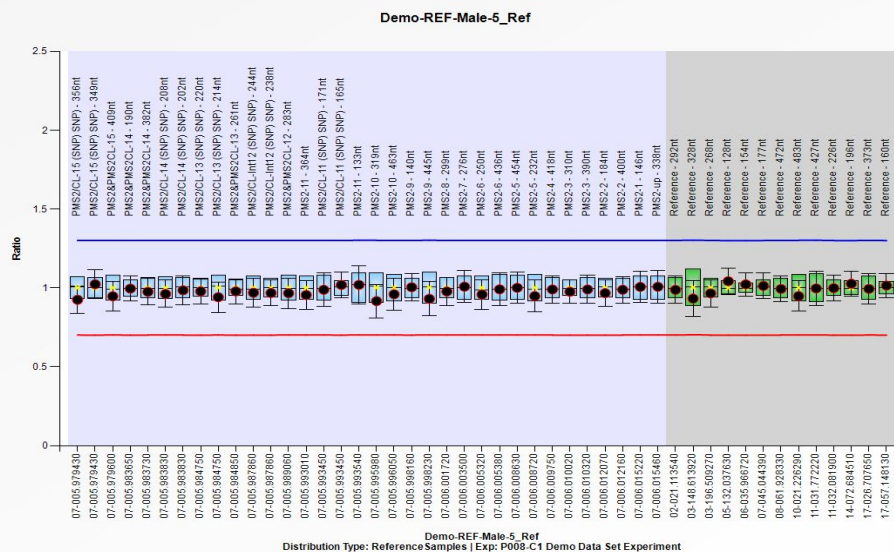
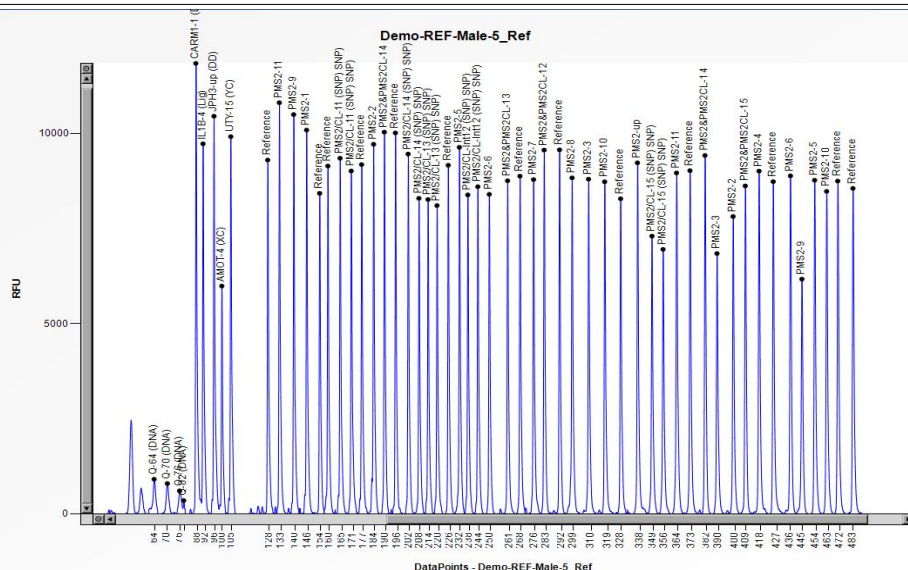


MLPA probe mix:	P008-PMS2	Nr of test probes:	47/47	FRSS:	OK 100%
Lot number:	C1-0523	Nr of ref probes	13/13	FMRS:	OK 90%
Sheet date:	4/8/2026 1:37:23 PM	DNA concentration:	OK	PSLP:	OK -3%
Control fragments:	CF-003-[brown] QDX2 (A2-0	DNA denaturation	OK	RSQ:	OK
Analysis method:	Block SSC: Off	Expected gender:	Male	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	6953	76443	0.92	0.04	=	60	0.02	SNP
349	PMS2/CL-15 (S	07p22.1	07-005.979430	7301	79030	1.02	0.05	=	54	0.08	SNP
409	PMS2&PMS2CL-15	07p22.1	07-005.979600	8621	91364	0.95	0.05	=	57	0.01	-
190	PMS2&PMS2CL-14	07p22.1	07-005.983650	10035	92153	1	0.04	=	50	0.00	-
382	PMS2&PMS2CL-14	07p22.1	07-005.983730	9421	97146	0.98	0.04	=	46	0.03	-
208	PMS2/CL-14 (S	07p22.1	07-005.983830	8295	80042	0.96	0.04	=	57	0.01	SNP
202	PMS2/CL-14 (S	07p22.1	07-005.983830	9459	88447	0.99	0.05	=	49	0.00	SNP
220	PMS2/CL-13 (S	07p22.1	07-005.984750	8104	77297	0.98	0.04	=	47	0.02	SNP
214	PMS2/CL-13 (S	07p22.1	07-005.984750	8263	79992	0.94	0.05	=	49	0.06	SNP
261	PMS2&PMS2CL-13	07p22.1	07-005.984850	8756	85794	0.98	0.04	=	59	0.07	-
244	PMS2/CL-Int12	07p22.1	07-005.987860	8600	85142	0.97	0.05	=	52	0.06	SNP
238	PMS2/CL-Int12	07p22.1	07-005.987860	8385	82697	0.97	0.04	=	46	0.04	SNP
283	PMS2&PMS2CL-12	07p22.1	07-005.989060	9563	93546	0.97	0.05	=	55	0.03	-
364	PMS2-11	07p22.1	07-005.993010	8962	102044	0.96	0.05	=	61	0.04	-
171	PMS2/CL-11 (S	07p22.1	07-005.993450	9013	88326	0.99	0.05	=	58	0.01	SNP
165	PMS2/CL-11 (S	07p22.1	07-005.993450	9348	90571	1.02	0.04	=	54	0.02	SNP
133	PMS2-11	07p22.1	07-005.993540	10808	101184	1.02	0.06	=	53	0.05	-
319	PMS2-10	07p22.1	07-005.995980	8730	89815	0.92	0.05	=	66	0.02	-
463	PMS2-10	07p22.1	07-005.996050	8477	96331	0.96	0.05	=	58	0.00	-
140	PMS2-9	07p22.1	07-005.998160	10496	98364	1	0.04	=	45	0.00	-
