM and two unknown people unrelated to the SUSPECT. 5. The SUSPECT (item 2-1) also cannot be excluded as the source of a male-specific DNA profile (Y-STR Profile B; see Testing Summary) from the VICTIM's underwear (item 3-1) The Y-STR DNA results are further estimated to be 2000 times more likely if it originates from the SUSPECT than if it originates from an unknown male, unrelated to him. Note: Close paternal male relatives of SUSPECT may also not be excluded. Database(s) Used: i. NIST Asian, ii. NIST African American, iii. NIST Caucasian, iv. Srivastava et al. (2019) South Asian, v. [Location Identifying Database] 		

	TABLE 7			
WebCode-	Item 3 Methods & Results			
Test				
3WHFJ3 - 5902	 Method(s): Likelihood Ratio Stats Analysis: H1: The evidence originated from the Known Female, Known Male, and one unknown, unrelated individual. H2: The evidence originated from the Known Female and two unknown, unrelated individuals. The DNA profile obtained from Item 3 is approximately 320 quintillion times more likely if it originated from the Known Female, Known Male and one unknown, unrelated individual than if it originated from the Known Female and two unknown, unrelated individual than if it originated from the Known Female and two unknown, unrelated individuals. There is evidentiary support for the inclusion of the Known Male as a possible contributor to the DNA profile obtained as compared with the alternate explanation of the evidence. Database(s) Used: NIST revised: African American, Asian, Caucasian, Hispanic 			
44PDRD - 5901	Method(s): Likelihood Ratio Stats Analysis: Item 3.1 (Portion of "Questioned stain from victim's underwear" - Fraction 2): The DNA profile from this item was interpreted as a mixture of three individuals, with at least one male contributor and with FEMALE VICTIM as an assumed contributor. The DNA results are approximately 74.8 quintillion times more likely if they originated from MALE SUSPECT, FEMALE VICTIM and an unknown, unrelated individual than if they originated from FEMALE VICTIM and two unknown, unrelated individuals. Based on the likelihood ratio, this provides very strong support that MALE SUSPECT is a contributor to the DNA from this item. Item 3.1 (Portion of "Questioned stain from victim's underwear" - Fraction 1): The male DNA profile from this item was interpreted as a single-source profile. FEMALE VICTIM and MALE SUSPECT are excluded as possible contributors to the DNA from this item. Database(s) Used: NIST 1036 July 2017			
4BEUJW - 5902	 Method(s): Likelihood Ratio Stats Analysis: Sperm fraction: The Victim (Item 1) is Excluded as being the source of the DNA in this sample. The Suspect (Item 2) is Excluded as being the source of the DNA in this sample. Non-sperm fraction/Epithelial: Victim (Item 1): Assumed contributor Suspect (Item 2): The DNA evidence is estimated to be 100 billion times more likely to occur if the DNA originated from the victim and the suspect and one unknown, unrelated person selected randomly from the [Population], than if it originated from the victim and two unknown and unrelated people selected randomly from the [Population]. Database(s) Used: [Location Identifying Database] 			
4FBJY4 - 5902	Method(s): Likelihood Ratio Stats Analysis: Item 1: 5.754e+11. Item 2: 4.377e+22. Unknown male from sperm fraction: 8.087e+11. Database(s) Used: [Location Identifying Database]			
4UQG64 - 5901	Method(s): Likelihood Ratio Stats Analysis: Assuming Female Victim (K1), the evidence profile 3A-3PTNS is approximately 130 trillion times more likely if it originated from Female Victim (K1), Male Suspect (K2), and an unknown individual than if it originated from Female Victim (K1) and two unknown individuals. This provides very strong support for inclusion of Male Suspect (K2). The evidence DNA profile 3A-3PTSP is approximately 0 times more likely if it originated from Female Victim (K1) than if it originated from an unknown individual. This provides support for exclusion of Female Victim (K1). The evidence DNA profile 3A-3SP is approximately 0 times more likely if it originated from Male Suspect (K2) than if it originated from an unknown individual. This provides support for exclusion of Male Suspect (K2). Database(s) Used: 2015 FBI Expanded Population Database			

TABLE 7			
WebCode-	Item 3 Methods & Results		
Test			
4XQRZ4 - 5901	 Method(s): Likelihood Ratio Stats Analysis: For 3 e: Assuming Female Victim(K1), the evidence profile is approximately 234 trillion times more likely if it originated from Female Victim (K1), Male Suspect (K2) and an unknown individual than if it originated from Female Victim (K1) and two unknown individuals. This analysis provides very strong support that Male Suspect (K2) is a contributor to the DNA obtained from this sub-item. For 3 sp: Female Victim (K1) is excluded based on the LR of zero. Male Suspect (K2) is excluded based on the LR of zero. Database(s) Used: 2015 FBI Expanded Population Database 		
62K9Q4 - 5901	 Method(s): Likelihood Ratio Stats Analysis: For 3A-3NS: Assuming (K1), the evidence profile is approximately 115 trillion times more likely if it originated from Female Victim (K1), Male Suspect (K2), and an unknown individual than if it originated from Female Victim (K1) and two unknown individuals. This analysis provides very strong support for the proposition that Male Suspect (K2) is a contributor to the DNA obtained from this sub-item. For 3A-3SP: The evidence profile is approximately 0 times more likely if it originated from Female Victim (K1) than if it originated from an unknown individual. Female Victim (K1) is excluded based on the LR of zero. The evidence profile is approximately 0 times more likely if it originated from Male Suspect (K2) than if it originated from an unknown individual. Male Suspect (K2) is excluded based on the LR of zero. Database(s) Used: 2015 FBI Expanded Population 		
63VBC2 - 5901	Method(s): Likelihood Ratio Stats Analysis: For 3e: Assuming Female Victim (K1), the evidence profile is approximately 2 trillion times more likely if it originated from Female Victim (K1), Male Suspect (K2), and one unknown individual than if it originated from Female Victim (K1) and two unknown individuals. This analysis provides very strong support for the proposition that Male Suspect (K2) is included as a contributor to this DNA mixture. For 3sp: The evidence DNA profile is approximately 0 times more likely if it originated from Female Victim (K1) than if it originated from one unknown individual. The evidence DNA profile is approximately 0 times more likely if it originated from Male Suspect (K2) than if it originated from one unknown individual. Database(s) Used: 2015 FBI Expanded Population Database		

TABLE 7			
WebCode- Test	Item 3 Methods & Results		
6PTH6Y - 5902	 Method(s): Likelihood Ratio Stats Analysis: The first fraction from the victim's underwear from item 3 contained a mixture of DNA from at least three individuals. A DNA match was identified between the data obtained from the first fraction from the victim's underwear from item 3 and the victim. A match between the data obtained from the first fraction from the victim's underwear from item 3 and the victim is approximately one sextillion times more probable than a coincidental match to an unrelated person in the population. A DNA match was identified between the data obtained from the first fraction from the victim's underwear from item 3 and the suspect. A match between the data obtained from the first fraction from the victim's underwear from item 3 and the suspect is approximately eighty quadrillion times more probable than a coincidental match to an unrelated person in the population. The DNA obtained from the second fraction from the victim's underwear from item 3 and the victim is approximately eight quadrillion times more probable than a coincidental match to an unrelated person in the second fraction from the victim's underwear from item 3 and the victim is approximately four nonillion times less probable than a coincidental match to an unrelated person in the population. The DNA obtained from the support from the second fraction from the victim's underwear from item 3 and the victim is approximately four nonillion times less probable than a coincidental match to an unrelated person in the population. The Victim is excluded from the data obtained from the second fraction from the victim's underwear from item 3 and the suspect is approximately six hundred undecillion times less probable than a coincidental match to an unrelated person in the population. The victim's underwear from item 3 and the suspect is approximately six hundred undecillion times less probable than a coincidental match to an unrelated person in the population. The Second fraction from the victim's underwear from item		
77A6RY - 5901	Method(s) : Likelihood Ratio Stats Analysis: 3.0457E19 Database(s) Used: The FBI's extended database for African Americans was used.		
786NUZ - 5901	 Method(s): Likelihood Ratio Stats Analysis: The DNA profile obtained from epithelial fraction of item # 3 is a mixture of at least 3 individuals. Following set of hypotheses were evaluated: A) Hp; item No. 1+2 unknown, Hd; 3 unknown, LR (MLE) = 30555. B) HP; Item No. 1+item No. 2 + 1 unknown, Hd; Item No. 1+ 2 Unknown, LR (MLE)= 1.373 e +18. C) Hp; Item No. 1+ Item No. 2+1 unknown, Hd; Item No. 1+ item No. 2 + 1 unknown, Hd; Item No. 1+ item No. 2 + 1 unknown, Hd; Item No. 1+ item No. 2 + 1 unknown, Hd; Item No. 1+ item No. 2 + 1 unknown, LR (MLE)=6.307 e +14 The DNA profile obtained from sperm fraction of item #3 is a mixture of at least 3 individuals. Following set of hypotheses were evaluated: A) Hp; item No. 1+2 unknown, Hd; 3 unknown, LR (MLE) = 1.421 e +8. B) Hp; item No.1+ item No. 2 + 1 unknown, Hd; Item #1 +2 unknown, LR (MLE) = 7.638 e +12. C) Hp; Item No. 1+ Item No. 2+1 unknown, Hd; Item No. 1+ item No. 2 + 1 unknown LR(MLE)=2.150 e +18 Database(s) Used: NIST Caucasian database 2017 with theta co-ancestry coefficient = 0.03 		

TABLE 7		
WebCode-	Item 3 Methods & Results	
Test		
7H4JRV - 5902	 Method(s): Likelihood Ratio Stats Analysis: A DNA match was identified between the data obtained from the first fraction from the questioned stain from victim's underwear (item 3) and the victim. A match between the data obtained from the first fraction from the questioned stain from victim's underwear (item 3) and victim is approximately twenty trillion times more probable than a coincidental match to an unrelated person in the population. A DNA match was identified between the data obtained from the first fraction from the questioned stain from victim's underwear (item 3) and the subject. A match between the data obtained from the first fraction from the questioned stain from victim's underwear (item 3) and subject is approximately seven hundred trillion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the second fraction from the questioned stain from victim's underwear (item 3) and the victim is approximately four nonillion times less probable than a coincidental match to an unrelated person in the population. The victim is excluded from the data obtained from the second fraction from the questioned stain from victim's underwear (item 3) and the victim is approximately four nonillion times less probable than a coincidental match to an unrelated person in the population. The victim is excluded from the data obtained from the second fraction from the questioned stain from victim's underwear (item 3). A DNA match between the data obtained from the second fraction from the questioned stain from victim's underwear (item 3). and the suspect is approximately six hundred undecillion times less probable than a coincidental match to an unrelated person in the population. The suspect is excluded from the data obtained from the data obtained from the questioned stain from victim's underwear (item 3). Database(s) Used: The National Institute of Standards and Technology's 1036 Revised U.S. combined popula	
8DDAHX - 5901	 Method(s): Likelihood Ratio Stats Analysis: The DNA profile of the sperm fraction was interpreted as originating from a single unknown male A. The DNA profile of the epithelial fraction was interpreted as a mixture of three individuals. Known Female Victim (1) was assumed as a contributor. The DNA profile is at least 1 trillion times more likely if it originated from Known Female Victim (1), Known Male Suspect (2), and one unknown individual than if it had originated from Known Female Victim (1) and two unknown, unrelated individuals. Statistical analysis provides very strong support for the inclusion of Known Male Suspect (2). Database(s) Used: FBI Extended CODIS core allele frequencies - Caucasian, African American, and Southwest Hispanic 	
8Y8DYV - 5902	 Method(s): Likelihood Ratio Stats Analysis: A DNA match was identified between the data obtained from the "questioned stainunderwear" (item #3) and the male suspect (item #2). A match between the data obtained from the "questioned stainunderwear" (item #3) and the male suspect (item #2) is approximately 2 sextilion times more probable than a coincidental match to an unrelated person in the population. A DNA match was identified between the data obtained from the "questioned stainunderwear" (item #1). A match between the data obtained from the "questioned stainunderwear" (item #3) and the female victim (item #1). A match between the data obtained from the "questioned stainunderwear" (item #3) and the female victim (item #1) is approximately 40 thousand times more probable than a coincidental match to an unrelated person in the population. Database(s) Used: The National Institute of Standards and Technology's 1036 Revised U.S. combined population database was used to generate DNA match statistics. The combined populations. Identical siblings will have identical DNA data and DNA match statistics. 	

TABLE 7			
WebCode-	Item 3 Methods & Results		
Test			
8YUL9Z - 5901	Method(s): Likelihood Ratio Stats Analysis: A likelihood ratio (LR) analysis was performed using the LRmix Studio software, using allele frequency data specific to the [Population]- based on own laboratory reference population database consisting of 5,000 genotypes from [Country]. The LR values for mixed DNA profiles were calculated with consideration for both drop-in and drop-out phenomena. The analysis assumed a drop-in probability of 0.05 and applied a minimum allele frequency threshold of 0.001. A subpopulation correction factor of 0.01 was also incorporated into the calculations. The resulting LR values were interpreted in terms of their probative value. The evaluated hypotheses included: Prosecution Hypothesis (Hp): The DNA mixture includes the genetic profiles of item1, item2, and one unknown, untested individual from the population. Alternative Hypothesis (Hd): The DNA mixture includes the genetic profile of item1 and two unknown, untested individuals from the population. The comparison of these hypotheses was used to assess the strength of the genetic evidence in the context of the case. Likelihood ratio on this sample is: 1.4024E12. Y-STR: Given the Haplotypes above and that there is 1 additional unknown contributor, the likelihood of observing the given trace under the hypothesis of the donorship of the suspect is approx. 135,377 times more likely than observing the given trace under the hypothesis of the non-donorship. This calculation is based on 298,405 haplotypes (Y17) in the worldwide database YHRD R69. Database(s) Used: STR laboratory population base of [Country]		
9CLYK9 - 5901	 Method(s): Likelihood Ratio Stats Analysis: Item 3.1 (Portion of "Questioned stain from victim's underwear" -Fraction 1): The male DNA profile from this item was interpreted as a single-source profile. The female victim and the male suspect are excluded as possible contributors to the DNA from this item. Item 3.1 (Portion of "Questioned stain from victim's underwear" - Fraction 2): The DNA profile from this item was interpreted as a mixture of three individuals, with at least one male contributor and with the female victim as an assumed contributor. The DNA results are approximately 2.32 nonillion times more likely if they originated from the female victim, the male suspect, and an unknown, unrelated individuals. Based on the likelihood ratio, this provides very strong support that the male suspect is a contributor to the DNA from this item. Database(s) Used: NIST 1036 July 2017 		
9MBR8Q - 5901	Method(s): Likelihood Ratio Stats Analysis: sp fraction - Item 1 (complainant) - Assumed contributor. Item 2 (known suspect) - >100 billion in favour of H1 (favouring contribution). e fraction - Item 1 (complainant) - Assumed contributor. Item 2 (known suspect) - >100 billion in favour of H1 (favouring contribution). Database(s) Used: [Location Identifying Database]		
9WLTTR - 5902	Method(s): Likelihood Ratio Stats Analysis: 3e: The DNA evidence is 30 billion times more likely if the suspect is a contributor. The DNA evidence is 4.9 billion times more likely if the victim is a contributor. 3sp: The DNA evidence is 3600 times more likely if the suspect is a contributor. The DNA evidence is 440 times more likely if the victim is a contributor. Database(s) Used: [Location Identifying Database]		
B7W38V - 5901	Method(s): Likelihood Ratio Stats Analysis: 1.2896E18 Database(s) Used: FBI Extended African American		

