

Genome: hg19
 Aberration Algorithm: ADM-1
 Threshold: 6.0
 Fuzzy Zero: ON
 GC Correction: ON
 Window Size: 2Kb
 Centralization (legacy): ON
 Centralization (legacy) Bin Size: 10
 Centralization (legacy) Threshold: 6.0
 Diploid Peak Centralization: ON
 Manually Reassign Peaks: OFF
 Combine Replicates (Intra Array): ON
 Combine Replicates (Inter Array): OFF
 Array Level Filter: NONE
 Aberration Filter: (Minimum Number of Probes for Amplification >= 8 AND Minimum Size (Kb) of Region >= 1000)
 Feature Level Filter: glsSaturated = true OR rlsSaturated = true OR glsFeatNonUnifOL = true OR rlsFeatNonUnifOL = true
 Design Level Filter: Homology = 0
 Genomic Boundary: OFF
 Show Flat Intervals: OFF

Aberration	Chr	Cytoband	Start	Stop	#Probes	Amplification	Deletion	pval
IPSC-20giu24-03_258633223258_S01_CGH_1100_Jul11_1_3								
1	chr1	p36.32	2567773	3149672	39	0	-0.3563	5.67E-20
2	chr1	q21.2	1.49E+08	1.49E+08	13	0	-0.59713	1.25E-18
3	chr2	p11.2	89143858	89312590	11	1.053887	0	8.87E-51
4	chr6	p21.32	32565005	32688989	10	0	-0.52186	1.85E-10
5	chr8	p11.22	39237438	39362887	11	5.30271	0	0.000
6	chr11	q11	55377910	55450788	8	0	-0.88965	7.65E-26
7	chr14	q32.33	1.06E+08	1.07E+08	17	3.686258	0	0.000
8	chr14	q32.33	1.07E+08	1.07E+08	27	0.799755	0	3.90E-72
9	chr15	q11.1 - q11	20564608	22558756	33	0	-0.64928	6.07E-58
10	chr17	q21.2	39093536	39594086	28	0	-0.2951	1.62E-13
11	chr21	q11.2	14678356	15372526	15	0	-0.44202	2.39E-11
12	chr22	q11.22	23056562	23245888	27	4.065959	0	0.000
13	chrX	p22.33	1019030	1458111	28	0.454671	0	1.56E-23
14	chrX	p11.23	47871771	47960608	8	0.630079	0	9.71E-16

Number of calls per sample:
 IPSC-20giu24-03_258633223258_S01_CGH_1100_Jul11_1_3 = 18

on for Amplification ≥ 0.0 AND Minimum Avg. Absolute Log Ratio for Amplification ≥ 0.00) OR (Minimum Avg. Absolute Log Ratio for Amplification ≥ 0.00 AND Minimum Avg. Absolute Log Ratio for Deletion ≥ 0.00) AND atNonUnifOL = true OR LogRatio = 0

Gene Nam Hs hg19 (Hs hg19 rHs hg19 PAR 20120403

ACTRT2, F CNV 2982 hsa-mir-4251

CNV_34420, CNV_3313, CNV_8328, CNV_2326, CNV_8924, CNV_8329, CNV_48117, CNV_0525, CNV_35869, CNV_35870, CNV_35871, CNV_35872, CNV_34427, CNV_35873

HLA-DQA1 CNV_3603, CNV_4493, CNV_64476, CNV_7567, CNV_37840, CNV_69421, CNV_31280, (ADAM5P, CNV_2749, CNV_36815, CNV_105114, CNV_112161, CNV_59473, CNV_36816, CNV_10OR4P4, OF CNV_8690, CNV_0308, CNV_3852, CNV_48051, CNV_2926, CNV_53140, CNV_37484, CKIAA0125, CNV_23307, CNV_10460, CNV_7026, CNV_113813, CNV_35212, CNV_35213, CNV_352LOC10013 CNV_23307, CNV_10460, CNV_7026, CNV_113813, CNV_35212, CNV_35213, CNV_352GOLGA6L CNV_3951 hsa-mir-3118-4, hsa-mir-3118-6, hsa-mir-1268

KRT23, KFCNV_72924, CNV_43948, CNV_72925, CNV_102455, CNV_67193, CNV_38355, CNV_77!
POTED, C; CNV_8890 hsa-mir-3156-3, hsa-mir-3118-5

