



## Correction to “Structure-Based in Silico Screening Identifies a Potent Ebolavirus Inhibitor from a Traditional Chinese Medicine Library”

Faraz Shaikh,<sup>ID</sup> Yuguang Zhao, Luis Alvarez, Maria Iliopoulou, Christopher Lohans,<sup>ID</sup> Christopher J. Schofield,<sup>ID</sup> Sergi Padilla-Parra, Shirley W. I. Siu, Elizabeth E. Fry, Jingshan Ren,<sup>\*ID</sup> and David I. Stuart<sup>\*ID</sup>

*J. Med. Chem.* 2019, 62, 6, 2928–2937. DOI: [10.1021/acs.jmedchem.8b01328](https://doi.org/10.1021/acs.jmedchem.8b01328)

Page 2928. Abstract: “database of natural compounds”, should be replaced with “database of natural compounds and derivatives”.

Page 2929. Line 5, right column, replace “Natural Compound” with “Natural Compound Derivative”.

Page 2929. Line 7, right column: replace “natural compound library” with “library of natural compounds and derivatives”.

Page 2929. Line 9, left column. Replace “traditional Chinese medicine (TCM) database” with “database of traditional Chinese medicines and derivatives”.

Page 2929. Figure 2 legend: Change title from “Natural compounds...” to “Compounds...”.

Page 2933. Experimental Section lines 1–2. Replace “natural ligand” with “ZINC”.

Page 2933. Experimental section line 4: remove “natural ligand”.

Page 2934. Left column, Virtual Screening and Binding Affinity Calculation section, line 2. Remove “natural”.

Published: October 28, 2019

