

**Significant association of the *CHRNA3-CHRNA6* gene cluster with nicotine
dependence in the Chinese Han population**

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Supplementary Table 1. Detailed information of selected SNPs in the
CHRNA6-CHRNA6 gene cluster

Gene	dbSNP ID	Assay ID	Chr. Pos.	Location	Alleles (Major/minor)	MAF
<i>CHRNA6</i>	rs10958725	C__11868275_20	8: 42524584	5' near gene	G/T	0.26
	rs10958726	C__11868302_10	8: 42535909	5' near gene	T/G	0.21
	rs13273442	C__11868306_10	8: 42544017	5' near gene	G/A	0.21
	rs4736835	C__26072327_10	8: 42547033	5' near gene	C/T	0.21
	rs1955186	C__11855540_10	8: 42549491	5' near gene	G/C	NA
	rs6474413	C__28976622_10	8: 42551064	5' near gene	T/C	0.21
	rs7004381	C__26072326_10	8: 42551161	5' near gene	G/A	0.21
	rs4950	C__8238995_20	8: 42552633	5'UTR	T/C	0.21
	rs1530848	C__1346774_10	8: 42552908	Intron	A/C	0.20
	rs13280604	C__272767_10	8: 42559586	Intron	A/G	0.21
	rs6474414	C__272765_10	8: 42560336	Intron	C/A	0.21
	rs6474415	C__29352729_10	8: 42562938	Intron	A/G	0.21
	rs4954	C__12110132_10	8: 42587796	Intron	A/G	0.18
	rs9298629	C__29905827_10	8: 42606186	3' near gene	G/T	0.22
<i>CHRNA6</i>	rs2304297	C__15974192_10	8: 42608199	3' UTR	G/C	0.22
	rs892413	C__9179088_10	8: 42614378	Intron	C/A	0.22
	rs10087172	C__29616685_20	8: 42616868	Intron	T/C	0.22
	rs2196128	C__15936015_20	8: 42618286	Intron	T/C	0.23
	rs2217732	C__9179113_10	8: 42618446	Intron	A/G	0.22
	rs1072003	C__8779199_1_	8: 42620001	Intron	C/G	0.22

Note: SNP, single nucleotide polymorphism; Chr. Pos., chromosomal position; MAF, minor allele frequency based on HapMap-HCB; NA, not available.

Supplementary Figure 1. Distribution of FTND scores in the sample

