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Author Correction: Donkey genomes provide new insights into domestication and selection for coat color

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Correction to: *Nat. Commun.* <https://doi.org/10.1038/s41467-020-19813-7>, published online 8 December 2020.

The original version of this Article contained an error in the author contributions, which incorrectly omitted the following: 'J.W. led the experiments to study the genetic mechanism underlying the emergence of the non-Dun coat colors of donkeys, including performing RNA-seq of croup skin samples of Dun and non-Dun donkeys, performing the immunohistochemistry and immunofluorescence assays against donkey TBX3 protein and bioinformatics analysis of the 1 bp deletion downstream the TBX3 gene. J.W. also contributed to the plotting of Figures'.

The original version of this Article also contained an error in the author affiliations. The affiliation of Antonia Noce with Leibniz-Institute for Farm Animal Biology (FBN), Dummerstorf 18196, Germany, was inadvertently omitted.

These have now been corrected in both the PDF and HTML versions of the Article.

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