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Why comprehensive datasets matter when inferring epidemic links or subgenotyping

Bram Vrancken, Mahmoudreza Pourkarim, Samad Amini-Bavil-Olyaee, Ali Aminy, Marc Van Ranst, Seyed Moaied Alavian

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Letter to the Editor

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Bram Vrancken^a, Mahmoudreza Pourkarim^{a,b}, Samad Amini-Bavil-Olyaee^c, Ali Aminy^d, Marc Van Ranst^a, and Seyed Moaied Alavian^{e*}

^aKU Leuven, Department of Microbiology and Immunology, Laboratory of Clinical and Epidemiological Virology, 3000, Leuven, Belgium

^bBlood Transfusion Research Centre, High Institute for Research and Education in Transfusion Medicine, Hemmat Exp. Way, 14665-1157, Tehran, Iran

^cBiosafety Development Group, Cellular Sciences Department, Amgen Inc., One Amgen Center Drive, Thousand Oaks, CA 91320, USA

^dFaculty of Medicine, University of Debrecen, Nagyerdei krt. 98, 4032 Debrecen, Hungary

^eBaqiyatallah Research Center for Gastroenterology and Liver Disease, Baqiyatallah University of Medical Sciences, Tehran, Iran.

*Correspondence to Seyed Moayed Alavian

Email: editor@hepatmon.com

Phone: +98 21 88352501-4

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