

Author Correction: Deep learning and genome-wide association meta-analyses of bone marrow adiposity in the UK Biobank

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 Check for updates

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In the version of this article initially published, in the fourth paragraph of the “Sex sensitivity analysis for meta-GWAS in the white population” section, in the sentence now reading “Notably, we identified three sex-specific genes: *AKNA* and *WHRN* were associated with total hip BMFF in males only, while *VMPI* was associated with spine BMFF in females only,” *AKNA* originally appeared as “*ANKA*.” The error has been corrected in the HTML and PDF versions of the article.

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