## **ERRATUM**

## **Erratum to: Genome-wide copy number variation** association study suggested VPS13B gene for osteoporosis in Caucasians

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Published online: 10 April 2010

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**Erratum to: Osteoporos Int (2010) 21:579–587** DOI 10.1007/s00198-009-0998-7

Table 3 unfortunately contained errors. The correct version is given here.

Table 3 Association between CNV 084 and bone phenotypes in the sample

Presented are mean (SD) [observation number]. In the total sample, age and gender were adjusted. In the genderstratified analyses, age was adjusted as a covariant. Marked in bold are data that remained significant after Bonferroni correction

Phenotype	Phenotypic value		P value		
	CNV state=1	CNV state=2	Total	Female	Male
SPN BMD	1.18 (0.22) [16]	1.03 (0.16) [965]	0.0001	0.0559	0.0003
Hip BMD	1.09 (0.20) [16]	0.97 (0.15) [963]	0.0002	0.0096	0.0071
FN BMD	0.92 (0.20) [16]	0.81 (0.14) [952]	0.0001	0.0032	0.0103
CT	0.18 (0.04) [16]	0.15 (0.03) [958]	0.0001	0.0029	0.0042
CSA	3.13 (0.77) [16]	2.83 (0.64) [958]	0.0030	0.0150	0.0510
BR	10.71 (2.92) [16]	12.04 (2.73) [958]	0.0170	0.1140	0.0710

The online version of the original article can be found at http://dx.doi. org/10.1007/s00198-009-0998-7.

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