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Author Correction: A consensus *S. cerevisiae* metabolic model Yeast8 and its ecosystem for comprehensively probing cellular metabolism

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In the original version of this Article, the calculation of the WAP score in the second paragraph of the “Methods” subsection “CLUMPS method to calculate p values of mutation-enriched PDB files” was incorrectly written as “Next, assuming that the uniform distribution of mutations across the protein residues covers the given structure, calculate each WAP score in 10 randomisations to obtain the null distribution.” It should read “Next, assuming that the uniform distribution of mutations across the protein residues covers the given structure, calculate each WAP score in 10^4 randomisations to obtain the null distribution.”

This has been corrected in both the PDF and HTML versions of this Article.

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