

Supplementary Table 4. Association analysis of *CHRNA3* and *CHRNA6* polymorphisms with the nicotine dependence in Korean male subjects

Gene	Loci	Allele	NDS	NS	OR (95%CI)	p
<i>CHRNA3</i>	<i>rs13280604</i>	A>G	0.173	0.196	0.99 (0.36–2.75)	0.98
	<i>rs16891561</i>	C>T	0.171	0.191	1.03 (0.36–2.97)	0.96
	<i>rs4954</i>	A>G	0.115	0.127	1.06 (0.29–3.84)	0.93
<i>CHRNA6</i>	<i>rs9298628</i>	C>T	0.188	0.206	1.07 (0.37–3.12)	0.91
	<i>rs16891604</i>	C>A	0.219	0.211	1.12 (0.45–2.79)	0.81
	<i>rs10109040</i>	C>G	0.180	0.186	0.87 (0.29–2.68)	0.81
	<i>ht1</i>	$\Delta ht1 > ht1$	0.588	0.574	0.92 (0.41–2.07)	0.84

OR and corresponding p-value were calculated by using logistic models whilst controlling by age and sex. NDS: nicotine dependent smoker, NS: non-smoker of population control, CHRN: cholinergic nicotinic receptor, OR: odds ratio