

## CORRECTION

# Correction: CTD: An information-theoretic algorithm to interpret sets of metabolomic and transcriptomic perturbations in the context of graphical models

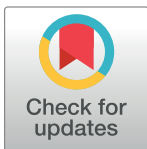
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There is a typo in Eq 1. A  $\log_2()$  is missing before the fraction  $\frac{P_A(S)}{P_0(S)}$ . The authors have provided the correct version of the equation below.

$$P_0 \left( \log_2 \left( \frac{P_A(S)}{P_0(S)} \right) > d \right) < 2^{-d}$$

## Reference

1. Thistlethwaite LR, Petrosyan V, Li X, Miller MJ, Elsea SH, Milosavljevic A (2021) CTD: An information-theoretic algorithm to interpret sets of metabolomic and transcriptomic perturbations in the context of graphical models. *PLoS Comput Biol* 17(1): e1008550. <https://doi.org/10.1371/journal.pcbi.1008550> PMID: [33513132](https://pubmed.ncbi.nlm.nih.gov/33513132/)



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