

Author Correction for Côté et al., The Genome-Wide Interaction Network of Nutrient Stress Genes in *Escherichia coli*

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Vol. 7, no. 6, doi: [10.1128/mBio.01714-16](https://doi.org/10.1128/mBio.01714-16), 2016. We report herein two referencing oversights in this article. We wish to emphasize here the important work by Ringlstetter (1), who first linked *yigM* to biotin transport in *Escherichia coli*. Our article overlooked the discovery of *yigM* as the coding gene for biotin transporter. Overlooked also was the work of Finkenwirth et al. (2), who created a $\Delta yigM \Delta bioH$ deletion mutant (*E. coli*) and characterized biotin transporters from various organisms. These corrections do not impact the integrity of the data presented, nor are the dominant conclusions of the study altered.

REFERENCES

1. Ringlstetter SL. 2010. Identification of the biotin transporter in *Escherichia coli*, biotinylation of histones in *Saccharomyces cerevisiae* and analysis of biotin sensing in *Saccharomyces cerevisiae*. Ph.D. thesis. University of Regensburg, Regensburg, Germany.
2. Finkenwirth F, Kirsch F, Eitinger T. 2013. Solitary BioY proteins mediate biotin transport into recombinant *Escherichia coli*. *J Bacteriol* 195:4105–4111. <http://dx.doi.org/10.1128/JB.00350-13>.

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