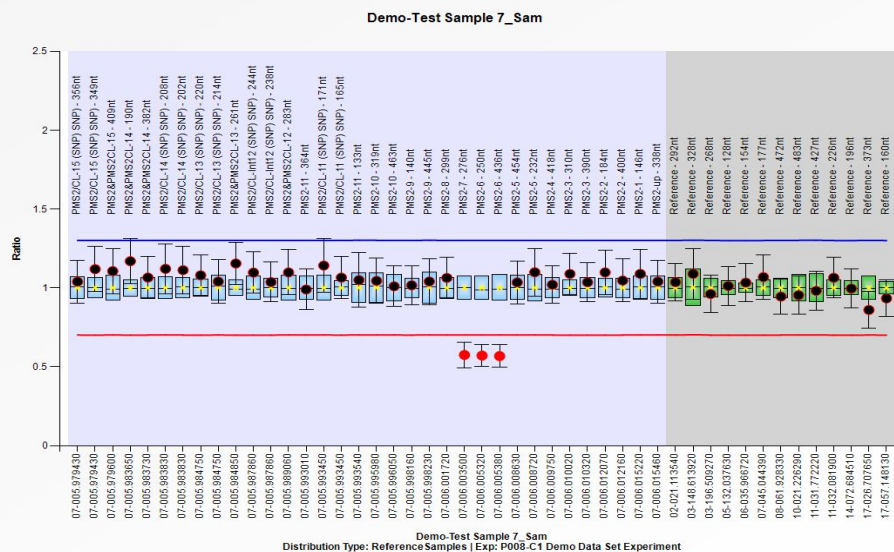
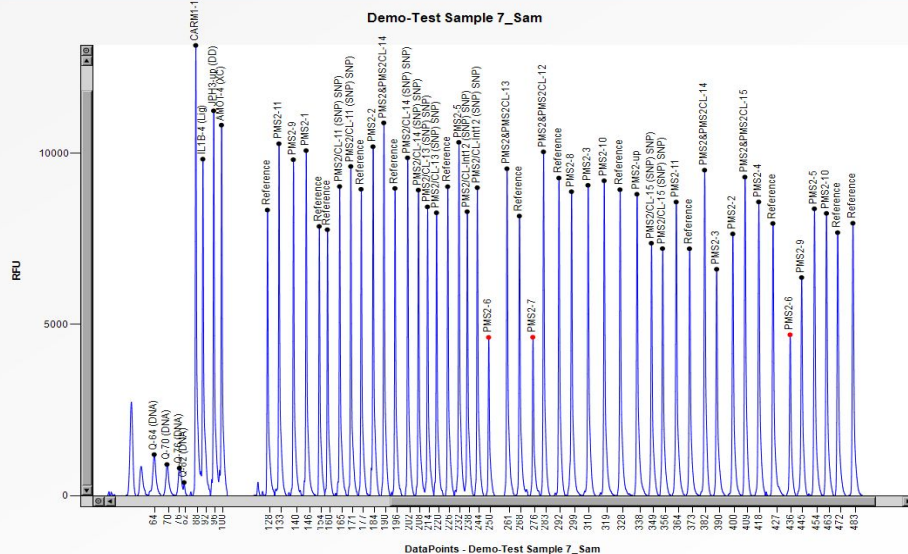


MLPA probe mix:	P008-PMS2	Nr of test probes:	47/47	FRSS:	OK 100%
Lot number:	C1-0523	Nr of ref probes	13/13	FMRS:	OK 100%
Sheet date:	4/8/2026 1:37:23 PM	DNA concentration:	OK	PSLP:	OK -5%
Control fragments:	CF-003-[brown] QDX2 (A2-0	DNA denaturation	OK	RSQ:	OK
Analysis method:	Block SSC: Off	Expected gender:	Female	RPQ:	OK
Used metric:	Peak height	Residual primer %	OK 13%	CAS:	OK 100%

Reference Samples: Demo-REF-Male-1_Ref | Demo-REF-Male-2_Ref | Demo-REF-Male-3_Ref | Demo-REF-Male-4_Ref | Demo-REF-Male-5_Ref | Demo-REF-Male-6_Ref | Demo-REF-Male-7_Ref | Demo-REF-Male-8_Ref |&...;



