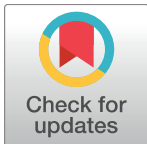


CORRECTION

Correction: Real-time cell toxicity profiling of Tox21 10K compounds reveals cytotoxicity dependent toxicity pathway linkage

Jui-Hua Hsieh, Ruili Huang, Ja-An Lin, Alexander Sedykh, Jinghua Zhao, Raymond R. Tice, Richard S. Paules, Menghang Xia, Scott S. Auerbach

The images for Figs 2 and 3 are incorrectly switched. The image that appears as Fig 2 should be Fig 3 and the image that appears as Fig 3 should be Fig 2. The figure captions appear in the correct order.



 OPEN ACCESS

Citation: Hsieh J-H, Huang R, Lin J-A, Sedykh A, Zhao J, Tice RR, et al. (2017) Correction: Real-time cell toxicity profiling of Tox21 10K compounds reveals cytotoxicity dependent toxicity pathway linkage. PLoS ONE 12(7): e0181291. <https://doi.org/10.1371/journal.pone.0181291>

Published: July 7, 2017

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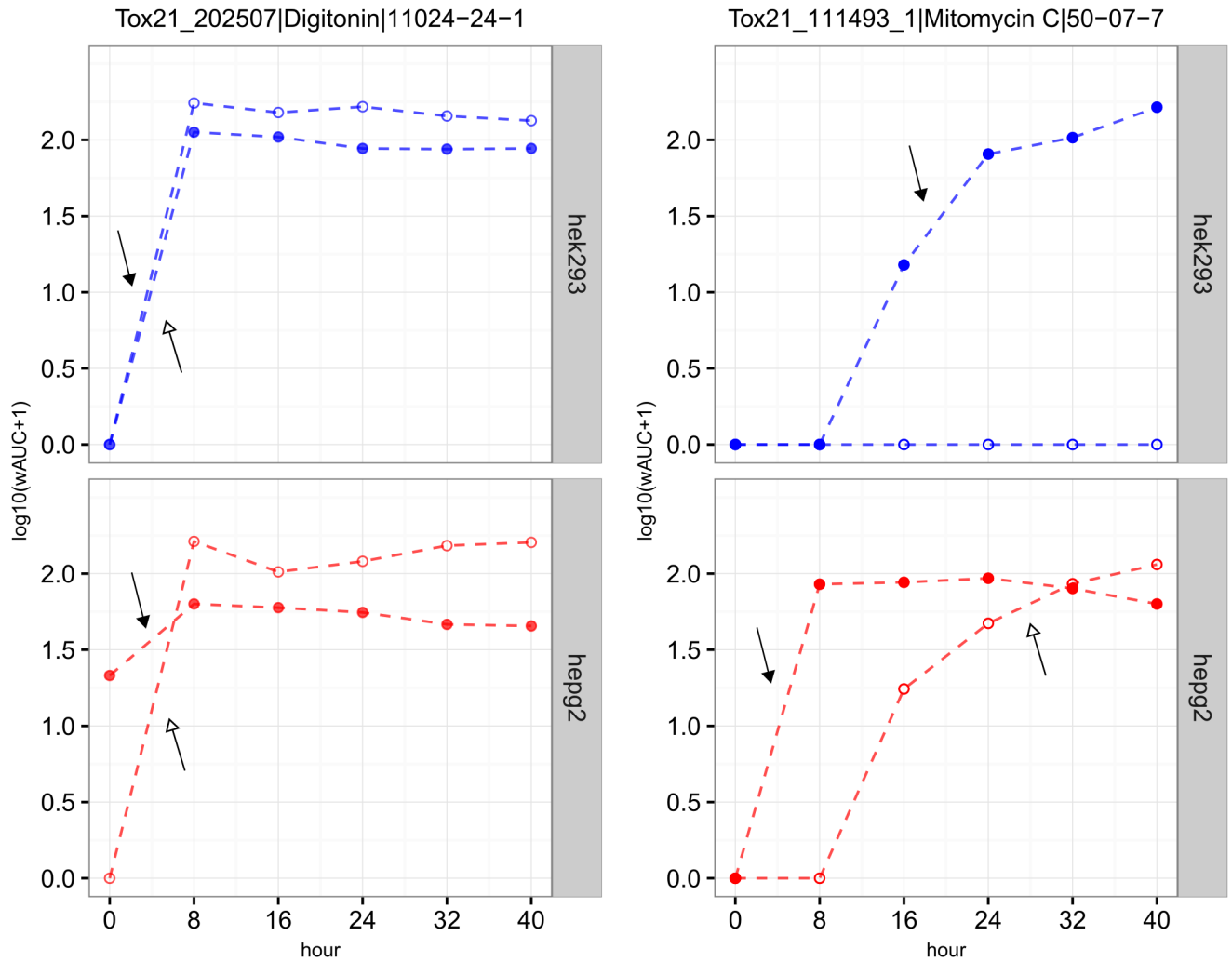


