

CORRECTION

Correction: Path Similarity Analysis: A Method for Quantifying Macromolecular Pathways

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Notice of Republication

This article was republished on June 11, 2018, to correct errors throughout the manuscript that were introduced due to a technical error made during the plotting of the data. The editors confirm the major conclusions of the manuscript remain accurate. Please download this article again to view the correct version.

Supporting information

S1 File. Republished, corrected article.

(PDF)

Reference

1. Seyler SL, Kumar A, Thorpe MF, Beckstein O (2015) Path Similarity Analysis: A Method for Quantifying Macromolecular Pathways. PLoS Comput Biol 11(10): e1004568. <https://doi.org/10.1371/journal.pcbi.1004568> PMID: 26488417



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