



Correction to: Association Study Between Genetic Variation in Whole Mitochondrial Genome and Ischemic Stroke

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The original version of this article unfortunately contained a mistake in Table 3. The first row of the table body consisting the entries [A, A5, 2, 3.64, 0, 0, 0.50, -], was inadvertently captured as part of the table heading.

The original article can be found online at <https://doi.org/10.1007/s12031-020-01778-3>.

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The corrected version is presented below:

Table 3 Distributions of DNA haplotypes in IS and control groups

Haplogroup	Control group		IS group		<i>P</i> value	<i>OR</i> (95% <i>CI</i>)	
	Cases	Proportion (%)	Cases	Proportion (%)			
A	A5	2	3.64	0	0	0.50	-
	A6	0	0	1	1.92	0.49	-
	A14	1	1.82	1	1.92	1	1.06 (0.06–17.38)
	A15	2	3.64	1	1.92	1	0.52 (0.05–5.91)
	A19	1	1.82	0	0	1	-
B	B4	3	5.45	3	5.77	1	1.06 (0.20–5.51)
	B5	2	3.64	2	3.85	1	1.06 (0.14–7.81)
C	C4	1	1.82	0	0	1	-
	C5	0	0	1	1.92	0.49	-
	C7	2	3.64	0	0	0.496	-
D	D4	13	23.64	16	30.77	0.41	1.44 (0.61–3.38)
	D5	4	7.27	4	7.69	1	1.06 (0.25–4.49)
	D6	1	1.82	0	0	1	-
F	F1	4	7.27	5	9.62	0.93	1.36 (0.34–5.36)
	F2	0	0	3	5.77	0.22	1,813,288,114(0-)
G	G1	2	3.64	0	0	0.50	-
	G2	2	3.64	3	5.77	0.95	1.62 (0.26–10.12)
	G3	2	3.64	0	0	0.50	-
M	M10	1	1.82	0	0	1	-
	M11	0	0	1	1.92	0.49	-
	M12	1	1.82	0	0	1	-
	M61	1	1.82	0	0	1	-
	M7	4	7.27	3	5.77	1	0.78 (0.17–3.67)
	M8	2	3.64	2	3.85	1	1.06 (0.14–7.81)
	M9	1	1.82	1	1.92	1	1.06 (0.06–17.38)
N9	N9	2	3.64	2	3.85	1	1.06 (0.14–7.81)
R9	R9	0	0	1	1.92	0.49	-
U	U2	0	0	1	1.92	0.49	-
Z	Z3	0	0	1	1.92	0.49	-
	Z4	1	1.82	0	0	1	-

The original article has been corrected.

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