

Table S1. Thirty Most Significant Association Results From the Analysis of Nicotine Dependence in the Dutch Sample

Chr	Location (bp)	SNP	Gene	SNP Type	P-value
15	76161636	rs11631180	<i>TBC1D2B</i>	Promoter	4.264 x 10 ⁻⁶
20	38475285	rs3092155			4.371 x 10 ⁻⁶
7	154997865	rs6459707			8.485 x 10 ⁻⁶
7	154996933	rs6971850			1.145 x 10 ⁻⁵
5	141168499	rs17097654			1.269 x 10 ⁻⁵
5	39249453	rs696748	<i>FYB</i>	Intron	1.531 x 10 ⁻⁵
10	28774361	rs7085106			1.695 x 10 ⁻⁵
15	76136405	rs12148365	<i>TBC1D2B</i>	Intron	2.060 x 10 ⁻⁵
16	51113243	rs8046979			2.300 x 10 ⁻⁵
18	7509001	rs576192			3.056 x 10 ⁻⁵
6	106644483	rs6923608	<i>PRDM1</i>	Intron	3.204 x 10 ⁻⁵
18	7478970	rs567338			3.700 x 10 ⁻⁵
11	2743637	rs231872	<i>KCNQ1</i>	Intron	3.720 x 10 ⁻⁵
15	96426216	rs12148124			3.759 x 10 ⁻⁵
3	188453990	rs3105782	<i>MASP1</i>	Intron	3.850 x 10 ⁻⁵
15	76131977	rs4886554	<i>TBC1D2B</i>	Intron	3.869 x 10 ⁻⁵
18	4016587	rs1372631			4.131 x 10 ⁻⁵
5	158920307	rs17729321			4.171 x 10 ⁻⁵
18	7481478	rs533682			4.237 x 10 ⁻⁵
12	104628953	rs4590982			4.240 x 10 ⁻⁵
5	160228811	rs12188645			4.421 x 10 ⁻⁵
11	2707296	rs2283214	<i>KCNQ1</i>	Intron	4.527 x 10 ⁻⁵
20	1558939	rs3746722	<i>SIRPB2</i>	Coding exon	4.571 x 10 ⁻⁵
12	12037208	rs12582693			4.596 x 10 ⁻⁵
7	147310468	rs11771941	<i>CNTNAP2</i>	Intron	4.874 x 10 ⁻⁵
7	147294425	rs2178771	<i>CNTNAP2</i>	Intron	4.990 x 10 ⁻⁵
12	74699761	rs7297654	<i>PHLDA1</i>	Downstream	5.087 x 10 ⁻⁵
11	2707861	rs231910	<i>KCNQ1</i>	Intron	5.102 x 10 ⁻⁵
20	1560819	rs1535883	<i>SIRPB2</i>	Intron	5.386 x 10 ⁻⁵
8	87695579	rs2362267	<i>CNGB3</i>	Intron	5.752 x 10 ⁻⁵