WebCode-	Item 3 Methods & Results
Test	
BDR7Z7 - 5901	 Method(s): Likelihood Ratio Stats Analysis: Fraction 1 of differential: Explanation 1: The DNA profile obtained originated from item 1 (1C), item 2 (1D), and an unknown, unrelated individual. Explanation 2: The DNA profile obtained originated from item 1 (1C) and two unknown, unrelated individuals. The DNA profile obtained is approximately 2E18 times more likely (very strong support) if it originated from the item 1 (1C), item 2 (1D), and an unknown, unrelated individual than if the DNA profile obtained originated from item 1 (1C) and two unknown, unrelated individuals. Fraction 2 of differential: No LR reported. Item 1 (1C) and item 2 (1D) were excluded. Ran STRmix for prob gen proportions only. Database(s) Used: Nist1036 (revised 2017)_Caucasian, Nist1036 (revised 2017)_African American, Nist1036 (revised 2017)_Hispanic
BKHHBU - 5901	Method(s): Likelihood Ratio Stats Analysis: 1.7158e27
	Database(s) Used: FBI Extended Atrican American
BNGT8U - 5902	Method(s): Likelihood Ratio Stats Analysis: Assuming 3 contributors for the non-sperm fraction of item 3, the evidence profile is 5.8 x 10^25 times more likely to be observed if the victim and two unknowns are the contributors than if three unknowns are the contributors. Assuming 3 contributors for the non-sperm fraction of item 3, the evidence profile is 6.2 x 10^20 times more likely to be observed if the suspect, victim, and one unknown are the contributors than if the victim and two unknowns are the contributors. Database(s) Used: NIST Databases - Forensic Sci. Int.: Genetics 31 (2017) e36–e40
CDTG9M - 5901	 Method(s): Likelihood Ratio, STRmix Stats Analysis: Epithelial Fraction Propositions for contributor 1: The propositions considered for ITEM 3 were: P1: The SUSPECT is one of three contributors to the mixed DNA profile obtained. P2: The SUSPECT is NOT a contributor to the mixed DNA profile obtained. Three unknown individuals, unrelated to the SUSPECT, are the contributors. For this calculation, it was assumed that three individuals have contributed to the DNA profile obtained. Likelihood Ratio: The evidence is at least 100 BILLION (2.9563E20) times more likely if the SUSPECT is one of three contributors to the mixed DNA profile than if the profile originated from three unknown individuals, unrelated to the SUSPECT, selected at random from the [Population]. Propositions for contributors to the mixed DNA profile obtained. P2: The VICTIM is one of three contributors to the mixed DNA profile obtained. P2: The VICTIM is NOT a contributor to the mixed DNA profile obtained. Three unknown individuals, unrelated to the VICTIM, are the contributors. For this calculation, it was assumed that three individuals have contributed to the DNA profile obtained. Likelihood Ratio: The evidence is at least 9 BILLION (9.2033E9) times more likely if the VICTIM is one of three unknown individuals, unrelated to the VICTIM, selected at random from the [Population]. Database(s) Used: [Location Identifying Database]

TABLE 7	
WebCode- Test	Item 3 Methods & Results
CKECAQ - 5902	 Method(s): Likelihood Ratio Stats Analysis: A DNA match was identified between the data obtained from the first fraction from the questioned stain from victim's underwear (Item 3) and the Victim. A match between the data obtained from the first fraction from the questioned stain from victim's underwear (Item 3) and the Victim is approximately 60 quadrillion times more probable than a coincidental match to an unrelated person in the population. A DNA match was identified between the data obtained from the first fraction from the questioned stain from victim's underwear (Item 3) and the Subject. A match between the data obtained from the first fraction from the questioned stain from victim's underwear (Item 3) and the Subject is approximately 700 quintillion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the subject is approximately 600 undecillion times less probable than a coincidental match to an unrelated person in the questioned stain from victim's underwear (Item 3) and the Subject is approximately 600 undecillion times less probable than a coincidental match to an unrelated person in the population. The Subject is excluded from the data obtained from the second fraction from the questioned stain from victim's underwear (Item 3). A DNA match between the data obtained from the second fraction from the questioned stain from victim's underwear (Item 3) and the Victim is approximately 400 nonillion times less probable than a coincidental match to an unrelated person in the population. The Victim is excluded from the data obtained from the second fraction from the questioned stain from victim's underwear (Item 3). Database(s) Used: All evidence genotypes were compared with all reference genotypes to compute likelihood ratio (LR) DNA match statistics. The National Institute of Standards and Technology's 1036 Revised U.S. combined population database was used to generate DNA match statistics. The combined
CRBAWU - 5901	Method(s): Likelihood Ratio Stats Analysis: The DNA evidence is at least one billion times more likely if the mixture of DNA originated from the Victim, the Suspect and an unknown unrelated individual rather than if it originated from the Victim, and two unknown, unrelated males Database(s) Used: [Location Identifying Database]
CRVQCW - 5901	Method(s): Likelihood Ratio Stats Analysis: 3A-1PT NS: Assuming Female Victim (K1), the evidence profile is 196 Trillion times more likely if it originated from Female Victim (K1), Male Suspect (K2) and an unknown individual than if it originated from Female Victim (K1) and two unknown individuals. This provides very strong support for inclusion. 3A-1PT SP: Male Suspect (K2) is excluded based on the LR of zero. Database(s) Used: 2015 FBI Expanded Population Database

TABLE 7	
WebCode- Test	Item 3 Methods & Results
CXYRCQ - 5902	 Method(s): Likelihood Ratio Stats Analysis: A DNA match was identified between the data obtained from the first fraction from the questioned stain from victim's underwear (Item 3) and Suspect. A match between the data obtained from the first fraction from the questioned stain from victim's underwear (Item 3) and Suspect is approximately fifty quadrillion times more probable than a coincidental match to an unrelated person in the population. A DNA match was identified between the data obtained from the first fraction from the questioned stain from victim's underwear (Item 3) and Victim. A match between the data obtained from the first fraction from the questioned stain from victim's underwear (Item 3) and Victim is approximately two trillion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the second fraction from the questioned stain from victim's underwear (Item 3) and Victim is approximately two trillion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the second fraction from the questioned stain from victim's underwear (Item 3) and Victim is approximately four nonillion times less probable than a coincidental match to an unrelated person in the population. Victim is excluded from the data obtained from the second fraction from the questioned stain from victim's underwear (Item 3) and Suspect is approximately six hundred undecillion times less probable than a coincidental match to an unrelated person in the population. Suspect is excluded from the data obtained from the second fraction from the questioned stain from victim's underwear (Item 3). Database(s) Used: The National Institute of Standards and Technology's 1036 Revised U.S. combined population database was used to generate DNA match statistics. The combined population database contains African American, Caucasian, Hispanic and Asian populations. Identical sibling
CY6DP6 - 5901	 Method(s): Likelihood Ratio Stats Analysis: Fraction 1: The male DNA profile from this item was interpreted as a single-source profile. The female victim and the male suspect are excluded as possible contributors to this DNA profile. Fraction 2: The DNA profile from this item was interpreted as a mixture of three individuals, with at least one male contributor and with the female victim as an assumed contributor. The DNA results are approximately 27.4 quintillion times more likely if they originated from the female victim, the male suspect, and an unknown, unrelated individual than if they originated from the female victim, and two unknown, unrelated individuals. Based on the likelihood ratio, this provides very strong support that the male suspect is a contributor to the DNA from this item. Database(s) Used: NIST 1036 July 2017
CZ6EPV - 5901	Method(s): Likelihood Ratio Stats Analysis: Item 3e: Assuming Female Victim (K1), the evidence profile is approximately 359 Trillion times more likely if it originated from the female victim (K1), male suspect (K2), and one unknown individual than if it originated from the female victim (K1) and two unknown individuals. This analysis provides very strong support for the proposition that the Male Suspect (K2) is a contributor to this mixture. Item 3sp: Female Victim (K1) is excluded based on the LR of zero. Male Suspect (K2) is excluded based on the LR of zero. Database(s) Used: 2015 FBI Expanded Population Database

TABLE 7	
WebCode-	Item 3 Methods & Results
Test	
E34ANN - 5902	Method(s): Likelihood Ratio Stats Analysis: A DNA match was identified between the data obtained from the first fraction from the stain from underwear (item 3) and Victim. A match between the data obtained from the first fraction from the stain from underwear (item 3) and Victim is approximately 5 billion times more probable than a coincidental match to an unrelated person in the population. A DNA match was identified between the data obtained from the first fraction from the stain from underwear (item 3) and Suspect. A match between the data obtained from the first fraction from the stain from underwear (item 3) and Suspect is approximately 10 quintillion times more probable than a coincidental match to an unrelated person in the population. A DNA match was identified between the data obtained from the second fraction from the stain from underwear (item 3) and Suspect is approximately 10 quintillion times more probable than a coincidental match to an unrelated person in the population. A DNA match was identified between the data obtained from the second fraction from the stain from underwear (item 3) and Suspect is approximately 100 thousand times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the second fraction from the stain from underwear (item 3) and Victim is approximately 100 nonillion times less probable than a coincidental match to an unrelated person in the population. Victim is excluded from the data obtained from the second fraction from the stain from underwear (item 3). Database(s) Used: NIST1036
EHXZJK - 5902	Method(s): Likelihood Ratio Stats Analysis: Sp: PP21 Item 1: Excluded. Item 2: Excluded. YFP Item 2: Excluded. E: PP21 Item 1: Assumed contributor. Item 2: Not excluded (LR = 100 billion). YFP Item 2: Not excluded (LR = 13,000). Database(s) Used: [Location Identifying Database]
ELJJPP - 5902	 Method(s): Likelihood Ratio Stats Analysis: A DNA match was identified between the data obtained from the first fraction of the stain from victim's underwear (item 3) and the male suspect (item 2). A match between the data obtained from the first fraction of the stain from victim's underwear (item 3) and the male suspect (item 2) is approximately 900 septillion times more probable than a coincidental match to an unrelated person in the population. A DNA match was identified between the data obtained from the first fraction of the stain from victim's underwear (item 3) and the female victim (item 1). A match between the data obtained from the first fraction of the stain from victim's underwear (item 3) and the female victim (item 1) is approximately 2 quadrillion times more probable than a coincidental match to an unrelated person in the population of the stain from victim's underwear (item 3) and the male suspect (item 2) is approximately 30 undecillion times less probable than a coincidental match to an unrelated person in the population. The male suspect (item 2) is excluded from the data obtained from the second fraction of the stain from victim's underwear (item 3) and the male suspect (item 2) is approximately 30 undecillion times less probable than a coincidental match to an unrelated person in the population. The male suspect (item 2) is excluded from the data obtained from the second fraction of the stain from victim's underwear (item 3). A DNA match between the data obtained from the second fraction of the stain from victim's underwear (item 3). and the female victim (item 1) is approximately 2 nonillion times less probable than a coincidental match to an unrelated person in the population. The female victim (item 1) is excluded from the data obtained from the second fraction of the stain from victim's underwear (item 3). and the female victim (item 1) is approximately 2 nonillion times less probable than a coincidental match to an unrelated person in the population.

TABLE 7	
WebCode-	Item 3 Methods & Results
Test	
EY6YTN - 5902	 Method(s): Likelihood Ratio Stats Analysis: The first fraction of the "victim's underwear" (item 3) contained a mixture of DNA from at least three individuals. A DNA match was identified between the data obtained from the first fraction of the "victim's underwear" (item 3) and the suspect. A match between the data obtained from the first fraction of the "victim's underwear" (item 3) and the suspect. A match between the data obtained from the first fraction of the "victim's underwear" (item 3) and the suspect is approximately 10 septillion times more probable than a coincidental match to an unrelated person in the population. A DNA match was identified between the data obtained from the first fraction of the "victim's underwear" (item 3) and the victim is approximately 5 quadrillion times more probable than a coincidental match to an unrelated person in the population of the "victim's underwear" (item 3) and the victim is approximately 5 quadrillion times more probable than a coincidental match to an unrelated person in the population. The second fraction of the "victim's underwear" (item 3) contained a mixture of DNA from at least two individuals. A DNA match between the data obtained from the second fraction of the "victim's underwear" (item 3) and the suspect is approximately 1 undecillion times less probable than a coincidental match to an unrelated person in the population. The suspect is excluded from the data obtained from the second fraction of the "victim's underwear" (item 3). A DNA match between the data obtained from the second fraction of the "victim's underwear" (item 3). A DNA match between the data obtained from the second fraction of the "victim's underwear" (item 3) and the suspect is approximately 1 undecillion times less probable than a coincidental match to an unrelated person in the population. The suspect is excluded from the second fraction of the "victim's underwear" (item 3) and the suspect is approximately 20 nonillion times less pr
F7H8UP - 5902	Method(s): Likelihood Ratio Stats Analysis: H1: The evidence originated from the victim, suspect and one unknown, unrelated individual. H2: The evidence originated from victim and two unknown, unrelated individuals. The DNA profile obtained from the non-sperm cell fraction from the victim's underwear is approximately 1.3 septillion times more likely if it originated from the victim, suspect and one unknown, unrelated individual than if it originated from the victim and two unknown, unrelated individuals. There is evidentiary support for the inclusion of the suspect as a possible contributor to the DNA profile obtained as compared with the alternate explanation of the evidence. Database(s) Used: NIST revised: African American, Asian, Caucasian, Hispanic
F8XQDM - 5902	 Method(s): Likelihood Ratio Stats Analysis: A DNA match was identified between the data obtained from the first fraction from the questioned stain from victim's underwear (item 3) and Victim. A match between the data obtained from the first fraction from the questioned stain from victim's underwear (item 3) and Victim is approximately 100 quintillion times more probable than a coincidental match to an unrelated person in the population. A DNA match was identified between the data obtained from the first fraction from the questioned stain from victim's underwear (item 3) and Suspect. A match between the data obtained from the first fraction from the questioned stain from victim's underwear (item 3) and Suspect is approximately 100 quintillion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the second fraction from the questioned stain from victim's underwear (item 3) and Victim is approximately 400 nonillion times less probable than a coincidental match to an unrelated person in the population. Victim is excluded from the data obtained from the second fraction from the questioned stain from victim's underwear (item 3) and Suspect is approximately 600 undecillion times less probable than a coincidental match to an unrelated person in the population. Suspect is excluded from the data obtained from the second fraction from the questioned stain from victim's underwear (item 3) and Suspect is approximately 600 undecillion times less probable than a coincidental match to an unrelated person in the population. Suspect is excluded from the data obtained from the second fraction from the questioned stain from victim's underwear (item 3). Database(s) Used: National Institute of Standards and Technology's 1036 Revised U.S. combined population database

