Correction to: A multi-target approach for discovery of antiviral compounds against dengue virus from green tea

Pooja Mahajan¹ · Shantanu Tomar² · Adarsh Kumar¹ · Neelesh Yadav³ · Aditya Arya¹ · Vivek Dhar Dwivedi¹

Published online: 23 July 2020

© Springer-Verlag GmbH Austria, part of Springer Nature 2020

Correction to:

Network Modeling Analysis in Health Informatics and Bioinformatics (2020) 9:20
<https://doi.org/10.1007/s13721-020-0222-4>

Publisher's Note Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.

In results and discussion section, the description of figures was revised as follows. In the case of NS1-Compound-1 complex, consider an additional contact of Glu12 (Fig. 1a), in case of NS1-Compound-2 complex, six instead of four residues were involved in contact (Fig. 1b). In case of RDRP-Compound-1 complex, seven instead of three hydrogen bonds were formed (Fig. 1d), while in case of RDRP Compound-2 complex nine instead of seven hydrogen bonds were recorded (Fig. 1e). In case of MTD-complex 1 consider an addition contact of Leu17 (Fig. 1g).

The original article can be found online at <https://doi.org/10.1007/s13721-020-0222-4>.

✉ Aditya Arya
contact.adityarya@gmail.com

✉ Vivek Dhar Dwivedi
vivek_bioinformatics@yahoo.com

¹ Center for Bioinformatics, Computational and Systems Biology, Pathfinder Research and Training Foundation, Knowledge Park-III, Greater Noida, India

² Department of Biotechnology, GLA University, Mathura, India

³ Information Technology Cell, Tropical Forest Research Institute, Jabalpur, India