

SUPPLEMENTARY TABLE 3: SNP genotype call rates

SNP	Missing data cases	Missing data controls	p HWE
rs2484676	0.08968	NA	0.3004
rs11205760	0.1168	NA	0.8304
rs4506508	0.1083	NA	0.3011
rs2814036	0.06176	NA	0.1105
rs12035082	0.05161	0.08696	0.903
rs10801047	0.1252	0.02767	0.03348
rs17419032	0.1261	NA	1
rs363617	0.08799	NA	0.005125
rs2322659	0.1193	NA	0.5904
rs9858542	0.1328	0.07065	0.03211
rs9870678	0.1269	NA	0.6901
rs1462651	0.1100	NA	0.6761
rs9993022	0.2132	NA	0.4204
rs17234657	0.05668	0.08449	0.009521
rs9292777	0.07783	0.1052	0.2744
rs10077785	0.04653	0.126	0.116
rs13361189	0.1159	0.1196	0.5228
rs4958847	0.08545	0.02668	1
rs1363670	0.1125	NA	0.5607
rs6887695	0.08883	0.1428	0.455
rs17309827	0.1582	NA	0.8854
rs12529198	0.1176	NA	0.8169
rs6908425	0.08968	NA	0.4938
rs6927210	0.07783	NA	0.4677
rs946227	0.1151	NA	0.6353
rs1558043	0.08037	NA	0.1331
rs12704036	0.1041	NA	0.4183
rs4871612	0.05668	NA	0.09334
rs3936503	0.08037	NA	0.6359
rs10761659	0.0423	NA	0.07061
rs7081330	0.1015	NA	0.1086
rs10883365	0.1311	0.1259	0.3358
rs1931047	0.0643	NA	0.4456
rs916977	0.09052	NA	0.04415
rs9895062	0.1024	NA	0.1845
rs2542151	0.1049	0.1215	0.5582
rs2836754	0.0753	0.1146	0.7498

Fraction of replication cases and controls (where applicable) with null genotype calls and HWE p value for replication cases for each of 37 genotyped SNPs.