

CORRECTION

Correction: Multiscale mutation clustering algorithm identifies pan-cancer mutational clusters associated with pathway-level changes in gene expression

The *PLOS Computational Biology* Staff

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Reference

1. Poole W, Leinonen K, Shmulevich I, Knijnenburg TA, Bernard B (2017) Multiscale mutation clustering algorithm identifies pan-cancer mutational clusters associated with pathway-level changes in gene expression. PLoS Comput Biol 13(2): e1005347. doi:[10.1371/journal.pcbi.1005347](https://doi.org/10.1371/journal.pcbi.1005347) PMID: [28170390](https://pubmed.ncbi.nlm.nih.gov/28170390/)



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