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Author Correction: Whole-genome resequencing of 472 *Vitis* accessions for grapevine diversity and demographic history analyses

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Correction to: *Nature Communications* <https://doi.org/10.1038/s41467-019-09135-8>, published online 13 March 2019.

In the original version of this Article, in the last paragraph of the first section of the Results, the ratios of SNPs located in the intergenic regions, SNPs located in the coding sequences, and indels located in the coding regions are incorrectly written as ‘71.2%’, ‘4.6%’, and ‘1.4%’, respectively. These ratios should read as ‘73.7%’, ‘4.0%’, and ‘1.3%’, respectively. In addition, the estimated ratio of indels in the coding regions that could cause frameshift mutations is incorrectly written as ‘66.7%’. It should read as ‘66.9%’. Furthermore, authors’ given names and surnames of Reference 47 are switched. The authors’ names should read ‘Grassi F., De Mattia F., Zecca G., Sala F., & Labra M.’. These have been corrected in both the PDF and HTML versions of the Article.

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