Supplementary Table: Schizophrenia-associated loci at a genome-wide level of significance ($P<5\times10^{-8}$) in the Li et al. (2017) schizophrenia GWAS dataset. Chr: chromosome, A1: minor allele, A2: major allele, OR: odds ratio, SE=standard error.

SNP	Chr.	Position	A1/A2	OR (SE)	*P* value
rs172531	1	8,495,590	A/G	0.931 (0.011)	5.05E-11
rs4949526	1	30,432,219	T/C	1.065 (0.010)	2.81E-10
rs2970610	1	44,097,530	T/C	1.072 (0.010)	1.24E-12
rs11210195	1	73,749,283	T/C	0.943 (0.010)	1.76E-09
rs1198589	1	98,550,411	T/C	0.907 (0.013)	1.19E-13
rs12129037	1	150,492,114	T/C	1.096 (0.016)	4.51E-09
rs6670165	1	177,280,121	T/C	1.073 (0.012)	3.37E-09
rs35978510	1	243,776,117	T/C	0.941 (0.010)	6.50E-10
rs12623170	2	28,020,157	T/C	0.943 (0.010)	2.52E-08
rs75575209	2	58,138,192	A/T	0.902 (0.018)	1.24E-08
rs1518395	2	58,208,074	A/G	0.862 (0.020)	3.78E-13
rs999494	2	73,157,395	T/C	0.924 (0.012)	2.40E-10
rs62152284	2	104,984,387	T/C	1.065 (0.011)	5.86E-09
rs6430491	2	134,840,967	A/G	0.936 (0.011)	9.55E-10
rs2381759	2	146,428,031	A/T	0.929 (0.013)	8.63E-09
rs7597593	2	185,533,580	T/C	1.066 (0.010)	9.30E-11
rs12619354	2	194,002,782	T/C	1.056 (0.010)	3.81E-08
rs895526	2	200,162,425	T/C	0.936 (0.011)	9.27E-10
rs796364	2	200,716,119	A/C	0.912 (0.011)	2.44E-16
rs17630293	2	201,146,776	T/G	1.124 (0.017)	3.32E-12
rs78681500	2	201,187,316	T/C	0.751 (0.045)	3.08E-10
rs11685299	2	225,391,296	A/C	0.934 (0.010)	4.70E-12
rs778371	2	233,743,109	A/G	0.923 (0.011)	2.17E-13
rs11708578	3	2,515,894	A/G	0.925 (0.012)	7.28E-11
rs4330281	3	17,859,366	T/C	0.942 (0.011)	3.04E-08
rs10510653	3	32,058,559	A/T	1.074 (0.013)	2.54E-08
rs3732386	3	36,871,993	T/C	1.075 (0.011)	2.88E-11
rs2073499	3	50,374,293	A/G	0.899 (0.019)	2.61E-08
rs2577831	3	52,628,056	A/C	0.936 (0.009)	4.53E-13
rs832190	3	63,842,629	T/C	0.948 (0.010)	3.93E-08
rs66691851	3	136,154,828	T/C	0.926 (0.010)	4.17E-15
rs6804239	3	161,780,488	T/C	0.943 (0.009)	8.41E-10
rs9859557	3	180,830,910	A/T	1.094 (0.012)	6.04E-14
rs9841616	3	181,167,585	A/T	0.928 (0.013)	5.35E-09
rs215411	4	23,423,603	A/T	1.069 (0.011)	6.37E-10
rs11722779	4	103,827,488	T/G	0.951 (0.009)	3.40E-08
rs7438	4	170,642,246	A/G	1.066 (0.010)	7.46E-11
rs6846161	4	176,866,459	A/G	1.067 (0.011)	9.96E-10

rs9292918	5	45,301,035	T/G	0.932 (0.011)	4.47E-11
rs10940346	5	49,806,042	A/G	0.949 (0.009)	1.11E-08
rs2247870	5	90,151,589	A/G	1.052 (0.009)	2.54E-08
rs4388249	5	109,036,066	T/C	1.066 (0.011)	8.38E-09
rs2764766	5	127,213,625	C/G	1.055 (0.009)	1.94E-08
rs13164092	5	137,840,860	A/G	0.943 (0.010)	9.79E-10
rs7733403	5	140,154,215	A/G	1.052 (0.009)	2.06E-08
rs113568682	5	152,299,754	A/T	1.064 (0.011)	3.77E-08
rs2910032	5	152,540,354	T/C	0.943 (0.010)	1.04E-09
rs11740474	5	153,680,747	A/T	0.941 (0.011)	1.81E-08
rs115329265	6	28,712,247	A/G	1.210 (0.015)	5.02E-36
rs111782145	6	30,873,508	C/T	0.813 (0.037)	1.80E-08
rs6903570	6	64,866,857	T/C	0.947 (0.010)	2.70E-08
rs217331	6	84,344,689	C/G	0.934 (0.009)	1.63E-13
rs160593	6	105,466,332	A/G	0.947 (0.009)	7.69E-09
rs9398171	6	108,983,527	T/C	0.935 (0.010)	1.85E-11
rs7757969	6	112,132,032	C/T	1.110 (0.019)	4.82E-08
rs4479915	6	165,075,601	G/C	0.876 (0.023)	4.82E-09
rs12668848	7	2,020,995	A/G	0.923 (0.010)	3.97E-17
rs112509803	7	24,735,004	C/G	1.090 (0.016)	4.80E-08
rs2944823	7	71,795,470	A/G	1.059 (0.010)	3.21E-08
rs323167	7	78,336,677	T/C	0.946 (0.010)	4.47E-08
rs12704290	7	86,427,626	A/G	0.908 (0.015)	2.61E-10
rs2057884	7	104,930,250	T/C	1.069 (0.010)	2.02E-12
rs214467	7	110,858,268	A/G	1.073 (0.010)	7.68E-13
rs11534004	7	113,467,444	G/A	0.890 (0.021)	1.71E-08
rs3735025	7	137,074,844	T/C	1.068 (0.010)	3.60E-12
rs13261217	8	4,183,057	A/G	0.934 (0.011)	2.12E-09
rs2980436	8	8,092,025	A/G	1.057 (0.010)	2.31E-08
rs12543645	8	10,246,325	C/G	1.058 (0.010)	3.32E-08
rs17687067	8	17,036,201	A/C	0.925 (0.011)	3.39E-12
rs73219805	8	26,272,768	A/G	1.078 (0.011)	1.94E-11
rs112537273	8	38,248,306	T/C	1.077 (0.011)	1.06E-11
rs867743	8	60,694,647	A/G	0.945 (0.010)	4.97E-09
rs7010876	8	89,264,751	A/T	0.939 (0.011)	1.48E-08
rs16880831	8	111,561,505	T/G	0.930 (0.012)	4.34E-10
rs13266463	8	143,403,693	A/G	0.947 (0.010)	2.25E-08
rs2767713	9	84,799,537	A/C	0.941 (0.011)	2.02E-08
rs7893279	10	18,745,105	T/G	1.117 (0.015)	3.28E-13
rs111364339	10	64,857,872	T/C	0.906 (0.017)	5.37E-09
rs11191424	10	104,625,886	A/G	0.917 (0.009)	4.22E-20
rs10786736	10	104,849,116	C/G	0.897 (0.013)	4.39E-17
rs17234749	11	24,381,898	A/G	0.943 (0.010)	5.39E-09

