PLOS GENETICS

CORRECTION

Correction: Filling gaps in bacterial amino acid biosynthesis pathways with high-throughput genetics

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The reference list for S1 Text is shown at the end of the S3 Text file. Please view the correct S1 Text and S3 Text files below, which include the correct reference lists for each file.

Supporting information

S1 Text. Testing the auxotroph predictions of D'Souza and colleagues.
(PDF)

S3 Text. Clear candidates whose mutants were not consistently auxotrophic.
(PDF)

Reference

1. Price MN, Zane GM, Kuehl JV, Melnyk RA, Wall JD, Deutschbauer AM, et al. (2018) Filling gaps in bacterial amino acid biosynthesis pathways with high-throughput genetics. PLoS Genet 14(1): e1007147. https://doi.org/10.1371/journal.pgen.1007147 PMID: 29324779

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