ISGC Parentage SNP	Chr	Mb Pos	Allele	MAF	Filter 1 Re-seq	Filter 2 Fluidigm	Filter 3 SQ_AgR	Filter 4 SQ_CLI	5k Chip	Filter problem
DU290101_408	1	7.8	A	0.337						3
DU518561_359	1	14.2	G	0.381						2
DU351298_316	1	69.6	A	0.445						
DU232924_365	1	95.8	G	0.250						
DU271929_382	1	97.5	A	0.483						4
DU502334_443	2	19.1	A	0.437						
DU469454_586	2	26.2	G	0.394						1
DU425907_184	2	50.1	G	0.358						
DU501115_497	2	62.8	A	0.239						
DU492516_411	2	63.4	T	0.478						
DU470875_383	2	91.5	G	0.357						
250506CS39010123000	2	100.9	G	0.345						
DU191879_495	2	157.6	A	0.335						
DU480434_533	2	192.2	A	0.480						
DU260201_585	2	226.7	A	0.422						
DU503161_123	2	237.2	A	0.352						
DU425259_620	3	21.4	A	0.461						4
DU231007_156	3	59.0	G	0.463						2
DU225323_218	3	91.0	A	0.467						
DU260081_579	3	108.8	A	0.383						
DU394537_289	3	181.6	G	0.371						
CL635241_413	3	181.9	A	0.455						
DU408817_431	3	205.0	A	0.343						
DU202116_405	4	58.2	A	0.444						
DU460511_423	4	61.1	G	0.443						
DU305004_417	4	70.1	A	0.270						
DU369175_467	4	73.0	G	0.375						
DU446213_412	5	12.5	A	0.394						
DU444709_372	5	56.0	A	0.489						
DU453259_440	5	64.8	G	0.346						
DU194639_560	6	56.7	G	0.442						
CZ925803_293	6	100.8	A	0.443						
DU337465_337	6	106.0	A	0.338						
CL635944_160	6	115.0	A	0.490						4
DU467751_524	7	10.6	A	0.429						
DU499587_509	7	74.0	A	0.325						
CZ920950_468	7	74.8	A	0.456						
DU530067_219	7	100.0	G	0.327						
DU213735_493	8	6.6	A	0.437						
DU411204_551	8	13.8	A	0.361						
DU189970_325	9	86.6	C	0.374						
DU471913_499	9	91.1	G	0.490						
DU364754_308	9	93.9	A	0.397						
DU372582_268	9	94.4	G	0.247						2
DU468275_284	10	33.1	A	0.352						
DU310747_445	10	38.2	G	0.470						
DU269694_582	11	1.9	A	0.473						
DU433863_261	11	15.5	A	0.419						
DU417675_79	11	19.6	A	0.344						
DU508448_227	11	25.3	A	0.485						4
DU326572_241	11	59.5	A	0.446						
DU314655_578	12	26.7	A	0.365						
DU310703_497	12	75.3	A	0.492						1
DU275428_276	13	10.9	A	0.460						
DU435573_466	13	30.1	A	0.449						
DU411403_398	13	41.3	G	0.427						
DU462008_263	14	44.6	A	0.330						
DU223894_556	14	57.5	G	0.449						1
DU381045_479	14	60.7	A	0.403						
DU464373_638	15	2.3	A	0.467						
DU426312_454	15	44.4	G	0.375						
DU301502_402	15	73.7	G	0.441						

DU241306_191	15	78.6	G	0.279							
DU324670_456	17	10.2	A	0.400							
DU206327_107	17	14.4	A	0.499							
DU378819_632	17	22.3	A	0.475							3
DU511222_139	17	27.4	A	0.351							
DU300156_445	17	38.0	G	0.456							
DU463532_137	17	56.0	A	0.443							
DU492379_209	18	3.9	A	0.385							
DU488903_267	18	21.4	G	0.334							
DU325612_517	18	25.4	A	0.433							1
DU440765_491	18	60.5	A	0.474							3
DU345394_399	18	61.1	A	0.450							4
DU264531_279	19	0.6	A	0.388							
DU258053_237	19	57.1	A	0.400							
DU411432_523	19	57.2	C	0.406							
DU183112_480	20	31.1	A	0.453							
DU442373_141	20	48.4	A	0.342							
DU380983_440	21	28.3	G	0.451							
DU383863_376	21	38.2	G	0.443							1
DU196132_525	21	42.7	G	0.388							
DU413316_575	22	13.1	A	0.419							
DU302760_528	23	11.6	G	0.494							
DU313102_671	23	17.3	G	0.484							
CZ920359_258	24	3.2	G	0.382							
DU455254_479	25	0.1	G	0.453							
DU512685_259	25	1.2	G	0.495							
oY1	Y	0.0	G	0.320							