

Author Correction: Systematic profiling of conditional pathway activation identifies context-dependent synthetic lethalties

Correction to: *Nature Genetics* <https://doi.org/10.1038/s41588-023-01515-7>, published online 25 September 2023.

<https://doi.org/10.1038/s41588-024-01680-3>

Published online: 23 February 2024

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In the version of the article initially published, Fig. 5f was originally divided into two panels, f and g, but these have now been combined to reflect the fact that they are from the same experiment. Panel g contained a duplicated NFT1 row from panel f, followed by the CSNK1A1 shRNA row, which now appears in panel f. Fig. 5 has been amended in the HTML and PDF versions of the article.

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Author Correction: GATK-gCNV enables the discovery of rare copy number variants from exome sequencing data

Correction to: *Nature Genetics* <https://doi.org/10.1038/s41588-023-01449-0>, published online 21 August 2023.

<https://doi.org/10.1038/s41588-024-01663-4>

Published online: 23 January 2024

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In the version of the article initially published, there were errors in equations in the Methods section. Equation 2, now reading $\lambda_{st} = d_s \mu_{st} c_{st} + d_s \epsilon_M$, previously read $\lambda_{st} = d_s c_{st} + d_s \epsilon_M$. The first line of Equation 3, now reading $\log(\mu_{st}) = m_t + \sum_{v=1}^D W_{tv} z_{vs} + \sum_{v=1}^K \bar{W}_{tv} \bar{z}_{vs}$, previously read $\log(\mu_{st}) = m_t + \sum_{v=1}^D W_{tv} z_{vs} + \sum_{v=1}^K \bar{W}_{tv} \bar{z}_{tv}$. In addition, the Supplementary Note has been updated. These changes have been made to the HTML and PDF versions of the article.

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