








# Virtual screening of bioactive anti-SARS-CoV natural products and identification of 3 $\beta$ ,12-diacetoxyabieta-6,8,11,13-tetraene as a potential inhibitor of SARS-CoV-2 virus and its infection related pathways by MD simulation and network pharmacology

A.K.M. Moyeenul Huq<sup>a,b,\*</sup> , Miah Roney<sup>a,c,\*</sup> , Syahrul Imran<sup>d,e</sup> , Shafi Ullah Khan<sup>f</sup> , Md. Nazim Uddin<sup>g</sup>, Thet Thet Htar<sup>h</sup>, Atif Amin Baig<sup>i#</sup>, Mohiuddin Ahmed Bhuiyan<sup>b</sup>, Zainul Amiruddin Zakaria<sup>j</sup>, Mohd Fadhilzil Fasihi Mohd Aluwi<sup>a,c</sup> and Saiful Nizam Tajuddin<sup>a,c</sup> 

<sup>a</sup>Bio Aromatic Research Centre, Universiti Malaysia Pahang, Kuantan, Pahang, Malaysia; <sup>b</sup>School of Medicine, Department of Pharmacy, University of Asia Pacific, Dhaka, Bangladesh; <sup>c</sup>Faculty of Industrial Sciences and Technology, Universiti Malaysia Pahang, Kuantan, Pahang, Malaysia; <sup>d</sup>Atta-ur-Rahman Institute for Natural Product Discovery (AuRIIns), Universiti Teknologi MARA Cawangan Selangor Kampus Puncak Alam, Puncak Alam, Selangor, Malaysia; <sup>e</sup>Faculty of Applied Science, Universiti Teknologi MARA (UiTM), Shah Alam, Selangor, Malaysia; <sup>f</sup>Product & Process Innovation Department, Qarshi Brands (Pvt) Ltd, Haripur, KPK, Pakistan; <sup>g</sup>Institute of Food Science and Technology, Bangladesh Council of Scientific and Industrial Research, Dhaka, Bangladesh; <sup>h</sup>School of Pharmacy, Monash University Malaysia, Subang Jaya, Selangor, Malaysia; <sup>i</sup>Faculty of Medicine, Universiti Sultan Zainal Abidin, Kuala Terengganu, Terengganu, Malaysia; <sup>j</sup>Department of Biomedical Sciences, Faculty of Medicine and Health Sciences, Universiti Malaysia Sabah, Jalan UMS, Kota Kinabalu, Sabah, Malaysia

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## ABSTRACT

Since the first prevalence of COVID-19 in 2019, it still remains the most devastating pandemic throughout the world. The current research aimed to find potential natural products to inhibit the novel coronavirus and associated infection by MD simulation and network pharmacology approach. Molecular docking was performed for 39 natural products having potent anti-SARS-CoV activity. Five natural products showed high binding interaction with the viral main protease for the SARS-CoV-2 virus, where 3 $\beta$ ,12-diacetoxyabieta-6,8,11,13 tetraene showed stable binding in MD simulation until 100 ns. Both 3 $\beta$ ,12-diacetoxyabieta-6,8,11,13 tetraene and tomentin A targeted 11 common genes that are related to COVID-19 and interact with each other. Gene ontology development analysis further showed that all these 11 genes are attached to various biological processes. The KEGG pathway analysis also showed that the proteins that are targeted by 3 $\beta$ ,12-diacetoxyabieta-6,8,11,13 tetraene and tomentin A are associated with multiple pathways related to COVID-19 infection. Furthermore, the ADMET and MDS studies reveals 3 $\beta$ ,12-diacetoxyabieta-6,8,11,13 as the best-suited compound for oral drug delivery.

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## KEYWORDS

COVID-19; docking; ADMET; natural products; MD simulation; network pharmacology

## 1. Introduction


Coronaviruses (CoVs) are the most frequent source of mild to acute respiratory illnesses. In 2003 and 2012, two especially virulent CoVs, the severe acute respiratory syndrome coronavirus (SARS-CoV) and the Middle East respiratory syndrome coronavirus (MERS-CoV), both transferred through animals to people, sparked global pandemics with significant mortality rates (Cavasotto & Di Filippo, 2021). In December 2019, the public health officials of China first reported to the World Health Organization (WHO) about an unknown virus from Wuhan which caused illness similar to pneumonia (Khan & Fahad, 2020). They quickly identified the virus as a member of the coronavirus (CoV) category, which was fast propagating outside from Wuhan to other places or countries and

was named novel coronavirus disease 2019 (COVID-19) (Hasan et al., 2021) with flue like symptoms including fever, cough, pneumonia, nausea and tiredness. WHO declared a worldwide emergency on 30<sup>th</sup> January 2020 and a pandemic on 11th March 2020 for the coronavirus illness (COVID-19) (Hua & Shaw, 2020). As of August 07, 2021, the global number of corona virus cases had reached 202.59 million, with 4.29 million deaths (<https://www.worldometers.info/coronavirus/>). COVID-19 is regarded as a serious public health threat worldwide. During the last 1.5 years, different variants of this virus have evolved with greater transmissibility and associated disease severely (<https://www.who.int/en/activities/tracking-SARS-CoV-2-variants/>). With the evolution of vaccines for SARS-CoV-2 and continuous vaccination through world, a significant decline in new transmission and death brought hope

**CONTACT** A. K. M. Moyeenul Huq  [moyeenul@ump.edu.my](mailto:moyeenul@ump.edu.my)  Bio Aromatic Research Centre, Universiti Malaysia Pahang, Kuantan, Pahang, Malaysia; Mohd Fadhilzil Fasihi Mohd Aluwi  [fasihi@ump.edu.my](mailto:fasihi@ump.edu.my)  Bio Aromatic Research Centre, Universiti Malaysia Pahang, Kuantan, Pahang, Malaysia.

\*Both authors contributed equally.

<sup>#</sup>University Institute of Public Health, Faculty of Allied Health Sciences, The University of Lahore, Pakistan.

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