

Erratum

CYP46A1 Functional Promoter Haplotypes Decipher Genetic Susceptibility to Alzheimer's Disease

Yan Lia, Leung-Wing Chu, Binbin Wang, Ping-Yiu, Yik, Huriletemuer, Dong-Yan Jin, Xu Ma and You-Qiang Song

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On page 1317, in Table 3, there are two wrong values in the table. Genotype CT; AD, *n* (%) says 63. This should be 49. Genotype CT; Control, *n* (%), it says 70. This should be 43. The correct table is presented below.

Table 3
 Frequency distribution of CYP46A1 promoter SNPs in Mongolian AD data set

	rs8003602	AD, <i>n</i> (%)	Control, <i>n</i> (%)	<i>P</i> - value	OR (95% CI)	rs3783320	AD, <i>n</i> (%)	Control, <i>n</i> (%)	<i>P</i> - value	OR (95% CI)
Allele	C	137 (64.0)	103 (51.5)	0.01	1.67 (1.13–2.48)	G	137 (64.0)	103 (51.5)	0.01	1.67 (1.13–2.48)
	T	77 (36.0)	97 (48.5)			A	77 (36.0)	97 (48.5)		
Genotype	CC	44 (41.1)	30 (30.0)	0.031	2.83 (1.27–6.26)	GG	44 (41.1)	30 (30.0)	0.031	2.83 (1.27–6.26)
	CT	49 (45.8)	43 (43.0)			GA	49 (45.8)	43 (43.0)		
	TT	14 (13.1)	27 (27.0)			AA	14 (13.1)	27 (27.0)		