

A genetic variant of *G6PC2* is associated with type 2 diabetes and fasting plasma glucose level in the Chinese population

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Unfortunately, the authors made a mistake when calculating HOMA-B, and the values reported in Table 3 and ESM Tables 3 and 4 were incorrect by a factor of 20. This mistake does not affect the findings of this study. The corrected version of Table 3 is reproduced here, and the corrected versions of ESM Tables 3 and 4 are available to authorised users.

Table 3 Associations of the rs16856187 genotype with clinical features related to glucose metabolism in the participants with normal glucose regulation

	AA (n=866)	AC (n=708)	CC (n=176)	p value
Fasting plasma glucose (mmol/l)	4.96±0.49	4.99±0.50	5.13±0.49	0.0002
2 h plasma glucose (mmol/l)	5.46±1.22	5.46±1.20	5.43±1.10	0.6929
Fasting insulin (pmol/l)	40.86 (27.66–59.64)	41.58 (28.14–59.34)	40.98 (28.44–61.62)	0.4021
2 h insulin (pmol/l)	217.86 (135.42–324.42)	210.12 (132.18–329.82)	229.74 (140.94–351.00)	0.2324
HOMA-IR	1.47 (0.98–2.22)	1.51 (0.99–2.19)	1.58 (1.01–2.23)	0.1884
HOMA-B	101.25 (70.96–155.33)	101.88 (69.62–147.82)	98.39 (64.59–142.70)	0.3727

Data are shown as mean±SD or median (interquartile range)
p values were adjusted for age, sex and BMI in an additive genetic model

The online version of the original article can be found at <http://dx.doi.org/10.1007/s00125-008-1241-3>.

Electronic supplementary material The online version of this article (doi:10.1007/s00125-009-1286-y) contains the corrected versions of Tables 3 and 4, and is available to authorised users.

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