Fig 2. Examples of two chemicals (digitonin vs. mitomycin C) with different kinetics of cytotoxicity. Blue: HEK293; red: HepG2. Filled circle (*glo*); hollow circle (*flor*); the arrow represents the earliest time interval where the maximum cytotoxic effect was obtained (filled arrow head: *glo*; hollow arrow head: *flor*).

<https://doi.org/10.1371/journal.pone.0181291.g001>

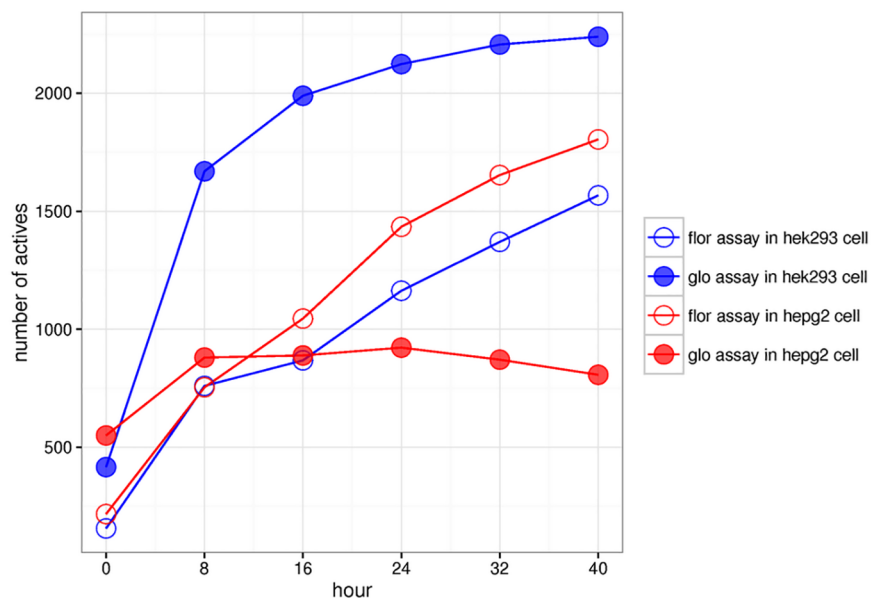


Fig 3. Comparison of number of actives in four assays. The number of actives detected in the four assays at the six different time points. Blue: HEK293 cell line; red: HepG2 cell line. Filled circle: *glo* assay technology; hollow circle: *flor* assay technology.

<https://doi.org/10.1371/journal.pone.0181291.g002>

Reference

1. Hsieh J-H, Huang R, Lin J-A, Sedykh A, Zhao J, Tice RR, et al. (2017) Real-time cell toxicity profiling of Tox21 10K compounds reveals cytotoxicity dependent toxicity pathway linkage. PLoS ONE 12(5): e0177902. <https://doi.org/10.1371/journal.pone.0177902> PMID: 28531190