



Sample report: 2 MCH P198 P16048 1-5

Sample type: Sample | Project: P198 | Experiment: 20160210_rozc1-5 | Dye: 6-FAM | Performed by: Admin
 Machine: ABI-3130 | Report date: 2016-02-11 | Run date: 2016-02-11 | Software Version: v.140721.1958 | Normal range: 0,7 - 1,3

Authorization
Date

MLPA probe mix: P198-FH
 Lot number: A3-1014
 Sheet date: 2015-12-21 14:48:24
 Control fragments: CF-003-[brown] QDX2 (A2-0)
 Analysis method: Block SSC: On
 Used metric: Peak height

Nr of test probes: 19/19
 Nr of ref probes: 9/9
 DNA concentration: OK
 DNA denaturation: OK
 Expected gender: Female
 Residual primer %: OK 8%

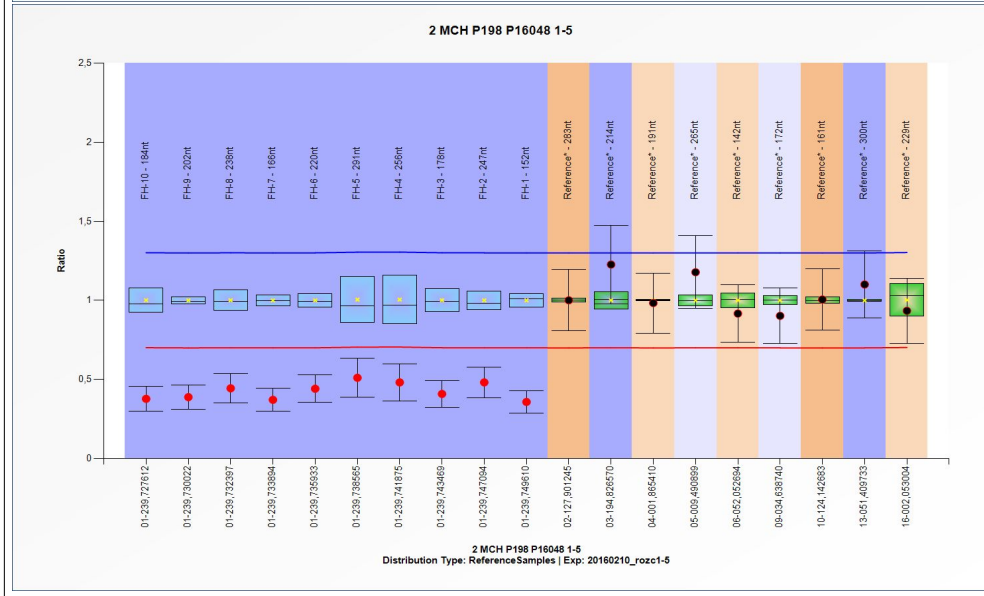
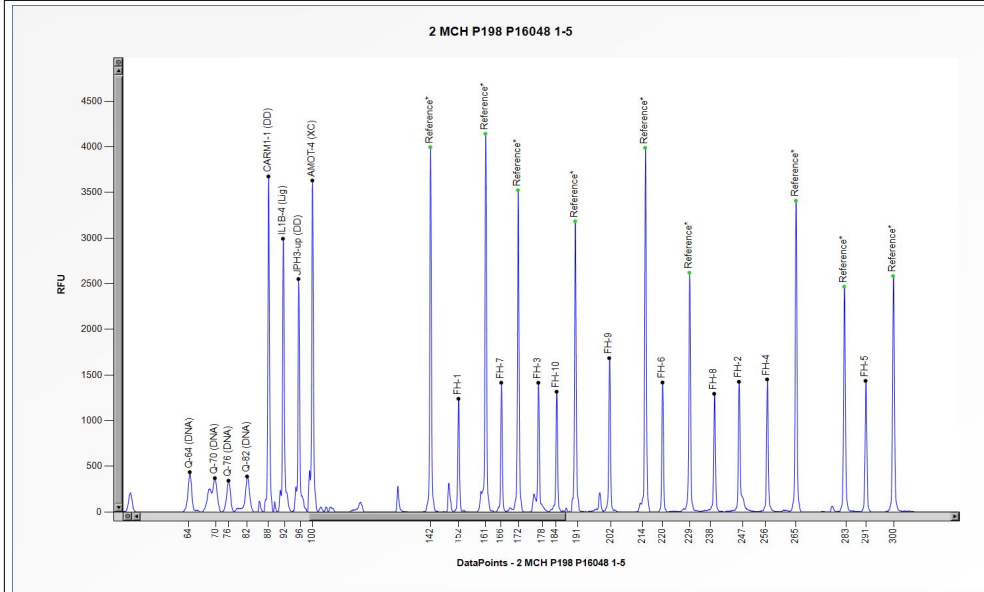
FRSS: OK 100%
 FRMS: OK 100%
 PSLP: Warning -16%
 RSO: OK
 RPQ: Warning
 CAS: OK? 85%

Reference Samples: 2 MCH P198 PC1 1-5 | 2 MCH P198 PC2 1-5 | 2 MCH P198 PC3 1-5

D [nt]	Gene-Exon	Chr.band	hg18 loc.	Height	Area	Ratio ^H	Stdev	[REF]	[Sam]	Width	d[nt]
184	FH-10	01q43	01-239,727612	1317	8785	0,38	0,04	<<*	?	39	0,0
202	FH-9	01q43	01-239,730022	1685	11050	0,39	0,04	<<*	?	39	0,1
238	FH-8	01q43	01-239,732397	1294	9324	0,44	0,05	<<*	<*	56	0,1
166	FH-7	01q43	01-239,733894	1416	8553	0,37	0,04	<<*	?	29	0,0
220	FH-6	01q43	01-239,735933	1418	8803	0,44	0,04	<<*	<*	30	0,0
291	FH-5	01q43	01-239,738565	1436	10071	0,51	0,06	<<*	<*	31	0,1
256	FH-4	01q43	01-239,741875	1452	10032	0,48	0,06	<<*	<*	60	0,1
178	FH-3	01q43	01-239,743469	1415	9210	0,41	0,04	<<*	?	36	0,0
247	FH-2	01q43	01-239,747094	1426	12059	0,48	0,05	<<*	<*	85	0,0
152	FH-1	01q43	01-239,749610	1240	7630	0,36	0,04	<<*	?	22	-0,1
283	Reference*	02q14.3	02-127,901245	2468	17524	1	0,1	=	=	43	0,1
214	Reference*	03q29	03-194,826570	3990	26271	1,23	0,12	=	=	46	0,1
191	Reference*	04p16.3	04-001,865410	3184	20822	0,98	0,09	=	=	42	0,0
265	Reference*	05p15.31	05-009,490899	3409	22963	1,18	0,12	=	=	34	0,0
142	Reference*	06p12.2	06-052,052694	3997	24648	0,92	0,09	=	=	51	0,0
172	Reference*	09p13.3	09-034,638740	3525	22197	0,9	0,09	=	=	34	0,0
161	Reference*	10q26.13	10-124,142683	4144	28052	1,01	0,1	=	=	55	0,0
300	Reference*	13q14.3	13-051,409733	2585	19400	1,1	0,11	=	=	56	0,1
229	Reference*	16p13.3	16-002,053004	2621	17857	0,93	0,1	=	=	59	0,1

Median value all probe values:

1685	12059	0,51*	0,06	42	0,03
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Normal range: 0,7 - 1,3 Essential information on the use of this product is present in the product description which is available on <http://www.mlpa.com>. For questions, <mailto:info@mlpa.com>.
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