

Erratum

Involvement of Human Histamine N-Methyltransferase Gene Polymorphisms in Susceptibility to Atopic Dermatitis in Korean Children

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Hee Seon Lee, Seung-Hyun Kim, Kyung Won Kim, Ji Young Baek, Hae-Sim Park, Kyung Eun Lee, Jung Yeon Hong, Mi Na Kim, Won II Heo, Myung Hyun Sohn, Kyu-Earn Kim. Involvement of human histamine N-methyltransferase gene polymorphisms in susceptibility to atopic dermatitis in Korean children. Allergy Asthma Immunol Res 2012;4:31-6. http://dx.doi.org/10.4168/aair.2012.4.1.31

We found an error in our published article. We apologize for any inconvenience this may have caused. The correct table is as below:

Corrected Table 3

Table 3. Genotype and allele frequencies of histamine N-methyltransferase (*HNMT*) polymorphisms in children with atopy

Locus	Genotype	Atopic eczema N=396, N (%)	Atopic control N=146, N (%)	<i>P</i> value OR (95% CI)
-465T>C	П	182 (46.5)	69 (47.3)	0.896
	TC	167 (42.7)	63 (43.2)	
	CC	42 (10.7)	14 (9.6)	
-413C>T	CC	331 (83.8)	123 (84.2)	0.322
	CT	63 (15.9)	22 (15.1)	
	Π	1 (0.3)	1 (0.7)	
314C>T	CC	359 (90.7)	136 (93.2)	1.000
	CT	37 (9.3)	9 (6.2)	
	Π	0 (0)	1 (0.7)	
939A>G	AA	208 (52.9)	79 (54.5)	0.048
	AG	167 (42.5)	51 (35.2)	0.415
	GG	18 (4.6)	15 (10.3)	(0.203-0.848)

P values were calculated using dominant models. Values in bold indicate statistical significance. For multiple comparisons of genotype and allele frequencies, Bonferroni's multiple adjustment was applied to the level of significance, which was set at P<0.0125 (0.05/4). Logistic regression analysis was applied to control for age and gender as covariables.

N, number of patients.

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