CORRECTION

Correction: Positively selected amino acid replacements within the RuBisCO enzyme of oak trees are associated with ecological adaptations

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The images for Figs 1 and 2 are incorrectly switched. The image that appears as Fig 1 should be Fig 2, and the image that appears as Fig 2 should be Fig 1. The figure captions appear in the correct order.

There are also errors in the figure captioned “*Quercus* large dataset Bayesian phylogram based on 158 *rbcL* sequences.” Please see the correct Figs 1 and 2 below.

OPEN ACCESS

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Fig 1. *Quercus* large dataset Bayesian phylogram based on 158 *rbcL* sequences. Numbers above branches correspond to Bayesian posterior probabilities. The figure was edited using FigTree Version 1.4.0 [77].

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Fig 2. Quercus small dataset Bayesian phylogeny based on 45 sequences of *rbCL*, 43 *matK* and 42 microsatellites. Numbers above branches correspond to Bayesian posterior probabilities. The figure was edited using FigTree Version 1.4.0 [77].

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Reference

1. Raza FA, Rehman Su, Khalid R, Ahmad J, Ashraf S, Iqbal M et al. (2014) Demographic and Clinico-Epidemiological Features of Dengue Fever in Faisalabad, Pakistan. PLoS ONE 9(3): e89868. <https://doi.org/10.1371/journal.pone.0089868> PMID: 24595236