



Manufacturer's Information

Test No. 21-5881: DNA Interpretation

Each sample pack contained digital files consisting of electropherograms from DNA profiles of two known samples (Items 1 & 2) and two questioned samples (Items 3 & 4). Participants were requested to evaluate the electropherograms and interpret the data using their existing protocols.

SAMPLE PREPARATION: Item 1 was created using blood collected from a female donor. Item 2 was created using blood collected from a male donor. The Item 3 mixture was created by combining one part of blood from the Item 1 female donor and three parts of blood from a 3rd-party male donor. The Item 4 mixture was created by combining three parts of blood from the Item 2 male donor and five parts of blood from a 3rd-party male donor (same as used in Item 3).

SAMPLE SET ASSEMBLY: Once sample preparation and verification was completed, the digital upload was checked to ensure all items were accessible.

VERIFICATION: Laboratories that conducted predistribution testing of the electropherograms reported consistent results for all loci. All associations were consistent amongst the predistribution laboratories.

Consensus results on the following pages were determined by ensuring at least 10 participants returned results for the locus. Each allele listed was determined by ensuring that at least 75% of participants that returned data for that specific locus and item had reported the same allele.

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Manufacturer's Information, continued
Test No. 21-5881: DNA Interpretation

Amelogenin and STR Results						
<i>Results compiled by predistribution laboratories and a consensus of participants.</i>						
Item	D1S1656	D2S1338	D2S441	D3S1358	D5S818	D7S820
	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		
1	15,17	19,25	11,11	17,18	11,13	8,10
	11,13	14,14	18,22	12,13	11,11	14,15
	12,13	29,30	11,16	X,X	12,12	20,25
	*	*	14,16	9,9	8,11	16,19
	NM	NM	NM	NM		
2	12,12	17,18	11,15	16,17	12,14	8,8
	10,13	14,16	22,22	8,12	11,13	12,24
	14,14	28,28	11,14	X,Y	11,12	20,24
	*	*	18,25.2	9.3,9.3	8,8	17,19
	10	*	*	2		
3	15,15.3,17	19,20,25	11,11.3	16,17,18	11,12,13	8,10,11
	11,13	13,14,15	18,20,22	12,13	11,12	14,15,19,22
	12,13	29,30,32.2	11,16	X,Y	11,12	20,21,24.2,25
	*	*	14,16,29.2	7,9,9.3	8,11	14,16,19
	10	*	*	2		
4	12,15,15.3	17,18,20,25	11,11.3,15	16,17,18	12,14†	8,10,11
	10,13	13,14,15,16	18,20,22	8,12	11,12,13	12,19,22,24†
	13,14	28,30,32.2	11,14,16	X,Y	11,12	20,21,24,24.2
	*	*	18,25.2,29.2	7,9.3	8,8	14,16,17,19
	10	*	*	2		

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

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

† Additional alleles may be present depending on laboratory thresholds and/or amplification kit used.

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YSTR Results									
<i>Results compiled from predistribution laboratories and a consensus of participants.</i>									
Item	DYF387S	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
2	37,41	15	13,17	13	29	23	10	11	12
	15	9	12	21	31	16	14	10	23
	40	12	*	17	15	17	21	*	12
3	37,38	14	13,14	13	31	24	10	11	12
	16	7	11	19	28	15	19	11	23
	40	12	*	16	16	16	22	*	13
4	37,38,41	14,15	13,14,17	13	29,31	23,24	10	11	12
	15,16	7,9	11,12	19,21	28,31	15,16	14,19	10,11	23
	40	12	*	16,17	15,16	16,17	21,22	*	12,13

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