WebCode-	Item 3 Methods & Results
Test	
FPXE6L - 5902	 Method(s): Likelihood Ratio Stats Analysis: 1. The SUSPECT (item Q02-01) cannot be excluded as a contributor to the following (see Testing Summary): • Mixture 1 and Y-STR Profile B from a blood and semen stained area on the underwear (item Q03-1) from the VICTIM. The STR DNA results for Mixture 1 are estimated to be greater than one trillion times more likely if they originate from the VICTIM, the SUSPECT and the donor of STR Profile 1 than if they originate from the VICTIM, the donor of STR Profile 1 and one unknown person unrelated to the SUSPECT. Note: A supplementary statistic can be calculated for the male-specific Y-STR profile that adds further weight to the association reported. Database(s) Used: i. NIST Asian, ii. NIST African American, iii. NIST Caucasian, iv. Srivastava et al. (2019) South Asian, v. [Location Identifying Database]
GCH7HK - 5902	 Method(s): Likelihood Ratio Stats Analysis: A DNA match was identified between the data obtained from the first fraction from the questioned stain from victim's underwear (item 3) and the victim. A match between the data obtained from the first fraction from the questioned stain from victim's underwear (item 3) and the victim is approximately 50 septillion times more probable than a coincidental match to an unrelated person in the population. A DNA match was identified between the data obtained from the first fraction from the questioned stain from victim's underwear (item 3) and the suspect. A match between the data obtained from the first fraction from the questioned stain from victim's underwear (item 3) and the suspect. A match between the data obtained from the first fraction from the questioned stain from victim's underwear (item 3) and the suspect is approximately 200 quintillion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the second fraction from the questioned stain from victim's underwear (item 3) and the victim is approximately 400 nonillion times less probable than a coincidental match to an unrelated person in the population. The victim is excluded from the data obtained from the second fraction from the questioned stain from victim's underwear (item 3). A DNA match between the data obtained from the second fraction from the questioned stain from victim's underwear (item 3) and the suspect is approximately 600 undecillion times less probable than a coincidental match to an unrelated person in the population. The suspect is excluded from the data obtained from the second fraction from the questioned stain from victim's underwear (item 3). Database(s) Used: National Institute of Standards and Technology's 1036 Revised U.S. combined population
H484WJ - 5902	 Method(s): Likelihood Ratio Stats Analysis: A DNA match was identified between the data obtained from the first fraction from the tan fabric (item 3) and the male subject. A match between the data obtained from the first fraction from the tan fabric (item 3) and the male subject is approximately 100 sextillion times more probable than a coincidental match to an unrelated person in the population. A DNA match was identified between the data obtained from the first fraction from the tan fabric (item 3) and the data obtained from the first fraction from the tan fabric (item 3) and the female victim. A match between the data obtained from the first fraction from the tan fabric (item 3) and the female victim is approximately 7 billion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the first fraction from the tan fabric (item 3) and the female victim is approximately 7 billion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the farst fraction from the tan fabric (item 3) and the female victim is approximately 400 nonillion times less probable than a coincidental match to an unrelated person in the population. The female victim is excluded from the data obtained from the second fraction from the tan fabric (item 3). A DNA match between the data obtained from the second fraction from the tan fabric (item 3) and the male subject is approximately 100 undecillion times less probable than a coincidental match to an unrelated person in the tan fabric (item 3) and the male subject is approximately 100 undecillion times less probable than a coincidental match to an unrelated person in the population. The male subject is excluded from the data obtained from the second fraction from the tan fabric (item 3) and the male subject is approximately 100 undecillion times less probable than a coincidental match to an unrelated person in the population.

TABLE 7	
WebCode-	Item 3 Methods & Results
Test	
HEQHYL - 5902	 Method(s): Likelihood Ratio Stats Analysis: A DNA match was identified between the data obtained from the first fraction from the victim's underwear (item 3) and the suspect. A match between the data obtained from the first fraction from the victim's underwear (item 3) and the suspect is approximately 3 sextillion times more probable than a coincidental match to an unrelated person in the population. The comparison between the victim and the data obtained from the first fraction from the victim's underwear (item 3) is inconclusive because reproducible match statistics were not obtained. Therefore, no conclusions will be offered regarding the victim as a possible contributor to the DNA mixture obtained from the first fraction from the victim's underwear (item 3). A DNA match between the data obtained from the second fraction from the victim's underwear (item 3) and the suspect is approximately 600 undecillion times less probable than a coincidental match to an unrelated person in the population. The suspect is excluded from the data obtained from the second fraction from the data obtained from the second fraction from the victim's underwear (item 3) and the suspect is excluded from the data obtained from the second fraction from the victim is approximately 400 nonillion times less probable than a coincidental match to an unrelated person in the population. The victim is excluded from the data obtained from the second fraction from the victim's underwear (item 3). Database(s) Used: Revised NIST 1036
HJX7CN - 5901	Method(s) : Likelihood Ratio Stats Analysis: OVERALL LR=9.2694E11 BY ANALYSIS WITH LRmixStudio Database(s) Used: [Location Identifying Database]
J37QNH - 5902	Method(s): Likelihood Ratio Stats Analysis: Item 1 (complainant) Assumed contributor for both FF and MF (minor contributor). Item 2 (suspect) and FF. Not Excluded LR = 100 billion item 2 (suspect) and MF. Excluded Database(s) Used: [Location Identifying Database]
J772JG - 5902	Method(s): Likelihood Ratio Stats Analysis: PP21 MF/SP fraction: COMP (Item 1) Excluded; SUS (Item 2) Excluded. YFP MF/SP fraction: SUS (Item 2): Excluded. PP21 FF/E fraction: COMP (Item 1) Assumed Contributor, SUS (Item 2) Not Excluded (LR=100 billion). YFP FF/E fraction: SUS (Item 2) Not Excluded (LR can't be calculated as unresolved mixture). Database(s) Used: [Location Identifying Database]
JHLQHM - 5902	Method(s): Likelihood Ratio Stats Analysis: It is 4,3e18 times more likely to observe the DNA profile if the mixed stain on the underwear (ITEM 3) originates from ITEM 2 (Suspect) and two unknown persons, than if it originated from three unknown persons, unrelated to ITEM 2 (Suspect). Theta is 0.01 and probability of drop-in is 0.05. It is 1,8e13 times more likely to observe the DNA profile if the mixed stain on the underwear (ITEM 3) originates from ITEM 1 (Victim) and two unknown persons, than if it originated from three unknown persons, unrelated to ITEM 1 (Victim). Theta is 0.01 and probability of drop-in is 0.05. It is 7,8e21 times more likely to observe the DNA profile if the mixed stain on the underwear (ITEM 3) originates from unknown sperm donor (in sperm fraction) and two unknown persons, than if it originated from three unknown persons, unrelated to unknown sperm donor. Theta is 0.01 and probability of drop-in is 0.05. Database(s) Used: [Location Identifying Database]. Rare allele frequency is 0.0007

TABLE 7	
WebCode-	Item 3 Methods & Results
Test	
JPGUBX - 5901	Method(s): Likelihood Ratio Stats Analysis: Item 3e: The DNA profile from this item was interpreted as a mixture of three individuals, with at least two male contributors and with the female victim as an assumed contributor. The DNA results are approximately 2.60 nonillion times more likely if they originated from the female victim, the male suspect and an unknown, unrelated individual than if they originated from the female victim and two unknown, unrelated individuals. Based on the likelihood ratio, this provides very strong support that the male suspect is a contributor to the DNA from this item. Item 3sp: The DNA profile from this item was interpreted as a mixture of two individuals with at least one male contributor. The DNA results are approximately 46.5 quadrillion times more likely if they originated from two unknown, unrelated individuals than if they originated from the female victim and an unknown, unrelated individual. Based on the likelihood ratio, this provides very strong support that the female victim is not a contributor to the DNA from this item. The DNA results are approximately 2.66 million times more likely if they originated from the male suspect and an unknown, unrelated individual than if they originated from the male suspect individuals. Based on the likelihood ratio, this provides very strong support that the female victim is not a contributor to the DNA from the male suspect individuals. Based on the likelihood ratio, this provides very strong support that the male suspect is a contributor to the DNA from this item. Database(s) Used: NIST 1036 July 2017
K9ECCE - 5902	Method(s): Likelihood Ratio Stats Analysis: Item 3 - Sperm Fraction: The DNA evidence is 14 times more likely if Item 2 (Suspect) is a contributor. The DNA evidence is more likely if Item 1 (Victim) is not a contributor. Item 3 - Epithelial Fraction: The DNA evidence is 630 million times more likely if Item 2 (Suspect) is a contributor. The DNA evidence is 15 billion times more likely if Item 1 (Victim) is a contributor. Database(s) Used: [Location Identifying Database]
KPDQAK - 5902	Method(s): Likelihood Ratio Stats Analysis: NSF: H1 = The evidence originated from Victim, Suspect, and one unknown, unrelated individual. H2 = The evidence originated from Victim and two unknown, unrelated individuals. The DNA profile obtained from the non-sperm cell fraction of the underwear stain is approximately 1.4 quintillion times more likely if it originated from Victim, Suspect, and one unknown, unrelated individual than if it originated from Victim and two unknown, unrelated individuals. There is evidentiary support for the inclusion of Suspect as a possible contributor to the DNA profile obtained as compared with the alternate explanation of the evidence. Database(s) Used: NIST revised: African American, Asian, Caucasian, Hispanic
L67JVL - 5901	Method(s): Likelihood Ratio Stats Analysis: 3e: Assuming Female Victim (K1), the evidence profile is approximately 272 trillion times more likely if it originated from Female Victim (K1), Male Suspect (K2), and an unknown individual than if it originated from Female Victim (K1) and two unknown individuals. This analysis provides very strong support for inclusion. 3sp: Female Victim (K1) is excluded based on the LR of zero. Suspect (K2) is excluded based on the LR of zero. Database(s) Used: Allele frequencies from the 2015 FBI Expanded Population Database were used for STR statistical calculations.
L8XEJE - 5901	Method(s): Likelihood Ratio Stats Analysis: Sperm Fraction: Item 1 - Assumed contributor, Item 2 - LR = 3.281E2 (320 support for contribution). Epithelial Fraction: Item 1 - Assumed contributor, Item 2 - LR = 4.4071E19 (>100 billion support for contribution). Database(s) Used: [Participant did not return a database used.]

TABLE 7	
WebCode- Test	Item 3 Methods & Results
LBAKVF - 5902	Method(s): Likelihood Ratio Stats Analysis: SUSPECT (Q02-1) cannot be excluded as a contributor to Mixture 1 from the underwear (Q03-1E) from the VICTIM. The STR DNA results are estimated to be greater than a trillion times more likely if they originate from VICTIM, SUSPECT and one unknown person than they originate from VICTIM and two unknown people, unrelated to SUSPECT. Database(s) Used: i. NIST Asian, ii. NIST African American, iii. NIST Caucasian, iv. Srivastava et al. (2019) South Asian, v.[Location Identifying Database]
LG3PNJ - 5901	Method(s): Likelihood Ratio Stats Analysis: Known contributors under Hp: victim and suspect. Known contributors under Hd: victim LR (total): 2,7E13. Database(s) Used: [Location Identifying Database]
LHUN3H - 5902	 Method(s): Likelihood Ratio Stats Analysis: A DNA match was identified between the data obtained from the first fraction from the stain from victim's underwear (Item 3) and Suspect. A match between the data obtained from the first fraction from the stain from victim's underwear (Item 3) and Suspect is approximately 9 sextillion times more probable than a coincidental match to an unrelated person in the population. A DNA match was identified between the data obtained from the first fraction from the stain from victim's underwear (Item 3) and Victim. A match between the data obtained from the first fraction from the stain from victim's underwear (Item 3) and Victim is approximately 100 quadrillion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the stain from victim's underwear (Item 3) and Victim is approximately 100 quadrillion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the second fraction from the stain from victim's underwear (Item 3) and Suspect is excluded from the stain from victim's underwear (Item 3) and Suspect is excluded from the data obtained from the second fraction from the stain from victim's underwear (Item 3). A DNA match between the data obtained from the second fraction from the stain from victim's underwear (Item 3) and Victim is approximately 300 octillion times less probable than a coincidental match to an unrelated person in the population. Victim is excluded from the data obtained from the second fraction from the stain from victim's underwear (Item 3). Database(s) Used: The National Institute of Standards and Technology's 1036 Revised U.S. combined population database was used to generate DNA match statistics. The combined population database contains African American, Caucasian, Hispanic and Asian populations. Identical Shings will have identical DNA data and DNA match statistics.

TABLE 7	
WebCode-	Item 3 Methods & Results
Test	
LUPUWF - 5901	Method(s) : Likelihood Ratio Stats Analysis: For the epithelial fraction: The LR value regarding the possible involvement of the female victim was calculated to be 2.52×10^{-15} to 1, which means it is about 2.52×10^{-15} to 1 times more likely that the observed DNA profile being a mixture originating from the female victim and two unknown individuals than if it originating from three unrelated individuals selected at random from the local [Population]. The LR value regarding the possible involvement of the male suspect was calculated to be 2.14×10^{-3} 0 to 1, which means it is about 2.14×10^{-3} 0 to 1 times more likely that the observed DNA profile being a mixture originating from the male suspect and two unknown individuals than if it originating from three unrelated individuals selected at random from the local [Population]. For the sperm fraction: The LR value regarding the possible involvement of the female victim was calculated to be 1.36×10^{-3} to 1, which means it is about 1.36×10^{-3} to 1 times more likely that the observed DNA profile being a mixture originating from the female victim and two unknown individuals than if it originating from three unrelated individuals selected at random from the local [Population]. The LR value regarding the possible involvement of the male suspect was calculated to be 1.22×10^{-6} to 1, which means it is about 1.22×10^{-6} to 1 times more likely that the observed DNA profile being a mixture originating from the male suspect and two unknown individuals than if it originating from three unrelated individuals selected at random from the local [Population]. The LR value regarding the possible involvement of the male suspect was calculated to be 1.22×10^{-6} to 1, which means it is about 1.22×10^{-6} to 1 times more likely that the observed DNA profile being a mixture originating from the male suspect and two unknown individuals than if it originating from three unrelated individuals selected at random fr
LYKQJL - 5901	 Method(s): Likelihood Ratio Stats Analysis: Non-sperm Fraction: Assuming Female Victim (K1), the evidence DNA profile is approximately 26 trillion times more likely if it originated from Female Victim (K1), Male Suspect (K2), and an unknown individual (Hypothesis 1) than if it originated from Female Victim (K1) and two unknown individuals (Hypothesis 2). This analysis provides very strong support for the proposition that Male Suspect (K2) is a contributor to the DNA obtained from this sub-item. Sperm Fraction: The evidence DNA profile is approximately 0 times more likely if it originated from Male Suspect (K2) (Hypothesis 1) than if it originated from an unknown individual (Hypothesis 2). This analysis provides an exclusion for the proposition that Male Suspect (K2) (Hypothesis 1) than if it originated from an unknown individual (Hypothesis 2). This analysis provides an exclusion for the proposition that Male Suspect (K2) is a contributor to the DNA obtained from this sub-item. Database(s) Used: 2015 FBI Expanded Population Database
MW99ZF - 5902	 Method(s): Likelihood Ratio Stats Analysis: A DNA match was identified between the data obtained from the first fraction from the questioned stain from the victim's underwear (Item 3) and suspect. A match between the data obtained from the first fraction from the questioned stain from the victim's underwear (Item 3) and suspect is approximately 4 sextillion times more probable than a coincidental match to an unrelated person in the population. A DNA match was identified between the data obtained from the first fraction from the questioned stain from the victim's underwear (Item 3) and victim. A match between the data obtained from the first fraction from the questioned stain from the victim's underwear (Item 3) and victim. A match between the data obtained from the first fraction from the guestioned stain from the victim's underwear (Item 3) and victim is approximately 6 trillion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the second fraction from the questioned stain from the victim's underwear (Item 3) and suspect is approximately 6 undecillion times less probable than a coincidental match to an unrelated person in the population. Suspect is excluded from the data obtained from the second fraction from the questioned stain from the victim's underwear (Item 3) and victim is approximately 400 nonillion times less probable than a coincidental match to an unrelated person in the population. Victim is excluded from the data obtained from the second fraction from the questioned stain from the victim's underwear (Item 3). Database(s) Used: National Institute of Standards and Technology's 1036 Revised U.S. combined population database.

WebCode-	Item 3 Methods & Results
Test	
NWNRKD - 5901	 Method(s): Likelihood Ratio Stats Analysis: For the epithelial (e) fraction: The calculated LR value for the possible involvement of the victim in this mixture was 1.27E+7 to 1. This means it is about 1.27E+7 times more likely that the observed DNA profile being a mixture originating from the victim and two unknown persons than if it originating from three unrelated persons selected at random from the local [Population]. The calculated LR value for the possible involvement of the suspect in this mixture was 1.97E+15 to 1. This means it is about 1.97E+15 to 1. This means it is about 1.97E+15 times more likely that the observed DNA profile being a mixture originating from the suspect and two unknown persons than if it originating from the suspect and two unknown persons than if it originating from the suspect and two unknown persons than if it originating from the suspect and two unknown persons than if it originating from the suspect and two unknown persons than if it originating from the suspect and two unknown persons than if it originating from the suspect and two unknown persons than if it originating from the suspect and two unknown persons than if it originating from three unrelated persons selected at random from the local [Population]. Database(s) Used: [Location Identifying Database]
P3QH3A - 5901	Method(s): Likelihood Ratio Stats Analysis: Item 3e: Item 1 - conditioned on. Item 2 - support for contribution greater than 100 billion (8.29E17). Unresolved unknown contributor. Item 3sp: Item 1 - conditioned on. Item 2 - support for contribution greater than 100 billion (2.21E15). Resolved unknown male profile. Database(s) Used: [Location Identifying Database]
P8KEPF - 5902	Method(s): Likelihood Ratio Stats Analysis: H1:The evidence originated from Victim, suspect, and one unknown, unrelated individual. H2: The evidence originated from Victim and two unknown, unrelated individuals. The DNA profile obtained from the non-sperm cell fraction of Item 3 is approximately 15 quintillion times more likely if it originated from Victim, Suspect and one unknown, unrelated individual than if it originated from the Victim and two unknown, unrelated individuals. There is evidentiary support for the inclusion of Suspect as a possible contributor to the DNA profile obtained as compared with the alternate explanation of the evidence.
	Database(s) Used: NIST revised: African American, Asian, Caucasian, Hispanic
P8TP7Q - 5901	Method(s): Likelihood Ratio Stats Analysis: Item 3.1 (Portion of "Questioned stain from victim's underwear" - Fraction 2): The DNA profile from this item was interpreted as a mixture of three individuals, with at least one male contributor and with the female victim as an assumed contributor. The DNA results are approximately 2.81 quintillion times more likely if they originated from the female victim, the male suspect, and an unknown, unrelated individual than if they originated from the female victim and two unknown, unrelated individuals. Based on the likelihood ratio, this provides very strong support that the male suspect is a contributor to the DNA from this item. Item 3.1 (Portion of "Questioned stain from victim's underwear" - Fraction 1): The male DNA profile from this item was interpreted as a single-source profile. The female victim and the male suspect are excluded as possible contributors to the DNA from this item. Database(s) Used: NIST 1036 July 2017
PD88ZF - 5902	 Method(s): Likelihood Ratio Stats Analysis: The DNA profile obtained from the sperm fraction was interpreted as a mixture of two individuals. The DNA profile is 27 times more likely if it originated from the Suspect (2) and one unknown individual than if it had originated from two unknown, unrelated individuals. Statistical analysis provides limited support for the inclusion of the Suspect (2). The Victim (1) is excluded as a contributor to the DNA profile. The DNA profile obtained from the epithelial fraction was interpreted as a mixture of three individuals. The Victim (1) was assumed as a contributor. The DNA profile is at least 1 trillion times more likely if it originated from the Victim (1), the Suspect (2), and one unknown individual than if it had originated from the Victim (1) and two unknown, unrelated individuals. Statistical analysis provides very strong support for the inclusion of the Suspect (2). Database(s) Used: The FBI Extended CODIS core allele frequencies - Caucasian, African American, and Southwest Hispanics

WebCode-	Item 3 Methods & Results
Test	
Q99BQE - 5901	Method(s): Likelihood Ratio Stats Analysis: [Participant did not return statistical analysis.] Database(s) Used: STRMIX 2.8.0 used Knowing the underwear belongs to the victim, the suspect's contribution in the mixture is calculated according to the following hypothesis : Hp : the mixture belongs to the victim, the suspect and 1 unknown person. Hd : the mixture belongs to the victim and 2 unknown people. LR = 7.0273E24.
R3MFPB - 5901	 Method(s): Likelihood Ratio Stats Analysis: For the epithelial fraction: The LR value for the involvement of female victim is 6.75E+20 to 1, which means it is about 6.75E+20 times more likely that the observed DNA profile being a mixture originating from the female victim and two unknown individuals than if it originating from three unrelated individuals selected at random from the local [Population]. The LR value for the involvement of male suspect is 2.02E+31 to 1, which means it is about 2.02E+31 times more likely that the observed DNA profile being a mixture originating from the reale suspect is 2.02E+31 to 1, which means it is about 2.02E+31 times more likely that the observed DNA profile being a mixture originating from the male suspect and two unknown individuals than if it originating from three unrelated individuals selected at random from the local [Population]. For the sperm fraction: The LR value for the involvement of female victim is 5.67E-1 to 1 which is uninformative. Hence, no conclusion could be made as to whether or not the female victim is a possible source of the DNA profile. The LR value for the involvement of male suspect is 7.22E+12 to 1, which means it is about 7.22E+12 times more likely that the observed DNA profile being a mixture originating from the male suspect and two unknown individuals than if it originating from three unrelated individuals selected at random from the local [Population]. Database(s) Used: [Location Identifying Database]
R9RDYE - 5901	Method(s): Likelihood Ratio Stats Analysis: For 3e: Assuming Female Victim (K1), the evidence profile is approximately 229 Trillion times more likely if it originated from Female Victim (K1), Male Suspect (K2) and an unknown individual than if it originated from Female Victim (K1) and two unknown individuals. This provides very strong support for the inclusion of Male Suspect (K2). For 3SP: The evidence DNA profile is approximately 0 times more likely if it originated from Female Victim (K1) than if it originated from an unknown individual. Female Victim (K1) is excluded. For 3SP: The evidence DNA profile is approximately 0 times more likely if it originated from Male Suspect (K2) than if it originated from an unknown individual. Male Suspect (K2) is excluded. Database(s) Used: 2015 FBI Expanded Population Database.
RCPXZD - 5901	Method(s): Likelihood Ratio Stats Analysis: LR was calculated for the epithelial fraction of Item 3, assuming the mixture is comprised of three contributors. We assumed the presence of the victim on her own underwear is not in question. The DNA profile is approximately 5*10^17 (500 billiard) times more likely if it originated from the Victim, the Suspect and an unknown individual than if it originated from the Victim and two unknown individuals. Database(s) Used: [Location Identifying Database]
TMDEV4 - 5902	Method(s): Likelihood Ratio Stats Analysis: FBI_EXTENDED_CAUC Database(s) Used: [Participant did not return a database used.]

TABLE 7	
WebCode-	Item 3 Methods & Results
Test	
TZBZVA - 5901	Method(s): Likelihood Ratio Stats Analysis: For the epithelial(e) fraction: The LR value calculated for the possible involvement of the female victim was 2.18E7 to 1, which means it is about 2.18E7 times more likely that the observed DNA profile being a mixture originating from the female victim and two unknown individuals than if it originating from three unrelated individuals selected at random from the local [Population]. The LR value calculated for the possible involvement of the male suspect was 4.93E18 to 1, which means it is about 4.93E18 times more likely that the observed DNA profile being a mixture originating from the male suspect and two unknown individuals than if it originating from three unrelated individuals selected at random from the local [Population]. Database(s) Used: [Location Identifying Database]
UNVR99 - 5902	 Method(s): Likelihood Ratio Stats Analysis: A DNA match was identified between the data obtained from the first fraction from the questioned stain from victim's underwear (Item 3) and victim. A match between the data obtained from the first fraction from the questioned stain from victim's underwear (Item 3) and victim is approximately 600 million times more probable than a coincidental match to an unrelated person in the population. A DNA match was identified between the data obtained from the first fraction from the questioned stain from victim's underwear (Item 3) and suspect. A match between the data obtained from the first fraction from the first fraction from the questioned stain from victim's underwear (Item 3) and suspect is approximately 300 quintillion times more probable than a coincidental match to an unrelated person in the population. Due to the complexity of the mixture, no conclusive determination can be made as to the evidence genotypes from the data obtained from the second fraction from the questioned stain from victim's underwear (Item 3). Database(s) Used: The National Institute of Standards and Technology's 1036 Revised U.S. combined population database.
VFHZHA - 5902	Method(s): Likelihood Ratio Stats Analysis: Hypothesis 1 (H1): The evidence originated from Victim, Suspect and one unknown, unrelated individual. Hypothesis 2 (H2): The evidence originated from Victim and two unknown, unrelated individuals. The DNA profile obtained from the non-sperm cell fraction of the underwear is approximately 66 quintillion times more likely if it originated from Victim, Suspect, and one unknown, unrelated individual than if it originated from Victim and two unknown, unrelated individuals. Database(s) Used: NIST revised: African American, Asian, Caucasian, Hispanic.
VK9834 - 5902	Method(s): Likelihood Ratio Stats Analysis: Semen fraction: H1: The DNA on Item 3 (Semen fraction) came from the suspect. H2: The DNA on Item 3 (Semen fraction) came from an unknown, unrelated person. LR= 0. H1: The DNA on Item 3 (Semen fraction) came from the victim. H2: The DNA on Item 3 (Semen fraction) came from an unknown, unrelated person. LR= 0. Epithelial fraction: H1: The DNA on Item 3 (Epithelial fraction) came from the suspect and two unknown, unrelated persons. H2: The DNA on Item 3 (Epithelial fraction) came from three unknown, unrelated persons. LR= 2.0093E17. H1: The DNA on Item 3 (Epithelial fraction) came from three unknown, unrelated persons. H2: The DNA on Item 3 (Epithelial fraction) came from three unknown, unrelated persons. LR= 2.2010E15. H1: The DNA on Item 3 (Epithelial fraction) came from three unknown, unrelated persons. LR= 2.2010E15. H1: The DNA on Item 3 (Epithelial fraction) came from three unknown, unrelated persons. LR= 3.5888E12 DNA on Item 3 (Epithelial fraction) came from three unknown, unrelated persons. H2: The DNA on Item 3 (Epithelial fraction) came from three unknown, unrelated persons. LR= 3.5888E12 Database(s) Used: [Location Identifying Database]

TABLE 7		
WebCode- Test	Item 3 Methods & Results	
W2RLKZ - 5901	 Method(s): Likelihood Ratio Stats Analysis: Item 3.1 (Portion of "Questioned stain from victim's underwear" - Fraction 1): The male DNA profile from this item was interpreted as a single-source profile. The Female Victim and The Male Suspect are excluded as possible contributors to this DNA profile. Item 3.1 (Portion of "Questioned stain from victim's underwear" - Fraction 2): The DNA profile from this item was interpreted as a mixture of three individuals with at least one male contributor and with The Female Victim as an assumed contributor. The DNA results are approximately 3.97 sextillion times more likely if they originated from The Female Victim, The Male Suspect and an unknown, unrelated individual than if they originated from The Female Victim and two unknown, unrelated individuals. Based on the likelihood ratio, this provides very strong support that The Male Suspect is a contributor to the DNA from this item. Database(s) Used: NIST 1036 July 2017 	
WRGKYZ - 5901	 Method(s): Likelihood Ratio Stats Analysis: Item 3.1 (Portion of "Questioned stain from victim's underwear" - Fraction 2): The DNA profile from this item was interpreted as a mixture of three individuals, with at least one male contributor and with Female Victim as an assumed contributor. The DNA results are approximately 42.4 quintillion times more likely if they originated from Male Suspect, Female Victim and an unknown, unrelated individual than if they originated from Female Victim and two unknown, unrelated individuals. Based on the likelihood ratio, this provides very strong support that Male Suspect is a contributor to the DNA from this item. Item 3.1 (Portion of "Questioned stain from victim's underwear" - Fraction 1): The male DNA profile from this item was interpreted as a single-source profile. Female Victim and Male Suspect are excluded as possible contributors to the DNA from this item. Database(s) Used: NIST 1036 July 2017 	
WU8M28 - 5901	 Method(s): Likelihood Ratio Stats Analysis: The DNA profile obtained from the sperm fraction was interpreted as originating from a single unknown male individual A. The DNA profile obtained from the epithelial fraction was interpreted as a mixture of three individuals. Assuming female victim (1) and the corresponding sperm fraction as contributors, the mixture contains one unknown individual. The DNA profile is at least 1 trillion times more likely if it originated from female victim (1), male suspect (2), and one unknown individual than if it had originated from female victim (1) and two unknown, unrelated individuals. Statistical analysis provides very strong support for the inclusion of male suspect (2). Database(s) Used: FBI Extended CODIS Core Allele Frequencies- Caucasian, African American, and Southwest Hispanics 	
X28843 - 5901	Method(s): Likelihood Ratio Stats Analysis: Spermatozoa Fraction: Item 1 - approximately 830 million supporting non-contribution; Item 2 - approximately 3500 supporting non-contribution. Epithelial Fraction: Item 1 - assumed contributor; Item 2 - greater than 100 billion supporting contribution. Database(s) Used: [Location Identifying Database]	
YGJ6L9 - 5901	Method(s): Likelihood Ratio Stats Analysis: For 3e - Assuming the victim (item 1), the evidence profile is approximately 247 trillion times more likely if it originated from the victim (item 1), suspect (item 2), and one unknown individual than if it originated from the victim (item 1) and two unknown individuals. This provides very strong support for inclusion of suspect (item 2). For 3s - Victim (item 1) is excluded with a likelihood ratio of 0. Suspect (item 2) is excluded with a likelihood ratio of 0. Database(s) Used: 2015 FBI Expanded Population Database	

WebCode- Test	Item 3 Methods & Results
YJMUQ3 - 5901	Method(s): Likelihood Ratio Stats Analysis: Assuming three contributors, including at least two males, the DNA typing results were interpreted using probabilistic genotyping software. The Victim (Item 1) is assumed to be a contributor to this mixture. The DNA typing results are 29.6 nonillion times more likely to be observed if they originated from the Victim (Item 1), the Suspect (Item 2), and one unknown, unrelated individual, rather than if they originated from the Victim (Item 1) and two unknown, unrelated individuals. This analysis provides very strong support for the proposition that includes the Suspect (Item 2) as one of the contributors to this mixture. Database(s) Used: NIST 1036 (2017)
Z8LUQ4 - 5902	 Method(s): Likelihood Ratio Stats Analysis: A DNA match was identified between the data obtained from the first fraction from the cutting from the tan fabric (item 3) and the subject. A match between the data obtained from the first fraction from the cutting from the tan fabric (item 3) and the subject is approximately 600 septillion times more probable than a coincidental match to an unrelated person in the population. A DNA match was identified between the data obtained from first fraction from the cutting from the tan fabric (item 3) and the victim. A match between the data obtained from first fraction from the cutting from the tan fabric (item 3) and the victim is approximately 600 quadrillion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from first fraction from the cutting from the tan fabric (item 3) and the victim is approximately 600 quadrillion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the second fraction from the cutting from the tan fabric (item 3) and the victim is approximately 400 nonillion times less probable than a coincidental match to an unrelated person in the population. The victim is excluded from the data obtained from the second fraction from the cutting from the tan fabric (item 3). A DNA match between the data obtained from the second fraction from the cutting from the tan fabric (item 3) and the subject is approximately 600 undecillion times less probable than a coincidental match to an unrelated person in the population. The subject is excluded from the data obtained from the second fraction from the cutting from the tan fabric (item 3). A DNA match between the data obtained from the second fraction from the cutting from the tan fabric (item 3). Database(s) Used: The National Institute of Standards and Technology's 1036 Revised U.S. combined population database

Statistical Analysis for Item 4

TABLE 8		
WebCode-	Item 4 Methods & Results	
Test		
2QGMC2 - 5902	 Method(s): Likelihood Ratio Stats Analysis: A DNA match was identified between the data obtained from the questioned stain from victim's pants (item 4) and the male suspect. A match between the data obtained from the questioned stain from victim's pants (item 4) and the male suspect is approximately one septillion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the questioned stain from victim's pants (item 4) and the questioned stain from victim's pants (item 4) and the victim is approximately two hundred sextillion times less probable than a coincidental match to an unrelated person in the population. The victim is excluded from the data obtained from the questioned stain from victim's pants (item 4). Database(s) Used: The National Institute of Standards and Technology's 1036 Revised U.S. combined population database 	
34G3B3 - 5902	Method(s): Likelihood Ratio Stats Analysis: Assuming three contributors, Female Victim is excluded (LR=0) as being a contributor to the mixture detected for this item. Assuming three contributors, the evidence profile is 1.3E+24 times more likely to be observed if Male Suspect and two unknown individuals are the contributors than if three unknown individuals are the contributors. Database(s) Used: NIST Databases - Forensic Sci Int.: Genetics 31 (2017) e36-e40	
37J2CU - 5901	Method(s): Likelihood Ratio Stats Analysis: [Participant did not return statistical analysis.] Database(s) Used: [Location Identifying Database]	
3MQZUY - 5902	Method(s): Likelihood Ratio Stats Analysis: 4. The SUSPECT (item 2-1) cannot be excluded as a contributor to Mixture 2 (see Testing Summary) from the VICTIMS pants (item 4-1) The STR DNA results are estimated to be greater than one trillion times more likely if they originate from the SUSPECT and two unknown people than if they originate from three unknown people unrelated to the SUSPECT. Database(s) Used: i. NIST Asian, ii. NIST African American, iii. NIST Caucasian, iv. Srivastava et al. (2019) South Asian, v. [Location Identifying Database]	
3WHFJ3 - 5902	 Method(s): Likelihood Ratio Stats Analysis: H1: The evidence originated from the Known Male and two unknown, unrelated individuals. H2: The evidence originated from three unknown, unrelated individuals. The DNA profile obtained from Item 4 is approximately 520 quadrillion times more likely if it originated from the Known Male and two unknown, unrelated individuals than if it originated from three unknown, unrelated individuals. There is evidentiary support for the inclusion of the Known Male as a possible contributor to the DNA profile obtained as compared with the alternate explanation of the evidence. Database(s) Used: NIST revised: African American, Asian, Caucasian, Hispanic 	
44PDRD - 5901	Method(s): Likelihood Ratio Stats Analysis: Item 4.1 (Portion of "Questioned stain from victim's pants"): The DNA profile from this item was interpreted as a mixture of three individuals, with at least two male contributors. The DNA results are approximately 106 sextillion times more likely if they originated from MALE SUSPECT and two unknown, unrelated individuals than if they originated from three unknown, unrelated individuals. Based on the likelihood ratio, this provides very strong support that MALE SUSPECT is a contributor to the DNA from this item. FEMALE VICTIM is excluded as a possible contributor to the DNA from this item. Database(s) Used: NIST 1036 July 2017	

TABLE 8		
WebCode- Test	Item 4 Methods & Results	
4BEUJW - 5902	Method(s): Likelihood Ratio Stats Analysis: Victim (Item 1): Excluded as being a contributor to the DNA in this sample. Suspect (Item 2): The DNA evidence is estimated to be 100 billion times more likely to occur if the DNA originated from the suspect and and two unknown, unrelated people selected randomly from the [Population], than if it originated from three unknown and unrelated people selected randomly from the [Population]. Database(s) Used: [Location Identifying Database]	
4FBJY4 - 5902	Method(s): Likelihood Ratio Stats Analysis: Item 1: no statistical calculation performed due to number of missmatches. Item 2: 4.524e+15. Unknown male from sperm fraction: no statistical calculation performed due to number of missmatches. Database(s) Used: [Location Identifying Database]	
4UQG64 - 5901	Method(s): Likelihood Ratio Stats Analysis: The evidence DNA profile 4A-2PT is approximately 0 times more likely if it originated from Female Victim (K1) and two unknown individuals than if it originated from three unknown individuals. This provides support for exclusion of Female Victim (K1). The evidence DNA profile 4A-2PT is approximately 572 trillion times more likely if it originated from Male Suspect (K2) and two unknown individuals than if it originated from three unknown individuals. This provides very strong support for inclusion of Male Suspect (K2). Database(s) Used: 2015 FBI Expanded Population Database	
4XQRZ4 - 5901	Method(s): Likelihood Ratio Stats Analysis: Female Victim (K1) is excluded based on the LR of zero. The evidence profile is approximately 25 trillion times more likely if it originated from Male Suspect (K2) and two unknown individuals than if it originated from three unknown individuals. This analysis provides very strong support that Male Suspect (K2) is a contributor to the DNA obtained from this sub-item Database(s) Used: 2015 FBI Expanded Population Database	
62K9Q4 - 5901	Method(s): Likelihood Ratio Stats Analysis: For 4A-2: The evidence profile is approximately 0 times more likely if it originated from Female Victim (K1) and two unknown individuals than if it originated from three unknown individuals. Female Victim (K1) is excluded based on the LR of zero. The evidence profile is approximately 3 quadrillion times more likely if it originated from Male Suspect (K2) and two unknown individuals than if it originated from three unknown individuals. This analysis provides very strong support for the proposition that Male Suspect (K2) is a contributor to the DNA obtained from this sub-item. Database(s) Used: 2015 FBI Expanded Population	
63VBC2 - 5901	Method(s): Likelihood Ratio Stats Analysis: The evidence DNA profile is approximately 0 times more likely if it originated from Female Victim (K1) and two unknown individuals than if it originated from three unknown individuals. The evidence DNA profile is approximately 12 trillion times more likely if it originated from Male Suspect (K2) and two unknown individuals than if it originated from three unknown individuals. This analysis provides very strong support for the proposition that Male Suspect (K2) is included as a contributor to this DNA mixture. Database(s) Used: 2015 FBI Expanded Population Database	

TABLE 8	
WebCode-	Item 4 Methods & Results
Test	
6PTH6Y - 5902	 Method(s): Likelihood Ratio Stats Analysis: The victim's pants from item 4 contained a mixture of DNA from at least three individuals. A DNA match was identified between the data obtained from the victim's pants from item 4 and the suspect. A match between the data obtained from the victim's pants from item 4 and the suspect is approximately one septillion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the victim's pants from the victim's pants from item 4 and the victim is approximately two sextillion times less probable than a coincidental match to an unrelated person in the population. The victim is excluded from the data obtained from the victim's pants from item 4. Database(s) Used: The National Institute of Standards and Technology's 1036 Revised U.S. combined population database was used to generate DNA match statistics. The combined populations and the statistical DNA data and DNA match statistics.
77A6RY - 5901	Method(s) : Likelihood Ratio Stats Analysis: 2.9365E26 Database(s) Used: The FBI's extended database for African Americans was used.
786NUZ - 5901	Method(s): Likelihood Ratio Stats Analysis: The DNA profile obtained from item # 4 is a mixture of at least 3 individuals, following set of hypotheses were evaluated: A) Hp; item No. 1 + 2 unknown, Hd; 3 unknown, LR (MLE) = 1.698 e -5. B) Hp; item No. 2 + 2 unknown, Hd; 3 unknown, LR (MLE) = 3.627 e +17 Database(s) Used: NIST Caucasian database 2017 with theta co-ancestry coefficient = 0.03
7H4JRV - 5902	 Method(s): Likelihood Ratio Stats Analysis: A DNA match was identified between the data obtained from the questioned stain from victim's pants (item 4) and the subject. A match between the data obtained from the questioned stain from victim's pants (item 4) and subject is approximately one hundred quadrillion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the questioned stain from victim's pants (item 4) and the victim is approximately two hundred sextillion times less probable than a coincidental match to an unrelated person in the population. The victim is excluded from the data obtained from the questioned stain from victim's pants (item 4). Database(s) Used: The National Institute of Standards and Technology's 1036 Revised U.S. combined population database
8DDAHX - 5901	 Method(s): Likelihood Ratio Stats Analysis: The DNA profile of the epithelial fraction was interpreted as a mixture of three individuals. The DNA profile is at least 1 trillion times more likely if it originated from Known Male Suspect (2) and two unknown individuals than if it had originated from three unknown, unrelated individuals. Statistical analysis provides very strong support for the inclusion of Known Male Suspect (2). Known Female Victim (1) is excluded as a contributor to the DNA profile. Due to an insufficient amount of male DNA detected during quantitation, autosomal Short Tandem Repeat (STR) analysis was not performed on the sperm fraction. Database(s) Used: FBI Extended CODIS core allele frequencies - Caucasian, African American, and Southwest Hispanic

TABLE 8		
WebCode-	Item 4 Methods & Results	
Test		
8Y8DYV - 5902	 Method(s): Likelihood Ratio Stats Analysis: A DNA match was identified between the data obtained from the "questioned stainpants" (item #4) and the male suspect (item #2). A match between the data obtained from the "questioned stainpants" (item #4) and the male suspect (item #2) is approximately 1 septillion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the "questioned stainpants" (item #1) is approximately 500 quintillion times less probable than a coincidental match to an unrelated person in the population. The female victim (item #1) is approximately 500 quintillion times less probable than a coincidental match to an unrelated person in the population. The female victim (item #1) is excluded from the data obtained stainpants" (item #4). Database(s) Used: The National Institute of Standards and Technology's 1036 Revised U.S. combined population database was used to generate DNA match statistics. The combined populations. Identical siblings will have identical DNA data and DNA match statistics. 	
8YUL9Z - 5901	 Method(s): Likelihood Ratio Stats Analysis: A likelihood ratio (LR) analysis was performed using the LRmix Studio software, using allele frequency data specific to the [Population]- based on own laboratory reference population database consisting of 5,000 genotypes from [Country]. The LR values for mixed DNA profiles were calculated with consideration for both drop-in and drop-out phenomena. The analysis assumed a drop-in probability of 0.05 and applied a minimum allele frequency threshold of 0.001. A subpopulation correction factor of 0.01 was also incorporated into the calculations. The resulting LR values were interpreted in terms of their probative value. The evaluated hypotheses included: Prosecution Hypothesis (Hp): The DNA mixture includes the genetic profiles of item2, and two unknown, untested individual from the population. Alternative Hypothesis (Hd): The DNA mixture includes the genetic evidence in the context of the case. Likelihood ratio on this sample is: 4.6942E10. Y-STR: Given the Haplotypes above and that there is 1 additional unknown contributor, the likelihood of observing the given trace under the hypothesis of the donorship of the suspect is approx. 5,786 times more likely than observing the given trace under the hypothesis of the non-donorship. This calculation is based on 298,405 haplotypes (Y17) in the worldwide database YHRD R69. Database(s) Used: STR laboratory population base of [Country] 	
9CLYK9 - 5901	Method(s): Likelihood Ratio Stats Analysis: Item 4.1 (Portion of "Questioned stain from victim's pants"): The DNA profile from this item was interpreted as a mixture of three individuals with at least two male contributors. Based on the likelihood ratio result, the female victim is excluded as a possible contributor to the DNA from this item. The DNA results are approximately 35.1 sextillion times more likely if they originated from the male suspect and two unknown, unrelated individuals than if they originated from three unknown, unrelated individuals. Based on the likelihood ratio, this provides very strong support that the male victim is a contributor to the DNA from this item. Database(s) Used: NIST 1036 July 2017	
9MBR8Q - 5901	Method(s): Likelihood Ratio Stats Analysis: Item 1 (complainant) - Excluded. Item 2 (known suspect) - >100 billion in favour of H1 (favouring contribution). Database(s) Used: [Location Identifying Database]	
9WLTTR - 5902	Method(s): Likelihood Ratio Stats Analysis: The DNA evidence is greater than 100 billion times more likely if the suspect is a contributor. The victim is excluded as a contributor. Database(s) Used: [Location Identifying Database]	

WebCode-	Item 4 Methods & Results
Test	
B7W38V - 5901	Method(s): Likelihood Ratio Stats Analysis: 1.3551E21 Database(s) Used: FBI Extended African American
BDR7Z7 - 5901	Method(s): Likelihood Ratio Stats Analysis: Explanation 1: The DNA profile obtained originated from item 2 (1D), and two unknown, unrelated individuals. Explanation 2: The DNA profile obtained originated from three unknown, unrelated individuals. The DNA profile obtained is approximately 1E33 times more likely (very strong support) if it originated from the item 2 (1D) and two unknown, unrelated individuals than if the DNA profile obtained originated from three unknown, unrelated individuals. Database(s) Used: Nist1036 (revised 2017)_Caucasian, Nist1036 (revised 2017)_African American, Nist1036 (revised 2017)_Hispanic
BKHHBU - 5901	Method(s): Likelihood Ratio Stats Analysis: 2.8651e24 Database(s) Used: FBI Extended African American
BNGT8U - 5902	Method(s): Likelihood Ratio Stats Analysis: The LR calculations exclude the victim as being a contributor to the DNA mixture detected for item 4. Assuming 3 contributors for item 4, the evidence profile is 9.9 x 10 ^ 26 times more likely to be observed if the suspect and two unknowns are the contributors than if three unknowns are the contributors. Database(s) Used: NIST Databases - Forensic Sci. Int.: Genetics 31 (2017) e36–e40
CDTG9M - 5901	Method(s): Likelihood Ratio, STRmix Stats Analysis: Propositions for contributor 1: The propositions considered for ITEM 4 were: P1: The SUSPECT is one of three contributors to the mixed DNA profile obtained. P2: The SUSPECT is NOT a contributor to the mixed DNA profile obtained. Three unknown individuals, unrelated to the SUSPECT, are the contributors. For this calculation, it was assumed that three individuals have contributed to the DNA profile obtained. Likelihood Ratio: The evidence is at least 100 BILLION (5.5762E22) times more likely if the SUSPECT is one of three contributors to the mixed DNA profile than if the profile originated from three unknown individuals, unrelated to the SUSPECT, selected at random from the [Population]. Database(s) Used: [Location Identifying Database]
CKECAQ - 5902	 Method(s): Likelihood Ratio Stats Analysis: A DNA match was identified between the data obtained from the questioned stain from victim's pants (Item 4) and the Subject. A match between the data obtained from the questioned stain from victim's pants (Item 4) and the Subject is approximately 900 sextillion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the questioned stain from victim's pants (Item 4) and the questioned stain from victim's pants (Item 4) and the Victim is approximately 90 septillion times less probable than a coincidental match to an unrelated person in the population. The Victim is excluded from the data obtained from the questioned stain from the questioned stain from victim's pants (Item 4). Database(s) Used: All evidence genotypes were compared with all reference genotypes to compute likelihood ratio (LR) DNA match statistics. The National Institute of Standards and Technology's 1036 Revised U.S. combined population database was used to generate DNA match statistics. The combined population database contains African American, Caucasian, Hispanic and Asian populations. Identical siblings will have identical DNA data and DNA match statistics.

