

CORRECTION

Correction: A mechanistic pan-cancer pathway model informed by multi-omics data interprets stochastic cell fate responses to drugs and mitogens

The *PLOS Computational Biology* Staff

The supporting information item [S1 Code](#) is incomplete.

The final, peer-reviewed version of [S1 Code](#) can be viewed here:

The publisher apologizes for this error.

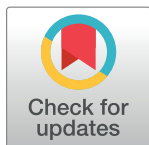
Supporting information

S1 Code. Matlab model code.

(ZIP)

Reference

1. Bouhaddou M, Barrette AM, Stern AD, Koch RJ, DiStefano MS, Riesel EA, et al. (2018) A mechanistic pan-cancer pathway model informed by multi-omics data interprets stochastic cell fate responses to drugs and mitogens. *PLoS Comput Biol* 14(3): e1005985. <https://doi.org/10.1371/journal.pcbi.1005985> PMID: [29579036](https://pubmed.ncbi.nlm.nih.gov/29579036/)



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Citation: The *PLOS Computational Biology* Staff (2018) Correction: A mechanistic pan-cancer pathway model informed by multi-omics data interprets stochastic cell fate responses to drugs and mitogens. *PLoS Comput Biol* 14(7): e1006189. <https://doi.org/10.1371/journal.pcbi.1006189>

Published: July 9, 2018

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