445	PMS2-9	07p22.1	07-005.998230	6171	72087	0.93	0.05	=	58	0.03	-
299	PMS2-8	07p22.1	07-006.001720	8832	96456	0.98	0.04	=	77	0.08	-
276	PMS2-7	07p22.1	07-006.003500	8788	89930	1.01	0.05	=	65	0.07	-
250	PMS2-6	07p22.1	07-006.005320	8399	82061	0.96	0.05	=	51	0.08	-
436	PMS2-6	07p22.1	07-006.005380	8885	96647	0.99	0.05	=	65	0.02	-
454	PMS2-5	07p22.1	07-006.008630	8770	97621	1	0.05	=	65	0.01	-
232	PMS2-5	07p22.1	07-006.008720	9636	96464	0.95	0.05	=	52	0.03	-
418	PMS2-4	07p22.1	07-006.009750	9009	101574	0.99	0.04	=	85	0.04	-
310	PMS2-3	07p22.1	07-006.010020	8798	90829	0.98	0.04	=	65	0.04	-
390	PMS2-3	07p22.1	07-006.010320	6844	76131	0.99	0.04	=	63	0.00	-
184	PMS2-2	07p22.1	07-006.012070	9715	96831	0.97	0.04	=	57	0.01	-
400	PMS2-2	07p22.1	07-006.012160	7817	83565	0.99	0.04	=	57	0.00	-
146	PMS2-1	07p22.1	07-006.015220	10088	97727	1.01	0.05	=	64	0.00	-
338	PMS2-up	07p22.1	07-006.015460	9227	99131	1.01	0.05	=	78	0.00	-
292	Reference	02p24.1	02-021.113540	9570	92600	0.99	0.04	=	52	0.10	-
328	Reference	03q24	03-148.613920	8283	86546	0.93	0.06	=	60	0.06	-
268	Reference	03q29	03-196.509270	8878	88706	0.97	0.04	=	49	0.03	-
128	Reference	05q31.1	05-132.037630	9301	86718	1.04	0.04	=	47	0.01	-
154	Reference	06p21.31	06-035.966720	8426	77491	1.02	0.04	=	40	0.02	-
177	Reference	07p13	07-045.044390	9182	89498	1.01	0.04	=	54	0.02	-
472	Reference	08q12.2	08-061.928330	8748	100876	0.99	0.04	=	58	0.07	-
483	Reference	10p12.31	10-021.226290	8555	97347	0.95	0.05	=	69	0.01	-
427	Reference	11p13	11-031.772220	8734	96607	1	0.05	=	81	0.00	-
226	Reference	11p13	11-032.081900	9163	88957	1	0.04	=	55	0.07	-
196	Reference	14q24.2	14-072.684510	10014	93014	1.03	0.04	=	60	0.02	-
373	Reference	17q11.2	17-026.707650	9022	97877	0.99	0.05	=	66	0.01	-
160	Reference	17q23.2	17-057.148130	9143	89187	1.01	0.04	=	51	0.04	-
Median value all probe values:				8832	90571	0.99	0.04		57	0.02	