

# Author Correction: *Cicer* super-pangenome provides insights into species evolution and agronomic trait loci for crop improvement in chickpea

Correction to: *Nature Genetics*  
<https://doi.org/10.1038/s41588-024-01760-4>,  
published online 23 May 2024

<https://doi.org/10.1038/s41588-024-01813-8>

Published online: 29 May 2024

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In the version of the article initially published, various numbers in the main text did not match the corresponding data in Table 1 and Supplementary Table 6. In the “Genome assembly and annotation of annual wild *Cicer* genomes” section, “25,761” was corrected to “25,546” in the sentence “...we predicted an average of 25,546 gene models for each *Cicer* species”; “23,486” and “29,642” have been corrected to “23,516” and “28,627” in the sentence “The number of genes ranged from 23,516 for *C. judaicum* to 29,627 for *C. reticulatum*”; “11,505” has been corrected to “11,590” in the sentence “...and 11,505 ribosomal RNA (rRNA) genes for each *Cicer* species”; and “1,292” has been corrected to “1,297” in the sentence “...with the highest number recorded for *C. reticulatum* (1,297)”. In the first sentence of the “Graph super-pangenome for *Cicer* genus and SV genotyping” section, “460,311” has been amended to “465,535”. These corrections have been made in the HTML and PDF versions of the article.

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