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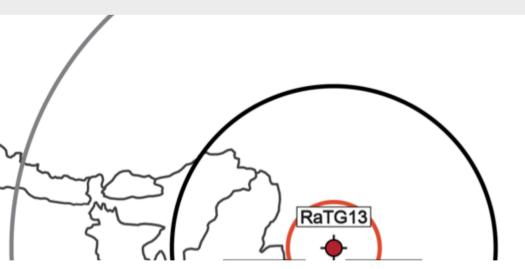
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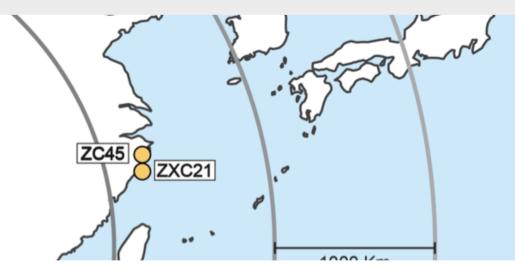
Inferring the ecological niche of bat viruses closely related to SARS-CoV-2 using phylogeographic analyses of *Rhinolophus* species

Alexandre Hassanin ⊡, Vuong Tan Tu, Manon Curaudeau & Gabor Csorba

Scientific Reports 11, Article number: 14276 (2021) Cite this article

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BIOTECHNOLOGY, COMMENTARIES, HEALTH AUGUST 2, 2021

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New Results

ACE2 binding is an ancestral and evolvable trait of sarbecoviruses

Digital Tyler N. Starr, Samantha K. Zepeda, Digital Alexandra C. Walls, Allison J. Greaney, Digital Veesler,

Jesse D. Bloom

doi: https://doi.org/10.1101/2021.07.17.452804

This article is a preprint and has not been certified by peer review [what does this mean?].

Molecular Evidence for SARS-CoV-2 in Samples Collected From Patients With Morbilliform Eruptions Since Late Summer 2019 in Lombardy, Northern Italy

27 Pages • Posted: 6 Aug 2021

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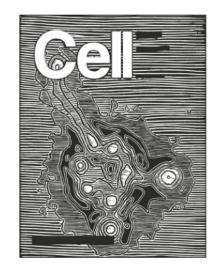
Temple University - Institute for Genomics and Evolutionary Medicine

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The Origins of SARS-CoV-2: A Critical Review

Edward C. Holmes, Stephen A. Goldstein, Angela L. Rasmussen, David L. Robertson, Alexander Crits-Christoph, Joel O. Wertheim, Simon J. Anthony, Wendy S. Barclay, Maciej F. Boni, Peter C. Doherty, Jeremy Farrar, Jemma L. Geoghegan, Xiaowei Jiang, Julian L. Leibowitz, Stuart J.D. Neil, Tim Skern, Susan R. Weiss, Michael Worobey, Kristian G. Andersen, Robert F. Garry, Andrew Rambaut



PII: S0092-8674(21)00991-0

DOI: https://doi.org/10.1016/j.cell.2021.08.017

Reference: CELL 12163

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The animal origin of SARS-CoV-2



Spyros Lytras¹, Wei Xia², Joseph Hughes¹, Xiaowei Jiang³, David L. Robertson¹

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