TABLE 8	
WebCode- Test	Item 4 Methods & Results
CRBAWU - 5901	Method(s): Likelihood Ratio Stats Analysis: The DNA evidence is at least one billion times more likely if the mixture of DNA originated from the Suspect and two unknown unrelated individuals rather than if it originated from three unknown, unrelated individuals Database(s) Used: [Location Identifying Database]
CRVQCW - 5901	Method(s): Likelihood Ratio Stats Analysis: The evidence DNA profile is 95 Trillion times more likely if it originated from Male Suspect (K2) and two unknown individuals than if it originated from three unknown individuals. This provides very strong support for inclusion. Female Victim (K1) is excluded based on the LR of zero. Database(s) Used: 2015 FBI Expanded Population Database
CXYRCQ - 5902	 Method(s): Likelihood Ratio Stats Analysis: A DNA match was identified between the data obtained from the questioned stain from victim's pants (Item 4) and Suspect. A match between the data obtained from the questioned stain from victim's pants (Item 4) and Suspect is approximately twenty septillion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the questioned stain from victim's pants (Item 4) and Suspect is approximately twenty septillion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the questioned stain from victim's pants (Item 4) and Victim is approximately twenty sextillion times less probable than a coincidental match to an unrelated person in the population. Victim is excluded from the data obtained from the questioned stain from victim's pants (Item 4). Database(s) Used: The National Institute of Standards and Technology's 1036 Revised U.S. combined population database was used to generate DNA match statistics. The combined population database contains African American, Caucasian, Hispanic and Asian populations. Identical siblings will have identical DNA data and DNA match statistics.
CY6DP6 - 5901	Method(s): Likelihood Ratio Stats Analysis: The DNA profile from this item was interpreted as a mixture of three individuals with at least two male contributors. The DNA results are approximately 2.98 quadrillion times more likely if they originated from the male suspect and two unknown, unrelated individuals than if they originated from three unknown, unrelated individuals. Based on the likelihood ratio, this provides very strong support that the male suspect is a contributor to the DNA from this item. Based on the likelihood ratio result, the female victim is excluded as a possible contributor to the DNA from this item. Database(s) Used: NIST 1036 July 2017
CZ6EPV - 5901	Method(s): Likelihood Ratio Stats Analysis: Item 4: The evidence DNA profile is approximately 82 Trillion times more likely if it originated from the male suspect (K2) and two unknown individuals than if it originated from three unknown individuals. This analysis provides very strong support for the proposition that the male suspect (K2) is a contributor to this DNA mixture. Female victim (K1) is excluded based on the LR of zero. Database(s) Used: 2015 FBI Expanded Population Database

TABLE 8		
WebCode- Test	Item 4 Methods & Results	
E34ANN - 5902	Method(s): Likelihood Ratio Stats Analysis: A DNA match was identified between the data obtained from the stain from pants (item 4) and Suspect. A match between the data obtained from the stain from pants (item 4) and Suspect is approximately 1 septillion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the stain from pants (item 4) and Victim is approximately 600 sextillion times less probable than a coincidental match to an unrelated person in the population. Victim is excluded from the data obtained from the stain from pants (item 4). Database(s) Used: NIST 1036	
EHXZJK - 5902	Method(s) : Likelihood Ratio Stats Analysis: PP21 Item 1: Excluded. Item 2: Not excluded (LR = 100 billion). YFP Item 2: Unable to be compared as mixture unresolvable. Database(s) Used: [Location Identifying Database]	
ELJJPP - 5902	 Method(s): Likelihood Ratio Stats Analysis: A DNA match was identified between the data obtained from the stain from victim's pants (item 4) and the male suspect (item 2). A match between the data obtained from the stain from victim's pants (item 4) and the male suspect (item 2) is approximately 10 quintillion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the stain from victim's pants (item 1) is approximately 500 quintillion times less probable than a coincidental match to an unrelated person in the population. The female victim (item 1) is excluded from the data obtained from the stain from victim's pants (item 4). Database(s) Used: National Institute of Standards and Technology's 1036 Revised U.S. combined population database 	
EY6YTN - 5902	 Method(s): Likelihood Ratio Stats Analysis: The "victim's pants" (item 4) contained a mixture of DNA from at least three individuals. A DNA match was identified between the data obtained from the "victim's pants" (item 4) and the suspect. A match between the data obtained from the "victim's pants" (item 4) and the suspect is approximately 700 quintillion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the from the from the "victim's pants" (item 4) and the victim is approximately 20 septillion times less probable than a coincidental match to an unrelated person in the population in the population. The victim is excluded from the data obtained from the "victim's pants" (item 4). Database(s) Used: The National Institute of Standards and Technology's 1036 Revised U.S. combined population database was used to generate DNA match statistics. 	
F7H8UP - 5902	Method(s): Likelihood Ratio Stats Analysis: H1: The evidence originated from the suspect and two unknown, unrelated individual. H2: The evidence originated from three unknown, unrelated individuals. The DNA profile obtained from the pants stain is approximately 580 sextillion times more likely if it originated from the suspect and two unknown, unrelated individual than if it originated from three unknown, unrelated individuals. There is evidentiary support for the inclusion of the suspect as a possible contributor to the DNA profile obtained as compared with the alternate explanation of the evidence. Database(s) Used: NIST revised: African American, Asian, Caucasian, Hispanic	

TABLE 8		
WebCode-	Item 4 Methods & Results	
Test		
F8XQDM - 5902	 Method(s): Likelihood Ratio Stats Analysis: A DNA match was identified between the data obtained from the questioned stain from victim's pants (item 4) and Suspect. A match between the data obtained from the questioned stain from victim's pants (item 4) and Suspect is approximately 30 quintillion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the questioned stain from victim is approximately 300 quintillion times less probable than a coincidental match to an unrelated person in the population. Nictim is excluded from the data obtained from the questioned stain from victim's pants (item 4). Database(s) Used: National Institute of Standards and Technology's 1036 Revised U.S. combined population database 	
FPXE6L - 5902	 Method(s): Likelihood Ratio Stats Analysis: The SUSPECT (item Q02-01) cannot be excluded as a contributor to the following: Mixture 2 from a bloodstained area on the pants (item Q04-1) from the VICTIM. The STR DNA results are estimated to be greater than one trillion times more likely if they originate from the SUSPECT and two unknown people than if they originate from three unknown people unrelated to the SUSPECT. Database(s) Used: i. NIST Asian, ii. NIST African American, iii. NIST Caucasian, iv. Srivastava et al. (2019) South Asian, v. [Location Identifying Database] 	
GCH7HK - 5902	 Method(s): Likelihood Ratio Stats Analysis: A DNA match was identified between the data obtained from the questioned stain from victim's pants (item 4) and the suspect. A match between the data obtained from the questioned stain from victim's pants (item 4) and the suspect is approximately 10 octillion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the questioned stain from victim is approximately 10 septillion times less probable than a coincidental match to an unrelated person in the population. The victim is excluded from the data obtained from the questioned stain from victim's pants (item 4). Database(s) Used: National Institute of Standards and Technology's 1036 Revised U.S. combined population 	
H484WJ - 5902	 Method(s): Likelihood Ratio Stats Analysis: A DNA match was identified between the data obtained from the blue fabric (item 4) and the male subject. A match between the data obtained from the blue fabric (item 4) and the male subject is approximately 10 quintillion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the blue fabric (item 4) and the female victim is approximately 90 octillion times less probable than a coincidental match to an unrelated person in the population. The female victim is excluded from the data obtained from the blue fabric (item 4). Database(s) Used: The National Institute of Standards and Technology's 1036 Revised U.S. combined population database was used to generate DNA match statistics. 	
HEQHYL - 5902	Method(s): Likelihood Ratio Stats Analysis: A DNA match was identified between the data obtained from the victim's pants (item 4) and the suspect. A match between the data obtained from the victim's pants (item 4) and the suspect is approximately 10 septillion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the victim's pants (item 4) and the victim is approximately 20 sextillion times less probable than a coincidental match to an unrelated person in the population. The victim is excluded from the data obtained from the victim's pants (item 4). Database(s) Used: Revised NIST 1036	

WebCode- Test	Item 4 Methods & Results
HJX7CN - 5901	Method(s) : Likelihood Ratio Stats Analysis: LR= 7.1650E12 BY ANALYSIS WITH LRmixStudio Database(s) Used: [Location Identifying Database]
J37QNH - 5902	Method(s) : Likelihood Ratio Stats Analysis: Item 1 (complainant) Excluded. Item 2 (suspect) Not Exlcuded. LR = 100 billion. Database(s) Used: [Location Identifying Database]
J772JG - 5902	Method(s): Likelihood Ratio Stats Analysis: PP21: COMP (Item 1) Excluded, SUS (Item 2) Not Excluded (LR=100 billion). YFP: SUS (Item 2) Not Excluded (LR can't be calculated as unresolved mixture). Database(s) Used: [Location Identifying Database]
JHLQHM - 5902	Method(s): Likelihood Ratio Stats Analysis: It is 4,4e26 times more likely to observe the DNA profile if the mixed stain on the victim's pants (ITEM 4) originates from ITEM 2 (Suspect) and two unknown persons, than if it originated from three unknown persons, unrelated to ITEM 2 (Suspect). Theta is 0.01 and probability of drop-in is 0.05. Database(s) Used: [Location Identifying Database]. Rare allele frequency is 0.0007
JPGUBX - 5901	Method(s): Likelihood Ratio Stats Analysis: The DNA profile from this item was interpreted as a mixture of three individuals with at least two male contributors. The DNA results are approximately 168 quadrillion times more likely if they originated from the male suspect and two unknown, unrelated individuals than if they originated from three unknown, unrelated individuals. Based on the likelihood ratio, this provides very strong support that the male suspect is a contributor to the DNA from this item. Based on the likelihood ratio result, the female victim is excluded as a possible contributor to the DNA from this item. Database(s) Used: NIST 1036 July 2017
K9ECCE - 5902	Method(s): Likelihood Ratio Stats Analysis: Item 4 - The DNA evidence is greater than 100 billion times more likely if Item 2 (Suspect) is a contributor. Item 1 (Victim) is excluded as a contributor. Database(s) Used: [Location Identifying Database]
KPDQAK - 5902	Method(s): Likelihood Ratio Stats Analysis: H1 = The evidence originated from Suspect and two unknown, unrelated individuals. H2 = The evidence originated from three unknown, unrelated individuals. The DNA profile obtained from the pants stain is approximately 16 septillion times more likely if it originated from Suspect and two unknown, unrelated individuals than if it originated from three unknown, unrelated individuals. There is evidentiary support for the inclusion of Suspect as a possible contributor to the DNA profile obtained as compared with the alternate explanation of the evidence. Database(s) Used: NIST revised: African American, Asian, Caucasian, Hispanic
L67JVL - 5901	Method(s): Likelihood Ratio Stats Analysis: Female Victim (K1) is excluded based on the LR of zero. The evidence DNA profile is approximately 2 quadrillion times more likely if it originated from Male Suspect (K2), and two unknown individuals than if it originated from three unknown individuals. This analysis provides very strong support for inclusion. Database(s) Used: allele frequencies from the 2015 FBI Expanded Population Database were used for STR statistical calculations.

WebCode- Test	Item 4 Methods & Results					
L8XEJE - 5901	Method(s): Likelihood Ratio Stats Analysis: Item 1 - 0 (excluded). Item 2 - LR = 1.4556E21 (>100 billion support for contribution). Database(s) Used: [Participant did not return database used.]					
LBAKVF - 5902	Method(s): Likelihood Ratio Stats Analysis: SUSPECT (Q02-1) cannot be excluded as a contributor to Mixture 2 from the pants (Q04-1) from the VICTIM. The STR DNA results are estimated to be 1.8 billion times more likely if they originate from SUSPECT and two unknown people than they originate from three unknown people, unrelated to SUSPECT. Database(s) Used: i. NIST Asian, ii. NIST African American, iii. NIST Caucasian, iv. Srivastava et al. (2019) South Asian, v. [Location Identifying Database]					
LG3PNJ - 5901	Method(s) : Likelihood Ratio Stats Analysis: Known contributors under Hp: suspect. Known contributors under Hd: - LR (total): 8,9E19. Database(s) Used: [Location Identifying Database]					
LHUN3H - 5902	 Method(s): Likelihood Ratio Stats Analysis: A DNA match was identified between the data obtained from the stain from victim's pants (Item 4) and Suspect. A match between the data obtained from the stain from victim's pants (Item 4) and Suspect is approximately 40 sextillion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the stain from the stain from victim's pants (Item 4) and Victim is approximately 300 quintillion times less probable than a coincidental match to an unrelated person in the population. Victim is excluded from the data obtained from the stain from victim's pants (Item 4). Database(s) Used: The National Institute of Standards and Technology's 1036 Revised U.S. combined population database was used to generate DNA match statistics. The combined population database contains African American, Caucasian, Hispanic and Asian populations. Identical siblings will have identical DNA data and DNA match statistics. 					
LUPUWF - 5901	Method(s): Likelihood Ratio Stats Analysis: The LR value regarding the possible involvement of the male suspect was calculated to be 1.39 x 10^17 to 1, which means it is about 1.39 x 10^17 to 1 times more likely that the observed DNA profile being a mixture originating from the male suspect and two unknown individuals than if it originating from three unrelated individuals selected at random from the local [Population]. Database(s) Used: [Location Identifying Database]					
LYKQJL - 5901	Method(s): Likelihood Ratio Stats Analysis: The evidence DNA profile is approximately 142 trillion times more likely if it originated from Male Suspect (K2) and two unknown individuals than if it originated from 3 unknown individuals. This analysis provides very strong support for the proposition that Male Suspect (K2) is a contributor to the DNA obtained from this sub-item. The evidence DNA profile is approximately 0 times more likely if it originated from Female Victim (K1) (Hypothesis 1) than if it originated from an unknown individual (Hypothesis 2). This analysis provides an exclusion for the proposition that Female Victim (K1) is a contributor to the DNA obtained from this sub-item. Database(s) Used: 2015 FBI Expanded Population Database					
TABLE 8						
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WebCode- Test	Item 4 Methods & Results					
MW99ZF - 5902	Method(s): Likelihood Ratio Stats Analysis: A DNA match was identified between the data obtained from the questioned stain from victim's pants (Item 4) and suspect. A match between the data obtained from the questioned stain from victim's pants (Item 4) and suspect is approximately 500 sextillion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the questioned stain from victim's pants (Item 4) and victim is approximately 7 sextillion times less probable than a coincidental match to an unrelated person in the population. Victim is excluded from the data obtained from the questioned stain from victim's pants (Item 4). Database(s) Used: National Institute of Standards and Technology's 1036 Revised U.S. combined population database.					
NWNRKD - 5901	Method(s): Likelihood Ratio Stats Analysis: The calculated LR value for the possible involvement of the suspect in this mixture was 4.25E+24 to 1. This means it is about 4.25E+24 times more likely that the observed DNA profile being a mixture originating from the suspect and two unknown persons than if it originating from three unrelated persons selected at random from the local [Population]. Database(s) Used: [Location Identifying Database]					
P3QH3A - 5901	Method(s): Likelihood Ratio Stats Analysis: Item 4: Item 1 - excluded. Item 2 - support for contribution greater than 100 billion (2.81E17). Unresolved unknown contributor. Unresolved unknown contributor. Database(s) Used: [Location Identifying Database]					
P8KEPF - 5902	Method(s): Likelihood Ratio Stats Analysis: H1:The evidence originated from Suspect, and two unknown, unrelated individuals. H2: The evidence originated from three unknown, unrelated individuals. The DNA profile obtained from Item 4 is approximately 110 sextillion times more likely if it originated from Suspect and two unknown, unrelated individuals than if it originated from three unknown, unrelated individuals. There is evidentiary support for the inclusion of Suspect as a possible contributor to the DNA profile obtained as compared with the alternate explanation of the evidence. Database(s) Used: NIST revised: African American, Asian, Caucasian, Hispanic					
P8TP7Q - 5901	Method(s): Likelihood Ratio Stats Analysis: Item 4.1 (Portion of "Questioned stain from victim's pants"): The DNA profile from this item was interpreted as a mixture of three individuals, with at least one male contributor. The DNA results are approximately 10.8 quintillion times more likely if they originated from the male suspect and two unknown, unrelated individuals than if they originated from three unknown, unrelated individuals. Based on the likelihood ratio, this provides very strong support that the male suspect is a contributor to the DNA from this item. Based on the likelihood ratio result, the female victim is excluded as a possible contributor to the DNA from this item. Database(s) Used: NIST 1036 July 2017					
PD88ZF - 5902	Method(s): Likelihood Ratio Stats Analysis: The DNA profile obtained from the epithelial fraction was interpreted as a mixture of three individuals. The DNA profile is at least 1 trillion times more likely if it originated from the Suspect (2) and two unknown individuals than if it had originated from three unknown, unrelated individuals. Statistical analysis provides very strong support for the inclusion of the Suspect (2). The victim (1) is excluded as a contributor to the DNA profile. Database(s) Used: The FBI Extended CODIS core allele frequencies - Caucasian, African American, and Southwest Hispanics					