D [nt]	Gene-Exon	Chr.band	hg18 loc.	Height	Area	Ratio ^H	Stdev	[REF]	Width	d[nt]	[Mut details]
356	PMS2/CL-15 (S	07p22.1	07-005.979430	7216	84410	1.04	0.07	=	69	0.03	SNP
349	PMS2/CL-15 (S	07p22.1	07-005.979430	7373	83345	1.12	0.07	=	61	0.01	SNP
409	PMS2&PMS2CL-15	07p22.1	07-005.979600	9303	108108	1.11	0.07	=	58	0.09	-
190	PMS2&PMS2CL-14	07p22.1	07-005.983650	10885	104406	1.17	0.07	=	53	0.03	-
382	PMS2&PMS2CL-14	07p22.1	07-005.983730	9505	108812	1.07	0.07	=	66	0.04	-
208	PMS2/CL-14 (S	07p22.1	07-005.983830	8923	88181	1.12	0.08	=	51	0.05	SNP
202	PMS2/CL-14 (S	07p22.1	07-005.983830	9865	95088	1.11	0.08	=	56	0.00	SNP
220	PMS2/CL-13 (S	07p22.1	07-005.984750	8261	80762	1.08	0.06	=	52	0.07	SNP
214	PMS2/CL-13 (S	07p22.1	07-005.984750	8433	83876	1.04	0.07	=	51	0.06	SNP
261	PMS2&PMS2CL-13	07p22.1	07-005.984850	9548	99607	1.16	0.07	=	61	0.05	-
244	PMS2/CL-Int12	07p22.1	07-005.987860	8994	91988	1.1	0.07	=	51	0.11	SNP
238	PMS2/CL-Int12	07p22.1	07-005.987860	8293	84539	1.04	0.06	=	54	0.04	SNP
283	PMS2&PMS2CL-12	07p22.1	07-005.989060	10039	104308	1.1	0.07	=	54	0.01	-
364	PMS2-11	07p22.1	07-005.993010	8575	103016	0.99	0.06	=	73	0.04	-
171	PMS2/CL-11 (S	07p22.1	07-005.993450	9614	92754	1.14	0.09	=	57	0.11	SNP
165	PMS2/CL-11 (S	07p22.1	07-005.993450	9027	87382	1.07	0.07	=	63	0.07	SNP
133	PMS2-11	07p22.1	07-005.993540	10277	99634	1.05	0.09	=	67	0.00	-
319	PMS2-10	07p22.1	07-005.995980	9196	100377	1.04	0.07	=	96	0.05	-
463	PMS2-10	07p22.1	07-005.996050	8244	100715	1.01	0.06	=	61	0.02	-
140	PMS2-9	07p22.1	07-005.998160	9811	94185	1.02	0.06	=	68	0.01	-
445	PMS2-9	07p22.1	07-005.998230	6372	81102	1.04	0.07	=	59	0.02	-
299	PMS2-8	07p22.1	07-006.001720	8877	101467	1.06	0.06	=	84	0.09	-
276	PMS2-7	07p22.1	07-006.003500	4629	49797	0.57	0.04	<<*	59	0.00	-
250	PMS2-6	07p22.1	07-006.005320	4625	46784	0.57	0.03	<<*	51	0.06	-
436	PMS2-6	07p22.1	07-006.005380	4701	56282	0.57	0.04	<<*	57	0.06	-
454	PMS2-5	07p22.1	07-006.008630	8378	100412	1.04	0.07	=	66	0.05	-
232	PMS2-5	07p22.1	07-006.008720	10317	106936	1.1	0.08	=	54	0.07	-
418	PMS2-4	07p22.1	07-006.009750	8577	102278	1.02	0.06	=	62	0.03	-
310	PMS2-3	07p22.1	07-006.010020	9063	98809	1.09	0.07	=	72	0.06	-
390	PMS2-3	07p22.1	07-006.010320	6613	79195	1.04	0.06	=	68	0.02	-
184	PMS2-2	07p22.1	07-006.012070	10190	104498	1.1	0.07	=	55	0.00	-
400	PMS2-2	07p22.1	07-006.012160	7644	89389	1.05	0.07	=	80	0.01	-
146	PMS2-1	07p22.1	07-006.015220	10075	99522	1.09	0.08	=	73	0.04	-
338	PMS2-up	07p22.1	07-006.015460	8802	100521	1.04	0.07	=	74	0.03	-
292	Reference	02p24.1	02-021.113540	9275	97050	1.04	0.06	=	67	0.08	-
328	Reference	03q24	03-148.613920	8935	101301	1.09	0.08	=	65	0.02	-
268	Reference	03q29	03-196.509270	8166	84779	0.96	0.06	=	51	0.04	-
128	Reference	05q31.1	05-132.037630	8341	78073	1.01	0.06	=	55	0.00	-
154	Reference	06p21.31	06-035.966720	7864	73779	1.03	0.06	=	41	0.03	-
177	Reference	07p13	07-045.044390	8947	87547	1.07	0.07	=	53	0.03	-
472	Reference	08q12.2	08-061.928330	7682	93880	0.94	0.06	=	60	0.06	-
483	Reference	10p12.31	10-021.226290	7956	98064	0.95	0.06	=	80	0.02	-
427	Reference	11p13	11-031.772220	7949	96125	0.98	0.06	=	84	0.06	-
226	Reference	11p13	11-032.081900	9023	88668	1.06	0.06	=	50	0.07	-
196	Reference	14q24.2	14-072.684510	8972	88155	1	0.06	=	55	0.09	-
373	Reference	17q11.2	17-026.707650	7212	84038	0.86	0.06	=	74	0.02	-
160	Reference	17q23.2	17-057.148130	7767	73215	0.93	0.06	=	51	0.03	-
Median value all probe values:				8802	93880	1.04	0.07		60	0.04	

Median value all probe values: