

CORRECTION

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Correction to: Transcriptional responses of *Escherichia coli* during recovery from inorganic or organic mercury exposure

Stephen P. LaVoie* and Anne O. Summers*

Correction

After publication of the original article [1] the authors noted that the key displayed in Figure 1a was incorrect, as the PMA and HgCl₂ conditions had been switched.

The correct information follows below:

MG1655 unexposed condition: blue line

MG1655 + 3μM PMA: green line

MG1655 + 3μM HgCl₂: red line.

The revised figure is included with this Correction (Fig 1).

Corrected Figures:

Revised Fig. 1:

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Reference

1. LaVoie SP, Summers AO. Transcriptional responses of *Escherichia coli* during recovery from inorganic or organic mercury exposure. *BMC Genomics*. 2018; 19:52. <https://doi.org/10.1186/s12864-017-4413-z>.

* Correspondence: slavoie5@uga.edu; slavoie5@gmail.com; summers@uga.edu

Department of Microbiology, University of Georgia, Athens, GA 30602, USA



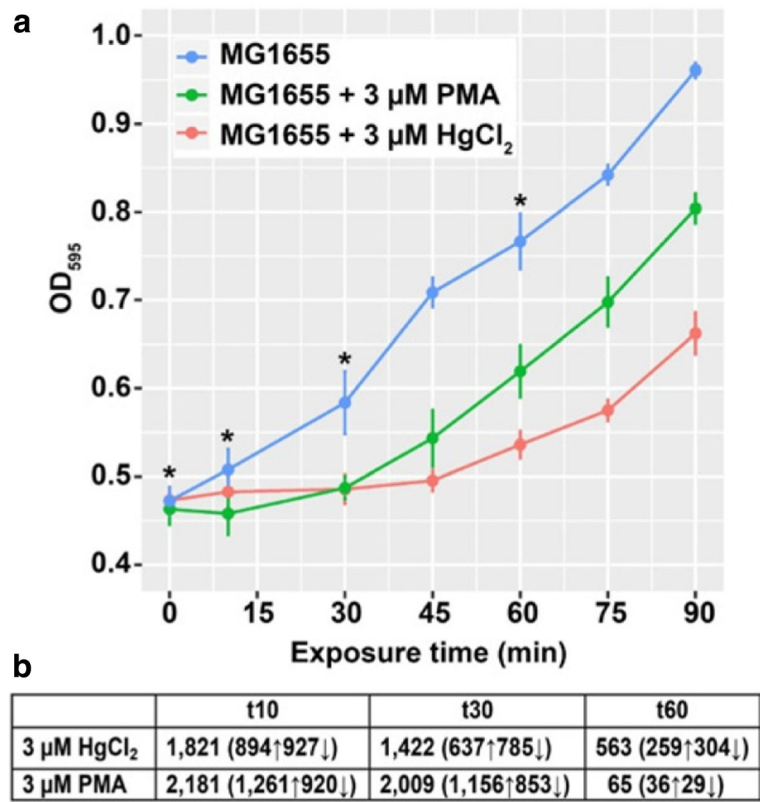


Fig. 1 Effects of sub-acute mercury exposure on growth of MG1655. **a** *E. coli* K12 MG1655 grown in MOPS minimal medium, unexposed (blue) or exposed to 3 μM HgCl₂ (red) or 3 μM PMA (green) during mid-log phase. Asterisks indicate sampling times for RNA-seq. Error bars are standard error (SEM) of 3 biological replicates for each culture condition. See Additional file 1: Figure S1 for full growth curve. **b** Significantly differentially expressed genes (DEG) counts (up-regulated or down-regulated) for HgCl₂ and PMA relative to unexposed control culture at each time point