CORRECTION

Estimation of non-additive genetic variance in human complex traits from a large sample of unrelated individuals

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In this paper, the mean estimates of SNP-based heritability and additive-by-additive variance across traits are $\overline{h}^2_{SNP} = 0.208 \pm 0.001$ (CI95% = [0.206;0.210]) and $\overline{\eta}^2_{SNP} = 0.055 \pm 0.032$ (CI95% = [-0.010;0.119]), as shown correctly in Figure 4. However, in the original text (summary and the 8th and 10th pages of the article), they were erroneously referred to as $\overline{h}^2_{SNP} = 0.207 \pm 0.001$ (CI95% = [0.205;0.208]) and $\overline{\eta}^2_{SNP} = 0.058 \pm 0.032$ (CI95% = [−0.005;0.122]). There was also a typo on the 9th page (0.0.157 instead of 0.157). These errors, which do not affect the conclusion of the paper, have been corrected online and in print. The authors apologize for these errors.