

Erratum to: Association study of H2AFZ with schizophrenia in a Japanese case–control sample

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Table 1 contains several errors as originally published. Table 1 should read as follows; the corrected values are shown in *italics*.

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Table 1 Allelic and genotypic analysis of the *H2AFZ* gene

SNP ID	rs number	Affection	<i>N</i>	HWE <i>P</i>	Allelic count		MAF	Allelic <i>P</i>	OR (95 % CI)	Genotypic count			Genotypic <i>P</i>	
					C	T				CC	CT	TT		
SNP01		SCZ	1,999	0.001	3,995	3	0.001	1.000	1.081		1,997	1	1	0.486
rs4594733		CON	2,160	1.000	4,317	3	0.001	(1.000)	(0.218–5.357)		2,157	3	0	(0.886)
					A	C				AA	AC	CC		
SNP02		SCZ	2,008	0.033	3,048	968	0.241	0.114	1.086		1,139	770	99	0.236
rs2276939		CON	2,163	0.158	3,347	979	0.226	(0.377)	(0.981–1.202)		1,283	781	99	(0.567)
					C	T				CC	CT	TT		
SNP03		SCZ	1,983	0.264	2,864	1,102	0.278	0.640	1.024		1,044	776	163	0.421
rs10014424		CON	2,155	0.551	3,133	1,177	0.273	(0.942)	(0.93–1.128)		1,133	867	155	(0.839)
					A	G				AA	AG	GG		
SNP04		SCZ	1,986	0.964	1,906	2,066	0.480	0.058	0.9191		458	990	538	0.107
rs6854536		CON	2,147	0.244	2,151	2,143	0.501	(0.188)	(0.8,431–1.002)		525	1,101	521	(0.325)

Bold *P* values indicate $P < 0.05$

N number of subjects, *HWE* Hardy–Weinberg equilibrium, *MAF* minor allele frequency, *OR* odds ratio, *95 % CI* 95 % confidence interval, *SCZ* schizophrenia, *CON* control