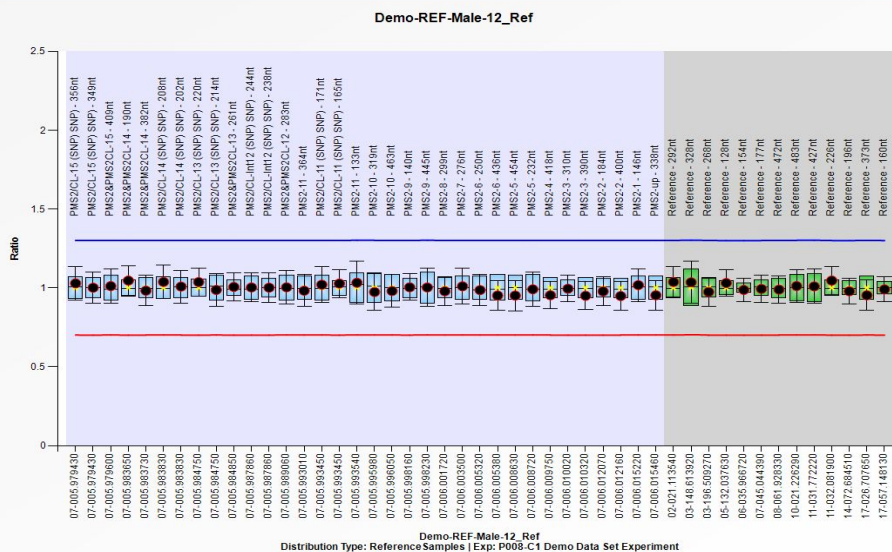
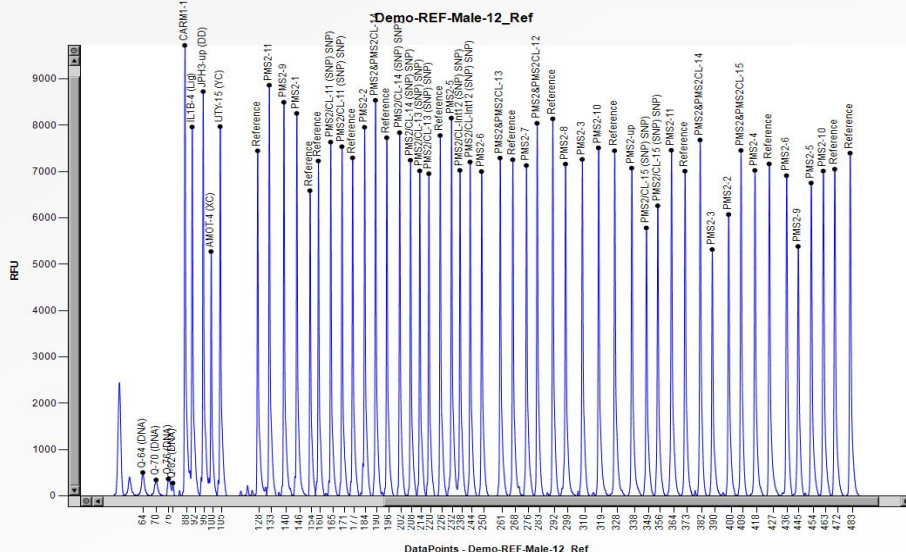


MLPA probe mix:	P008-PMS2	Nr of test probes:	47/47	FRSS:	OK 100%
Lot number:	C1-0523	Nr of ref probes	13/13	FMRS:	OK 100%
Sheet date:	4/8/2026 1:37:23 PM	DNA concentration:	OK	PSLP:	OK 2%
Control fragments:	CF-003-[brown] QDX2 (A2-0	DNA denaturation	OK	RSQ:	OK
Analysis method:	Block SSC: Off	Expected gender:	Male	RPQ:	OK
Used metric:	Peak height	Residual primer %	OK 12%	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	6262	71683	1.03	0.05	=	64	0.06	SNP
349	PMS2/CL-15 (S	07p22.1	07-005.979430	5783	65737	1	0.05	=	61	0.02	SNP
409	PMS2&PMS2CL-15	07p22.1	07-005.979600	7456	86278	1.01	0.05	=	56	0.09	-
190	PMS2&PMS2CL-14	07p22.1	07-005.983650	8540	83268	1.05	0.05	=	50	0.03	-
382	PMS2&PMS2CL-14	07p22.1	07-005.983730	7681	86560	0.98	0.05	=	55	0.12	-
208	PMS2/CL-14 (S	07p22.1	07-005.983830	7244	69348	1.04	0.05	=	50	0.00	SNP
202	PMS2/CL-14 (S	07p22.1	07-005.983830	7839	73289	1.01	0.05	=	51	0.00	SNP
220	PMS2/CL-13 (S	07p22.1	07-005.984750	6955	66611	1.04	0.04	=	47	0.05	SNP
214	PMS2/CL-13 (S	07p22.1	07-005.984750	7016	68062	0.99	0.05	=	49	0.07	SNP
261	PMS2&PMS2CL-13	07p22.1	07-005.984850	7293	76739	1.01	0.04	=	63	0.03	-
244	PMS2/CL-Int12	07p22.1	07-005.987860	7207	73065	1	0.05	=	51	0.03	SNP
238	PMS2/CL-Int12	07p22.1	07-005.987860	7026	71441	1	0.05	=	55	0.00	SNP
283	PMS2&PMS2CL-12	07p22.1	07-005.989060	8043	83749	1	0.05	=	54	0.09	-
364	PMS2-11	07p22.1	07-005.993010	7460	88834	0.98	0.05	=	73	0.03	-
171	PMS2/CL-11 (S	07p22.1	07-005.993450	7537	72618	1.02	0.06	=	59	0.04	SNP
165	PMS2/CL-11 (S	07p22.1	07-005.993450	7636	72377	1.03	0.04	=	57	0.05	SNP
133	PMS2-11	07p22.1	07-005.993540	8863	85705	1.03	0.07	=	57	0.00	-
319	PMS2-10	07p22.1	07-005.995980	7512	81853	0.97	0.06	=	68	0.00	-
463	PMS2-10	07p22.1	07-005.996050	7011	84384	0.98	0.05	=	60	0.11	-
140	PMS2-9	07p22.1	07-005.998160	8496	82013	1	0.04	=	62	0.05	-