CNV 5356 hsa-mir-650

CRLF2, CS CNV_3441 hsa-mir-36 PAR-chrX.p

ZNF630 CNV_4149, CNV_59879, CNV_3250, CNV_37025, CNV_0822, CNV_2276, CNV_34577, C

num Number of Probes for Deletion >= 8 AND Minimum Size (Kb) of Region for Deletion >= 0.0 AND Mi

IV_48118, CNV_48119, CNV_48120, CNV_6785, CNV_64896, CNV_64895, CNV_84553, CNV_30433,
3, CNV_22905, CNV_107994, CNV_35874, CNV_97914, CNV_104583, CNV_44037, CNV_78633, CNV
CNV_4767, CNV_111517, CNV_81340, CNV_93642, CNV_103651, CNV_103652, CNV_69420, CNV_6
5798, CNV_65114, CNV_106294, CNV_37462, CNV_7683, CNV_82574, CNV_95380, CNV_106595, C
NV_0419, CNV_34977, CNV_9663, CNV_29606, CNV_48826, CNV_48827, CNV_48828, CNV_53507,
14, CNV_4850, CNV_35215, CNV_113843, CNV_107992, CNV_0770, CNV_2176, CNV_0441, CNV_3
14, CNV_107992, CNV_0317, CNV_35218, CNV_35219, CNV_35220, CNV_22954, CNV_72040, CNV_
593, CNV_88455, CNV_88456, CNV_77594, CNV_88457, CNV_77595, CNV_9286, CNV_72927, CNV_

NV_8304, CNV_34416, CNV_83493, CNV_96641, CNV_67973, CNV_67975, CNV_7789, CNV_37576,

[illegible]

[illegible]

43, CNV_9861, CNV_53002, CNV_53003, CNV_53004, CNV_83491, CNV_8912, CNV_8305, CNV_8306

√_48131, CNV_1327, CNV_1331, CNV_1328, CNV_1329, CNV_1330, CNV_4254, CNV_1332, CNV_10
√V_78640, CNV_74111, CNV_74120, CNV_115389, CNV_115390, CNV_115391, CNV_103346, CNV_
√V_69413, CNV_103655, CNV_38775, CNV_111502, CNV_111541, CNV_93643, CNV_59329, CNV_6
4, CNV_48835, CNV_48836, CNV_53101, CNV_38079, CNV_9188, CNV_48053, CNV_4743, CNV_48
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414, CNV_72007, CNV_35225, CNV_35226, CNV_4856, CNV_113817, CNV_43896, CNV_72014, CN'
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_74107, CNV_74123, CNV_104585, CNV_115398, CNV_7286, CNV_37226, CNV_74122, CNV_37227,
i9462, CNV_107890, CNV_32794, CNV_69423, CNV_81338, CNV_81341, CNV_111498, CNV_111518

837, CNV_53573, CNV_48838, CNV_48839, CNV_113029, CNV_53178, CNV_75645, CNV_65956, CI
CNV_113844, CNV_72016, CNV_72050, CNV_72013, CNV_113832, CNV_113845, CNV_113846, CNV
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3791, CNV_1590, CNV_78641, CNV_59010, CNV_74121, CNV_23274, CNV_115401, CNV_78642, CNV_93650, CNV_69473, CNV_93641, CNV_103656, CNV_111501, CNV_111525, CNV_59330, CNV_2328

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CNV_7159, CNV_38806, CNV_104123, CNV_102458, CNV_30829, CNV_109535, CNV_0205, CNV_31

4113, CNV_89784, CNV_74104, CNV_74124, CNV_68082, CNV_74115, CNV_34620, CNV_115397, CNV_93651, CNV_93638, CNV_69409, CNV_81339, CNV_69428, CNV_69437, CNV_69463, CNV_6250

6762, CNV_76763, CNV_113850, CNV_0772, CNV_2177, CNV_108091, CNV_76764, CNV_76767, CNV_35231, CNV_35232, CNV_113859, CNV_72034, CNV_72079, CNV_5771, CNV_113860, CNV_7:

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CNV_10516, CNV_49858, CNV_37949, CNV_1964, CNV_9773, CNV_114403, CNV_67199, CNV_7760

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8, CNV_77609, CNV_88471, CNV