

Correction



Correction: An *In-Silico* Model of Lipoprotein Metabolism and Kinetics for the Evaluation of Targets and Biomarkers in the Reverse Cholesterol Transport Pathway

The *PLOS Computational Biology* Staff

The parameter k_f in Table 5 is incorrect. It should be given as 50000. Correspondingly the standard deviation for the parameter k_f should also be ± 15440 .

Reference

1. Lu J, Hübner K, Nanjee MN, Brinton EA, Mazer NA (2014) An In-Silico Model of Lipoprotein Metabolism and Kinetics for the Evaluation of Targets and Biomarkers in the Reverse Cholesterol Transport Pathway. *PLoS Comput Biol* 10(3): e1003509. doi:10.1371/journal.pcbi.1003509.

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