rs61882743	11	46,548,754	C/G	0.914 (0.013)	6.99E-12
rs9420	11	57,510,294	A/G	1.061 (0.010)	8.52E-09
rs12421382	11	109,378,071	T/C	0.945 (0.010)	1.51E-08
rs2514218	11	113,392,994	T/C	0.928 (0.011)	5.01E-11
rs77502336	11	123,394,636	C/G	1.063 (0.010)	1.35E-09
rs55661361	11	124,613,957	A/G	0.932 (0.010)	1.16E-12
rs10894308	11	130,891,895	A/G	0.947 (0.010)	2.09E-08
rs75059851	11	133,822,569	A/G	1.098 (0.013)	8.77E-13
rs619091	11	133,884,137	A/T	0.947 (0.009)	5.85E-09
rs2159100	12	2,346,393	T/C	1.102 (0.011)	3.45E-19
rs758117	12	2,513,309	T/C	0.941 (0.011)	1.68E-08
rs302321	12	29,928,388	A/C	0.937 (0.010)	2.07E-11
rs4240748	12	92,246,786	C/G	0.948 (0.010)	4.55E-08
rs4766428	12	110,723,245	T/C	1.065 (0.011)	2.60E-09
rs28607014	12	117,708,611	T/C	0.946 (0.010)	1.75E-08
rs61041384	12	123,644,043	T/C	0.891 (0.016)	5.50E-13
rs1615350	12	123,650,335	T/C	0.916 (0.010)	5.53E-18
rs10148671	14	29,469,373	T/C	0.947 (0.010)	4.46E-08
rs1191551	14	30,000,405	T/G	1.074 (0.013)	2.28E-08
rs2383377	14	33,257,914	A/G	1.081 (0.014)	2.36E-08
rs35607894	14	72,408,638	A/G	1.094 (0.014)	5.37E-11
rs8012642	14	84,669,481	C/G	1.055 (0.010)	4.66E-08
rs2693698	14	99,719,219	A/G	0.941 (0.011)	2.20E-08
rs35225048	15	61,856,263	T/C	0.935 (0.009)	7.76E-13
rs3825845	15	78,910,258	T/C	0.932 (0.011)	2.47E-10
rs783540	15	83,254,708	A/G	0.950 (0.009)	3.05E-08
rs950169	15	84,706,461	T/C	0.923 (0.012)	5.17E-12
rs758129	15	89,900,887	A/G	0.945 (0.010)	2.87E-08
rs6500596	16	4,470,027	T/G	0.939 (0.011)	5.24E-09
rs9922678	16	9,946,319	A/G	1.065 (0.011)	3.36E-09
rs7405404	16	13,749,859	T/C	1.073 (0.012)	8.06E-10
rs3814881	16	30,000,901	A/G	0.936 (0.009)	8.41E-13
rs8058130	16	64,371,163	A/G	0.948 (0.010)	4.77E-08
rs56007784	17	1,290,950	C/G	1.059 (0.009)	1.16E-09
rs8082590	17	17,958,402	A/G	0.936 (0.011)	7.40E-10
rs72843506	17	19,946,287	A/T	1.070 (0.012)	3.73E-08
rs35065479	17	55,736,735	A/G	1.066 (0.011)	2.31E-08
rs715170	18	53,795,514	T/C	0.928 (0.011)	5.21E-12
rs56775891	18	77,575,613	T/C	1.058 (0.010)	1.85E-08
rs28735056	18	77,622,879	A/G	0.941 (0.010)	4.60E-10
rs2053079	19	30,987,423	A/G	0.938 (0.011)	4.64E-09
rs5757730	22	39,967,430	A/G	0.933 (0.011)	1.84E-10
rs9607782	22	41,587,556	A/T	1.090 (0.012)	1.01E-12