

**Table S2. Thirty Most Significant Association Results From the Analysis of Alcohol Dependence in the Dutch NESDA Sample**

Chr	Location (bp)	SNP	Gene	SNP Type	P-value
6	90731172	rs16882302	<i>BACH2</i>	Intron	$2.278 \times 10^{-7}$
22	46049871	rs11090754			$2.161 \times 10^{-6}$
21	42006728	rs13051745	<i>C21orf129</i>	Intron (boundary)	$4.778 \times 10^{-6}$
13	83474030	rs9546562			$6.829 \times 10^{-6}$
10	43383818	rs2393938	<i>ZNF239</i>	5' UTR	$7.126 \times 10^{-6}$
10	43371825	rs10793442	<i>ZNF239</i>	3' UTR	$7.640 \times 10^{-6}$
10	43370721	rs7893169	<i>ZNF239</i>	Downstream	$7.640 \times 10^{-6}$
10	43374973	rs7894102	<i>ZNF239</i>	Intron	$7.640 \times 10^{-6}$
10	51231805	rs17178655	<i>MSMB</i>	Intron	$7.784 \times 10^{-6}$
13	83489863	rs9546566			$1.023 \times 10^{-5}$
22	19654308	rs178260	<i>AIFL</i>	Intron	$1.365 \times 10^{-5}$
14	68270943	rs8019389			$1.956 \times 10^{-5}$
1	218174540	rs12129498	<i>SLC30A10</i>	Promoter	$2.706 \times 10^{-5}$
11	16791086	rs216495	<i>PLEKHA7</i>	Intron	$2.880 \times 10^{-5}$
19	41939312	rs826969	<i>MGC62100</i>	Intron	$3.146 \times 10^{-5}$
4	109150488	rs4956145	<i>HADHSC</i>	Coding exon	$4.110 \times 10^{-5}$
7	4162514	rs619672	<i>SDK1</i>	Intron	$4.593 \times 10^{-5}$
3	110120120	rs2715689	<i>GUCA1C</i>	Intron	$4.905 \times 10^{-5}$
13	29542418	rs9314995			$5.051 \times 10^{-5}$
5	118701596	rs17164572			$5.286 \times 10^{-5}$
5	118703145	rs3813301			$5.408 \times 10^{-5}$
5	118695541	rs7704398			$5.408 \times 10^{-5}$
6	83427171	rs376096			$5.557 \times 10^{-5}$
1	94889745	rs947386			$5.675 \times 10^{-5}$
6	46050909	rs3777599	<i>CLIC5</i>	Intron	$5.699 \times 10^{-5}$
14	40625514	rs1954456			$5.729 \times 10^{-5}$
14	40627078	rs1959909			$5.729 \times 10^{-5}$
6	46055426	rs4714899	<i>CLIC5</i>	Intron	$6.538 \times 10^{-5}$
14	26880311	rs1456796			$6.861 \times 10^{-5}$
20	48368185	rs928072	<i>LOC284751</i>	Downstream	$6.939 \times 10^{-5}$