

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.

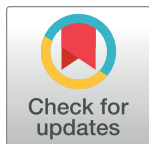
## Supporting information

**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



## OPEN ACCESS

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