(145)

TABLE 8					
WebCode-	Item 4 Methods & Results				
Test					
Q99BQE - 5901	Method(s): Likelihood Ratio Stats Analysis: [Participant did not return statistical analysis.] Database(s) Used: STRMIX 2.8.0 used The pants belongs to the victim. The victim (item 1) is not contributor to the questioned stain, but the suspect (item 2) is contributor to the questioned stain. The suspect's contribution in the mixture is calculated according to the following hypothesis : Hp : the mixture belongs to the suspect and 2 unknown people. Hd : the mixture belongs to 3 unknow people. LR = 2.7357E25.				
R3MFPB - 5901	Method(s): Likelihood Ratio Stats Analysis: The LR value for the involvement of male suspect is 8.65E+22 to 1, which means it is about 8.65E+22 times more likely that the observed DNA profile being a mixture originating from the male suspect and two unknown individuals than if it originating from three unrelated individuals selected at random from the local [Population]. Database(s) Used: [Location Identifying Database]				
R9RDYE - 5901	Method(s): Likelihood Ratio Stats Analysis: The evidence DNA profile is approximately 0 times more likely if it originated from Female Victim (K1) and two unknown individuals than if it originated from three unknown individuals. Female Victim (K1) is excluded. The evidence DNA profile is approximately 1 Quadrillion times more likely if it originated from Male Suspect and two unknown individuals than if it originated from three unknown individuals. This provides very strong support for the inclusion of Male Suspect (K2). Database(s) Used: 2015 FBI Expanded Population Database.				
RCPXZD - 5901	Method(s): Likelihood Ratio Stats Analysis: LR was calculated assuming the mixture is comprised of three contributors. The DNA profile is approximately 8*10 ^ 15 (8 billiard) times more likely if it originated from the Suspect and two unknown individuals than if it originated from the three unknown individuals. Database(s) Used: [Location Identifying Database]				
TMDEV4 - 5902	Method(s) : Likelihood Ratio Stats Analysis: [Participant did not return statistical analysis.] Database(s) Used: FBI_EXTENDED_CAUC				
TZBZVA - 5901	Method(s): Likelihood Ratio Stats Analysis: The LR value calculated for the possible involvement of the male suspect was 3.87E25 to 1, which means it is about 3.87E25 times more likely that the observed DNA profile being a mixture originating from the male suspect and two unknown individuals than if it originating from three unrelated individuals selected at random from the local [Population]. Database(s) Used: [Location Identifying Database]				
UNVR99 - 5902	 Method(s): Likelihood Ratio Stats Analysis: A DNA match was identified between the data obtained from the questioned stain from victim's pants (Item 4) and suspect. A match between the data obtained from the questioned stain from victim's pants (Item 4) and suspect is approximately 600 septillion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the questioned stain from victim is approximately 2 sextillion times less probable than a coincidental match to an unrelated from the questioned stain from victim's pants (Item 4) and victim is approximately 2 sextillion times less probable than a coincidental match to an unrelated person in the population. The victim is excluded from the data obtained from the questioned stain from victim's pants (Item 4). Database(s) Used: The National Institute of Standards and Technology's 1036 Revised U.S. combined population database. 				

TABLE 8						
WebCode- Test	Item 4 Methods & Results					
VFHZHA - 5902	Method(s): Likelihood Ratio Stats Analysis: Hypothesis 1 (H1): The evidence originated from Suspect and two unknown, unrelated individuals. Hypothesis 2 (H2): The evidence originated from three unknown, unrelated individuals. The DNA profile obtained from the stain from the victim's pants is approximately 1.6 septillion times more likely if it originated from Suspect and two unknown, unrelated individuals than if it originated from three unknown, unrelated individuals. Database(s) Used: NIST revised: African American, Asian, Caucasian, Hispanic.					
VK9834 - 5902	Method(s): Likelihood Ratio Stats Analysis: H1: The DNA on Item 4 came from suspect and two unknown, unrelated persons. H2: The DNA on Item 4 came from three unknown, unrelated persons. LR= 3.4804E24. H1: The DNA on Item 4 came from victim and two unknown, unrelated persons. H2: The DNA on Item 4 came from three unknown, unrelated persons. LR= 0 Database(s) Used: [Location Identifying Database]					
W2RLKZ - 5901	Method(s): Likelihood Ratio Stats Analysis: tem 4.1 (Portion of "Questioned stain from victim's pants"): The DNA profile from this item was interpreted as a mixture of three individuals with at least two male contributors. The DNA results are approximately 113 quintillion times more likely if they originated from The Male Suspect and two unknown, unrelated individuals than if they originated from three unknown, unrelated individuals. Based on the likelihood ratio, this provides very strong support that The Male Suspect is a contributor to the DNA from this item. Based on the likelihood ratio, The Female Victim is excluded as a possible contributor to this DNA profile Database(s) Used: NIST 1036 July 2017					
WRGKYZ - 5901	Method(s): Likelihood Ratio Stats Analysis: Item 4.1 (Portion of "Questioned stain from victim's pants"): The DNA profile from this item was interpreted as a mixture of three individuals with at least two male contributors. The DNA results are approximately 1.02 octillion times more likely if they originated from Male Suspect and two unknown, unrelated individuals than if they originated from three unknown, unrelated individuals. Based on the likelihood ratio, this provides very strong support that Male Suspect is a contributor to the DNA from this item. Female Victim is excluded as a possible contributor to the DNA from this item. Database(s) Used: NIST 1036 July 2017					
WU8M28 - 5901	 Method(s): Likelihood Ratio Stats Analysis: The DNA profile obtained from the sperm fraction is a mixture. No further conclusions can be drawn due to an insufficient amount of DNA for interpretation. The DNA profile obtained from the epithelial fraction was interpreted as a mixture of three individuals. The DNA profile is at least 1 trillion times more likely if it originated from male suspect (2) and two unknown individuals than if it had originated from three unknown, unrelated individuals. Statistical analysis provides very strong support for the inclusion of male suspect (2). Female victim (1) is excluded as a contributor to the DNA profile. Database(s) Used: FBI Extended CODIS Core Allele Frequencies- Caucasian, African American, and Southwest Hispanics 					
X28843 - 5901	Method(s): Likelihood Ratio Stats Analysis: Item 1 - excluded as a contributor of DNA. Item 2 - greater than 100 billion supporting contribution. Database(s) Used: [Location Identifying Database]					

	TABLE 8						
WebCode- Test	Item 4 Methods & Results						
YGJ6L9 - 5901	Method(s): Likelihood Ratio Stats Analysis: Victim (item 1) is excluded with a likelihood ratio of 0. The evidence DNA profile is approximately 56 trillion times more likely if it originated from the suspect (item 2) and 2 unknown individuals than if it originated from 3 unknown individuals. This provides very strong support for inclusion of suspect (item 2). Database(s) Used: 2015 FBI Expanded Population Database						
YJMUQ3 - 5901	Method(s): Likelihood Ratio Stats Analysis: Assuming three contributors, including at least three males, the DNA typing results were interpreted using probabilistic genotyping software. The DNA typing results are 552 sextillion times more likely to be observed if they originated from the Suspect (Item 2), and two unknown, unrelated individuals, rather than if they originated from three unknown, unrelated individuals. This analysis provides very strong support for the proposition that includes the Suspect (Item 2) as one of the contributors to this mixture. The Victim (Item 1) is excluded as a contributor to this mixture. Database(s) Used: NIST 1036 (2017)						
Z8LUQ4 - 5902	Method(s): Likelihood Ratio Stats Analysis: A DNA match was identified between the data obtained from the cutting from the blue fabric (item 4) and the subject. A match between the data obtained from the cutting from the blue fabric (item 4) and the subject is approximately 800 sextillion times more probable than a coincidental match to an unrelated person in the population. A DNA match between the data obtained from the cutting from the blue fabric (item 4) and the victim is approximately 2 sextillion times less probable than a coincidental match to an unrelated person in the population. The victim is excluded from the data obtained from the cutting from the blue fabric (item 4). Database(s) Used: The National Institute of Standards and Technology's 1036 Revised U.S. combined population database						

Additional Comments

TABLE 9

WebCode- Test	Additional Comments
2QGMC2 - 5902	Two peaks in the second fraction of item 3 (3 sp) were identified as possible elevated stutter; however, neither peak could be confirmed with re-amplification due to lower level data during the re-amp. The peaks are 27.2 at SE33 and 24 at D2S1338.
34G3B3 - 5902	For items 3 and 4, the results include all detected alleles along with the associated stutter peaks. This reflects the data that is imported into STRmix in order to perform probabilistic genotyping.
44PDRD - 5901	** Possible Stutter
4BEUJW - 5902	The results obtained and presented in this report were generated following lab procedures, however I am not authorised to report the following in casework: Y Profiling/YSTR testing. I can only determine if a Y quant has been detected. The quality of the DNA may not have been ideal in this test. The DNA results for Item 3 (epithelial/non-sperm fraction) had low levels of DNA (0.005ng/ul), requiring re-amp. In case work, I may have considered taking a second sample, however the very narrow time frame allowed for this test did not allow time for resampling.
77A6RY - 5901	Reported/listed alleles for evidence samples include stutter peaks. Our laboratory does not report DNA concentration or DNA proportions.
786NUZ - 5901	DNA mixture concentration and proportions: DNA mixture proportions are reported as computed by probabilistic genotyping software (EuroForMIX). DNA concentrations (ng/ul) for each contributor is reported as concentration of item obtained by real time PCR analysis multiplied by mixture proportion of that contributor, divided by 100.
8Y8DYV - 5902	PCR amplification procedures were not performed on the second fraction from the "questioned stainunderwear" (item 3) due to an insufficient amount of DNA detected using the Quantifiler Trio DNA Quantification Kit on the Applied Biosystems 7500 real-time PCR system.
9CLYK9 - 5901	NR=No results. **= Possible elevated stutter.
9MBR8Q - 5901	Item 3 - S#. Resolved contributor (C2) - Designated Unknown Male 1. Resolved contributor (C3) - Item 2 (known suspect).
9WLTTR - 5902	This laboratory truncates likelihood ratios favouring contribution at 100 billion. For reporting purposes any figure greater than is reported as 'greater than 100 billion'.
B7W38V - 5901	Reported/listed alleles for evidence samples include stutter peaks. Our laboratory does not report DNA concentration or proportion.
BDR7Z7 - 5901	DNA concentration for contributor positions is not reported. Reported 99% 1-sided Lower HPD for statistical analysis. Reported lowest HPD of the three databases used. Reported verbal scale endorsed by the Scientific Working Group on DNA Analysis Methods. NID= No Interpretable Data. No LR stat reported for fraction 2 of item 3 (1Af2) due to exclusions for item 1 and item 2. Carry over from f1 fraction (Penta E, D2S1338, D21, D12, and SE33) at 5 markers. Trace contributor (contributor 2) was unsuitable for comparison.
BKHHBU - 5901	Reported/listed alleles for evidence samples include stutter peaks. Our laboratory does not report DNA concentration or proportion.
BNGT8U - 5902	For items 3 and 4, the results include all detected alleles along with the associated stutter peaks. This reflects the data that is imported into STRmix in order to perform probabilistic genotyping.
CXYRCQ - 5902	Item 1 was tested using the RapidHIT ID System in accordance with internal approved policies. Note that per internal validated protocol, partial information was obtained at Amelogenin but is approved for use for comparison purposes and at D10S1248 which was marked as inconclusive and not suitable for comparison.

TABLE 9

WebCode- Test	Additional Comments
CY6DP6 - 5901	** = possible elevated stutter
HJX7CN - 5901	Item 4 : we reach to total 4 contributor (inclusive of sperm contributor) based upon the estimate of the DNA proportion concentration percentage.
LHUN3H - 5902	A "partial" profile was obtained for item 1 due to the homozygous allele at D10S1248 being below the stochastic threshold.
P3QH3A - 5901	Please note: DNA Proportions are obtained from the STRmix output file. [Laboratory] does not use the DNA proportions to calculate the DNA concentrations of each contributor.
P8TP7Q - 5901	** = possible elevated stutter.
PD88ZF - 5902	Y-STR DNA profiles of indistinguishable mixtures of two or more individuals are not interpreted per our laboratory procedures.
Q99BQE - 5901	Although the item 4 is supposed to be victim's pants, the genetic profil of the female victim (item 1) is not found in the mixture obtained from item 4. The unknown contributor of item 3 sp is not found in the mixutre of item 4.
UNVR99 - 5902	The second (sperm) faction is inconclusive due to the complexity of the mixture. The mixture weights of two concordant runs did not reflect the data. The data had a major contributor with some minor alleles. Review of the data in TrueAllele showed five alleles at SE33 making it an at least three person mixture. Two concordant runs in TrueAllele had mixture weights of ~52,46,2 and ~50,49,1 which, as stated above, is not reflective of the data. Item 4 was run on the RapidHIT system (twice). The allele at D10S1248 did not meet our stochastic threshold; therefore, it was not used for comparison or statistical calculations.
W2RLKZ - 5901	NR: No results. **: Possible Stutter peak.
WRGKYZ - 5901	** = Possible Elevated Stutter
XCU6U9 - 5902	Inconclusive Y-STR DNA results were obtained from the following item(s). These results are not suitable for comparison purposes due to the poor quality or complexity of the data. Item 1.3F1 - The results are best described as an indistinguishable mixture of 2 male individuals and no assumptions can be made. Item 1.3F2 - The results are best described as an indistinguishable mixture of 2 male individuals and no assumptions can be made. Item 1.4 - The results are best described as a mixture of >2 male individuals.
YJMUQ3 - 5901	The number of contributors could not be determined for the epithelial cell fraction of Item 3 due to low-level results. STRmix deconvolution was not conducted on this sample. ND = Not Detected. All potential alleles considered during deconvolution have been reported.

Test No. 25-5901: Probabilistic Genotyping

DATA MUST BE SUBMITTED BY April 14, 2025, 11:59 p.m. EDT TO BE INCLUDED IN THE REPORT

Participant Code: U1234A

WebCode: MCZ9PU

The Accreditation Release section can be accessed by using the "Continue to Final Submission" button above. This information can be entered at any time prior to submitting to CTS.

Scenario:

Police are investigating a sexual assault involving a female victim and male suspect. The victim reported to police that her boyfriend and several of his friends assaulted her when she was drunk. Her boyfriend, the male suspect, has been apprehended by police. The other men have not yet been identified. Police are submitting a stain from the underwear (Item 3) and a stain from the pants (Item 4) the victim was wearing at the time of the assault. Also provided are known standards from the female victim (Item 1) and male suspect (Item 2).