445	PMS2-9	07p22.1	07-005.998230	5384	68063	1	0.06	=	55	0.15	-
299	PMS2-8	07p22.1	07-006.001720	7162	80703	0.98	0.05	=	84	0.02	-
276	PMS2-7	07p22.1	07-006.003500	7132	77217	1.01	0.06	=	60	0.06	-
250	PMS2-6	07p22.1	07-006.005320	7000	71127	0.99	0.05	=	59	0.05	-
436	PMS2-6	07p22.1	07-006.005380	6911	81933	0.95	0.05	=	64	0.16	-
454	PMS2-5	07p22.1	07-006.008630	6753	80544	0.95	0.05	=	65	0.10	-
232	PMS2-5	07p22.1	07-006.008720	8154	86033	0.99	0.05	=	52	0.04	-
418	PMS2-4	07p22.1	07-006.009750	7025	84301	0.95	0.04	=	83	0.06	-
310	PMS2-3	07p22.1	07-006.010020	7263	78352	1	0.04	=	58	0.05	-
390	PMS2-3	07p22.1	07-006.010320	5317	61359	0.95	0.04	=	70	0.11	-
184	PMS2-2	07p22.1	07-006.012070	7956	81701	0.98	0.05	=	60	0.04	-
400	PMS2-2	07p22.1	07-006.012160	6073	70245	0.95	0.05	=	61	0.07	-
146	PMS2-1	07p22.1	07-006.015220	8257	81295	1.02	0.05	=	73	0.04	-
338	PMS2-up	07p22.1	07-006.015460	7075	80800	0.95	0.05	=	80	0.00	-
292	Reference	02p24.1	02-021.113540	8139	85064	1.04	0.05	=	55	0.05	-
328	Reference	03q24	03-148.613920	7450	84823	1.04	0.07	=	89	0.00	-
268	Reference	03q29	03-196.509270	7256	77335	0.97	0.05	=	74	0.03	-
128	Reference	05q31.1	05-132.037630	7448	71389	1.03	0.04	=	52	0.05	-
154	Reference	06p21.31	06-035.966720	6588	61291	0.99	0.04	=	39	0.01	-
177	Reference	07p13	07-045.044390	7296	72767	0.99	0.04	=	52	0.12	-
472	Reference	08q12.2	08-061.928330	7052	86238	0.99	0.04	=	65	0.06	-
483	Reference	10p12.31	10-021.226290	7397	90370	1.01	0.05	=	74	0.05	-
427	Reference	11p13	11-031.772220	7166	85053	1.01	0.05	=	77	0.12	-
226	Reference	11p13	11-032.081900	7778	76894	1.05	0.04	=	52	0.00	-
196	Reference	14q24.2	14-072.684510	7731	76447	0.98	0.04	=	56	0.04	-
373	Reference	17q11.2	17-026.707650	7008	80926	0.95	0.05	=	67	0.04	-
160	Reference	17q23.2	17-057.148130	7229	68362	0.99	0.04	=	49	0.06	-
Median value all probe values:				7256	78352	1	0.05		59	0.05	