Additions and Corrections

Vol. 29 (2006) 353–360

Electrocardiogram Screening for Left High R-Wave Predicts Cardiovascular Death in a Japanese Community-Based Population: NIPPON DATA90

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Page 353: Abstract, Line 8

"in 4.1% of the 4,275 normotensives" should be "in 6.4% of the 4,275 normotensives."

Page 353: Abstract, Line 8

"128 participants died due to cardiovascular disease." should be "133 participants died due to cardiovascular disease."

Page 355: Results, Line 7

"12.4% had left high R-wave" should be "14.6% had left high R-wave."

Page 355: Results, Line 8

"in only 4.1% of 4,275 normotensives" should be "in only 6.4% of 4,275 normotensives."

The authors apologize for errors.

Vol. 29 (2006) 611–619

Association of Sixty-One Non-Synonymous Polymorphisms in Forty-One Hypertension Candidate Genes with Blood Pressure Variation and Hypertension

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In the version of this article initially published, Table 4 and the relevant results section were incorrect. The correct Table 4 and the relevant results sections are followed. The authors apologize for errors.

Page 613: Right-hand column, Lines 13–25

Rs754615 of *CAST* was associated with hypertension in women (GG+GC *vs*. CC, odds ratio=4.02, 95% confidence interval=1.47–11.0, p=0.007). Rs9016 of *GC* was associated with hypertension in women (CC *vs*. CT+TT, odds ratio=3.20, 95% confidence interval=1.52–6.71, p=0.002). Rs1390938 of *SLC18A1* was associated with hypertension in women (TT+TC *vs*. CC, odds ratio=1.37, 95% confidence interval= 1.03–1.82, p=0.033). Rs5036 of *SLC4A1* was associated with hypertension in men (AA *vs*. AG+GG, odds ratio=1.43, 95% confidence interval=1.04–1.97, p=0.031). Rs1063856 of *VWF* was associated with hypertension in men (AA *vs*. AG+GG, odds ratio=1.55, 95% confidence interval=1.04–2.31, p=0.034).

Page 617: Table 4

Table 4. Allele Frequency and Odds Ratio of Presence of Hypertension by Genotypes of Polymorphisms

Gene	SNP amino acid change	Allele1/2 (allele freq.)	Sex	Genotype group	Odds ratios (95% CI)	p*
CAST	rs754615	G/C	Women	GG+GC	1	
	C408S	(0.865/0.135)		CC	4.02 (1.47–11.0)	0.007
CTLA4	rs231775	G/A	Men	GG+GA	1	
	A17T	(0.617/0.383)		AA	0.67 (0.45–1.00)	0.050
F5	rs6020	A/G	Women	AA+AG	1	
	K513R	(0.650/0.350)		GG	1.71 (1.14–2.58)	0.010
GC	rs9016	C/T	Women	CC	1	
	R445C	(0.975/0.025)		CT+TT	3.20 (1.52–6.71)	0.002
GHR	rs6180	C/A	Women	CC+CA	1	
	L544I	(0.556/0.444)		AA	1.43 (1.00–2.02)	0.048
LIPC	rs6078	G/A	Men	GG	1	
	V95M	(0.760/0.240)		GA+AA	0.70 (0.52–0.93)	0.016
PLA2G7	rs1805018	T/C	Women	TT+TC	1	
	I198T	(0.793/0.207)		CC	0.43 (0.21–0.87)	0.020
SLC18A1	rs1390938	T/C	Women	TT+TC	1	
	I136T	(0.256/0.744)		CC	1.37 (1.03–1.82)	0.033
SLC4A1	rs5036	A/G	Men	AA	1	
	K56E	(0.841/0.159)		AG+GG	1.43 (1.04–1.97)	0.031
TRH	rs5658	G/C	Women	GG+GC	1	
	V8L	(0.328/0.672)		CC	1.58 (1.02–2.46)	0.041
VWF	rs1063856	A/G	Women	AA	1	
	T789A	(0.928/0.072)		AG+GG	1.55 (1.04–2.31)	0.034

*Conditional logistic analysis, adjusted for age, body mass index (BMI), present illness (hyperlipidemia and diabetes mellitus), and lifestyle (smoking and drinking). SNP, single nucleotide polymorphism; CI, confidence intervals.