Items Submitted (Sample Pack G1 - Cloth Swatches):

Item 1: Known blood from the female victim

Item 2: Known blood from the male suspect

Item 3: Questioned stain from victim's underwear (tan)

Item 4: Questioned stain from victim's pants (blue)

Part I: SCREENING TESTS

Note: Laboratories submitting their results for ASCLD/LAB or NATA accreditation MUST identify any screening tests performed and report the test results.

Indicate the results of any screening tests performed on the questioned stains (Items 3 & 4).

Please use the abbreviations listed in this response key to fill in the Screening Test tables on this tab. This is not an all inclusive list of tests, and should not be used to determine what tests should be performed. TESTS NOT ON THIS LIST MAY BE USED FOR SCREENING.

Test	Abbreviation	Test	Abbreviation
Acid Phosphatase	AP	Alternate Light Source	ALS
Kastle Meyer	KM	Leucomalachite Green	LMG
Microscopic	Micro	Ortho-tolidine	O-tol
Phenolphthalein-Tetramethyl benzidine	PTMB	Prostate Specific Antigen	PSA
Rapid Stain Identification	RSID	Tetramethyl benzidine	ТМВ

Example:	<u>Positive</u>	<u>Negative</u>	<u>Inconclusive</u>	Not Tested	Test(s) Performed
Blood					KM, O-tol, PTMB
Semen					PSA
Saliva				۲	
Human Origin					
Y-Screening (male DNA)					
Other Specified Body Fluid					

Please indicate the Test(s) Performed on the corresponding line for each type of screening.

Screening data not reported. \square

Item 3:	<u>Positive</u>	<u>Negative</u>	Inconclusive	Not Tested	Test(s) Performed
Blood	\bigcirc	\bigcirc	\bigcirc	\bigcirc	
Semen	\bigcirc	\bigcirc	\bigcirc	\bigcirc	
Saliva	\bigcirc	\bigcirc	\bigcirc	\bigcirc	
Human Origin	\bigcirc	\bigcirc	\bigcirc	\bigcirc	
Y-Screening (male DNA)	\bigcirc	\bigcirc	\bigcirc	\bigcirc	
Other:	\bigcirc	\bigcirc	0	\bigcirc	

Item 4:	Positive	<u>Negative</u>	Inconclusive	Not Tested	Test(s) Performed
Blood	\bigcirc	\bigcirc	\bigcirc	\bigcirc	
Semen	\bigcirc	\bigcirc	\bigcirc	\bigcirc	
Saliva	\bigcirc	\bigcirc	\bigcirc	\bigcirc	
Human Origin	\bigcirc	\bigcirc	\bigcirc	\bigcirc	
Y-Screening (male DNA)	\bigcirc	\bigcirc	\bigcirc	\bigcirc	
Other:	0	\bigcirc	0	0	

Part II: DNA INTERPRETATION

Based on results obtained from DNA analysis, could the Victim (Item 1) and/or the Suspect (Item 2) be a contributor to the questioned stains (Items 3 & 4)?

	<u>Victim (It</u>	<u>em 1)</u>	Suspect (Item 2)		
	Item 3	Item 4		Item 3	Item 4
Yes			Yes		
No			No		
Inconclusive			Inconclusive		
No Interpretation			No Interpretation		

Part III: DNA Results for Known Item 1

- Report alleles in numerical order, separated by a comma.
 Follow your laboratory procedures for reporting homozygotes (i.e. X,X or X) and null responses.
 If your laboratory policy is to indicate minor or weaker alleles, please enclose them within brackets [].

STR Amplification Kit(s) Used:		Please check all the branc information in the blank p					
Identifiler®		🗖 GlobalFiler	☐ GlobalFiler™ Investig				
PowerPlex®		Other					
Report the Probabilistic Genotyping Software Used: STRmix TrueAllele Other							
Alleles below are	sorted in Defau	lt order.					
ITEM	D1S1656	D2S1338	D2S441	D3S1358	D5S818	D6S1043	
1							

1						
ITEM	D7S820	D8S1179	D10S1248	D12S391	D13S317	D16S539
1						
ITEM	D18S51	D19S433	D21S11	D22S1045	Amelogenin	CSF1PO
1						
ITEM	FGA	Penta D	Penta E	SE33	TH01	TPOX
1						
ITEM	vWA	DYS391	DYS570	DYS576	Y Indel	
1						

Part III (continued): DNA Results for Known Item 2

- Report alleles in numerical order, separated by a comma.
- Follow your laboratory procedures for reporting homozygotes (i.e. X,X or X) and null responses.

STR Amplification	on Kit(s) Used:	Please check all the bran information in the blank	nds that apply for this ite provided (i.e. 16, Plus,	em and record only add Direct, HS, Fusion, etc.	itional kit specific).	
Identifiler®		📃 GlobalFile	er™	Inv	estigator® 24plex	
PowerPlex®		Other				
Report the Probab	ilistic Genotyping	Software Used:				
STRmix		TrueAllele	2	Othe	r	
Illeles below are	sorted in Defa	ult order.	D25441	D3S1358	D55818	D6S1043
2						
ITEM	D7S820	D8S1179	D10S1248	D12S391	D13S317	D16S539
2						
ITEM	D18S51	D19S433	D21S11	D22S1045	Amelogenin	CSF1PO
2						
ITEM	FGA	Penta D	Penta E	SE33	TH01	TPOX
2						

DYS570

YSTR Amplification Kit(s) Used:

Please check all the brands that apply for this item and record only additional kit specific information in the blank provided (i.e. Plus, 23, etc.).

DYS576

I YFiler™

ITEM

2

PowerPlex® Y

DYS391

Other

Y Indel

Alleles below are sorted in Default order.

vWA

ITEM	DYF387S1	DYS19	DYS385	DYS389-1	DYS389-II	DYS390	DYS391	DYS392	DYS393
2									
ITEM	DYS437	DYS438	DYS439	DYS448	DYS449	DYS456	DYS458	DYS460	DYS481
2									
ITEM	DYS518	DYS533	DYS549	DYS570	DYS576	DYS627	DYS635	DYS643	Y GATA H4
2									

Part III (continued): DNA Results for Questioned Item 3

- Report alleles in numerical order, separated by a comma.Follow your laboratory procedures for reporting homozygotes (i.e. X,X or X) and null responses.

							,		
٠	If your laboratory	policy is to indicate	e minor	or weaker	^r alleles,	, please	enclose the	m within bracke	ts

	<u>on Kit(s) Usea:</u>	information in the blan	ik provided (i.	e. 16, Plus, Di	irect, HS, Fusion, etc.).		
Identifiler®		🗖 GlobalFi	iler™		lnv	vestigator® 24pl	ex	
PowerPlex®		Other						
Report the Probab	ilistic Genotypin	g Software Used:						
STRmix		TrueAlle	le		Othe	r		
Alleles below are	Did y sorted in Defa	vou perform a differ ault <i>order</i> .	ential ext	raction of	Item 3? YES	NO 🔍		
ITEM	D1S1656	D2S1338	D29	5441	D3S1358	D5S818		D6S1043
3								
3 e								
3 sp								
ITEM	D7S820	D8S1179	D105	51248	D12S391	D13S317	′ I	D16S539
3								
3 e								
3 sp								
ITEM	D18S51	D19S433	D2 ⁻	1S11	D22S1045	Amelogen	in	CSF1PO
3								
3 e								
3 sp								
ITEM	FGA	Penta D	Per	nta E	SE33	TH01		TPOX
3								
3 e								
3 sp								
ITEM	VWA	DYS391	DYS	5570	DYS576	Y Indel		
3								
3 e								
3 sp								
YSTR Amplificat	<u>ion Kit(s) Used</u>	Please check all the black information in the black	orands that ap	ply for this ite	em and record only ac	Iditional kit specifi	с	
YFiler™		PowerPl	ex® Y		0	ther		
	Did y	ou perform a differ	ential ext	raction of	Item 3? YES	○ NO ○		
lleles below are	sorted in Defa	ault order.						
ITEM DYF	387S1 DYS	DYS385	DYS389-1	DYS389-	II DYS390	DYS391	DYS392	DYS39
3								
3 e								
3 sp								
		12.0 D) (5.12.0	D)/C 4 40	D)/C 4 40		D)/C (F0	D)/C // C	

ITEM	DYF387S1	DYS19	DYS385	DYS389-1	DYS389-II	DYS390	DYS391	DYS392	DYS393
3									
3 e									
3 sp									
ITEM	DYS437	DYS438	DYS439	DYS448	DYS449	DYS456	DYS458	DYS460	DYS481
3									
3 e									
3 sp									
ITEM	DYS518	DYS533	DYS549	DYS570	DYS576	DYS627	DYS635	DYS643	Y GATA H4
3									
3 e									
3 sp									

Part III (continued): DNA Results for Questioned Item 4

- Report alleles in numerical order, separated by a comma.
 Follow your laboratory procedures for reporting homozygotes (i.e. X,X or X) and null responses.

				(, , , , , , , , , , , , , , , , , , ,	
٠	If your laboratory	y policy is to indicate	e minor or weaker alleles,	, please enclose them wit	hin brackets [].

STR Amplification	<u>on Kit(s) Used:</u>	Please check all the bran	ds that apply for this ite	m and record only add	itional kit specific	
Identifiler®		GlobalFile	r™		/estigator® 24plex	
PowerPlex®		Other			5	
Report the Probab	ilistic Genotyping So	ftware Used:				
				0.1		
STRMIX		IrueAllele		Othe	r	
	Did you	perform a differei	ntial extraction o	f Item 4? YES	NO O	
leles below are	sorted in Default	order.				
ITEM	D1S1656	D2S1338	D2S441	D3S1358	D5S818	D6S1043
4						
4 e						
4 sp						
ITEM	D7S820	D8S1179	D10S1248	D12S391	D13S317	D16S539
4						
4 e						
4 sp						
ITEM	D18S51	D19S433	D21S11	D22S1045	Amelogenin	CSF1PO
4						
4 e						
4 sp						
ITEM	FGA	Penta D	Penta E	SE33	TH01	TPOX
4						
4 e						
4 sp						
ITEM	vWA	DYS391	DYS570	DYS576	Y Indel	
4						
4 e						
4 sp						

Did you perform a differential extraction of Item 4? $\,$ YES \odot $\,$ NO $\,$

Alleles below are sorted in Default order.

ITEM	DYF387S1	DYS19	DYS385	DYS389-1	DYS389-II	DYS390	DYS391	DYS392	DYS393
4									
4 e									
4 sp									
ITEM	DYS437	DYS438	DYS439	DYS448	DYS449	DYS456	DYS458	DYS460	DYS481
4									
4 e									
4 sp									
ITEM	DYS518	DYS533	DYS549	DYS570	DYS576	DYS627	DYS635	DYS643	Y GATA H4
4									
4 e									
4 sp									

Part III (continued): DNA Analysis - Additional DNA

- Use this section to report results for loci not currently listed in other sections of the data sheet.
- Report alleles in numerical order, separated by a comma.
- If your laboratory policy is to indicate minor or weaker alleles, please enclose them within brackets [].
- Click "Add Row" to show another row of boxes for entry.

Did you perform a differential extraction of Item 3? $\,$ Yes $\,$ No $\,$

Did you perform a differential extraction of Item 4? $\,$ Yes $\,$ $\,$ No $\,$ $\,$

Locus	ltem 1	ltem 2	ltem 3	ltem 3e	ltem 3sp	ltem 4	ltem 4e	ltem 4sp

Item 3:

Part IV: Mixture Sample Analysis

NOTE: To allow functionality of this page please select an answer to differential extraction questions on previous tabs for Item 3 and Item 4.

For each item, select the Estimated Number of Contributors and Contributor Identification (Victim, Suspect, or Unknown Individual). Calculate the DNA Concentration and DNA Proportion for each contributor using your laboratory's protocols. For Unknown Individual(s), report the contributor determined to have the highest concentration of DNA first and remaining in descending order. Enter "DNA Concentration" in ng/uL and "DNA Proportion" in percentage.

Concentration and proportion data not reported. \square

Estimated number of contributors:	DNA Concentration (ng/uL)	DNA Proportion (%)
Contributor 1		
Item 4:		
Estimated number of contributors:		DNA
	DNA Concentration (ng/uL)	Proportion (%)
Contributor 1		

Part V: DNA Statistical Analysis

Item 3:

Likelihood Ratio (LR)	Other	
Note: Please use appropriate punctuation to indicond returns used for separation within your text will no and tabular formats to deliver information is also c	ate the end of sentences, sections, and staten t transfer and may cause your information to autioned against, as these do not transfer.	nents in the free-form space below. Extra spacing a be illegible in the Summary Report. The use of list
Please list any databases used in the statistical a	analysis of Item 3 below	
teuse tist any databases used in the statistical		
m 4: Statistical Analysis of Item 4 DNA Typing Select the statistical method(s) used by marking the assu	Results: ociated box and report these results in the spa	ce below:
m 4: Statistical Analysis of Item 4 DNA Typing Select the statistical method(s) used by marking the asso Likelihood Ratio (LR)	Results: ociated box and report these results in the spa Other	ce below:
 m 4: Statistical Analysis of Item 4 DNA Typing Select the statistical method(s) used by marking the assu Likelihood Ratio (LR) Note: Please use appropriate punctuation to indicate returns used for separation within your text will not and tabular formats to deliver information is also c 	Results: pociated box and report these results in the spa Other ate the end of sentences, sections, and staten t transfer and may cause your information to autioned against, as these do not transfer.	ce below: hents in the free-form space below. Extra spacing a be illegible in the Summary Report. The use of list
m 4: Statistical Analysis of Item 4 DNA Typing Select the statistical method(s) used by marking the asse Likelihood Ratio (LR) Note: Please use appropriate punctuation to indica returns used for separation within your text will no and tabular formats to deliver information is also c	Results: bociated box and report these results in the spa Other ate the end of sentences, sections, and staten t transfer and may cause your information to autioned against, as these do not transfer.	ce below: nents in the free-form space below. Extra spacing a be illegible in the Summary Report. The use of list

Part VI: ADDITIONAL COMMENTS

- Use this section to report comments regarding any part of this test.
- Written conclusions (including statistical information) for DNA analysis are not required.
- Note: Laboratories submitting their results for accreditation are asked to report any additional information that will assist in the review of their results. This includes an explanation of any deviations from a full completion of the test and/or unique findings such as elevated stutter.

Note: Please use appropriate punctuation to indicate the end of sentences, sections, and statements in the free-form space below. Extra spacing and returns used for separation within your text will not transfer and may cause your information to be illegible in the Summary Report. The use of lists and tabular formats to deliver information is also cautioned against, as these do not transfer.

RELEASE OF DATA TO ACCREDITATION BODIES

The Accreditation Release is accessed by pressing the "Continue to Final Submission" button online and can be completed at any time prior to submission to CTS.

CTS submits external proficiency test data directly to ANAB and/or A2LA. Please select one of the following statements to ensure your data is handled appropriately.

This participant's data is intended for submission to ANAB and/or A2LA. (Accreditation Release section below must be completed.)
 This participant's data is not intended for submission to ANAB and/or A2LA.

Have the laboratory's designated individual complete the following steps <u>only if your laboratory is accredited in this testing/calibration discipline</u> by one or more of the following Accreditation Bodies.

Step 1: Provide the applicable Accreditation Certificate Number(s) for your laboratory	
ANAB Certificate No.	
A2LA Certificate No.	
Step 2: Complete the Laboratory Identifying Information in its entirety	
Authorized Contact Person and Title	
Laboratory Name	
Location (City/State)	