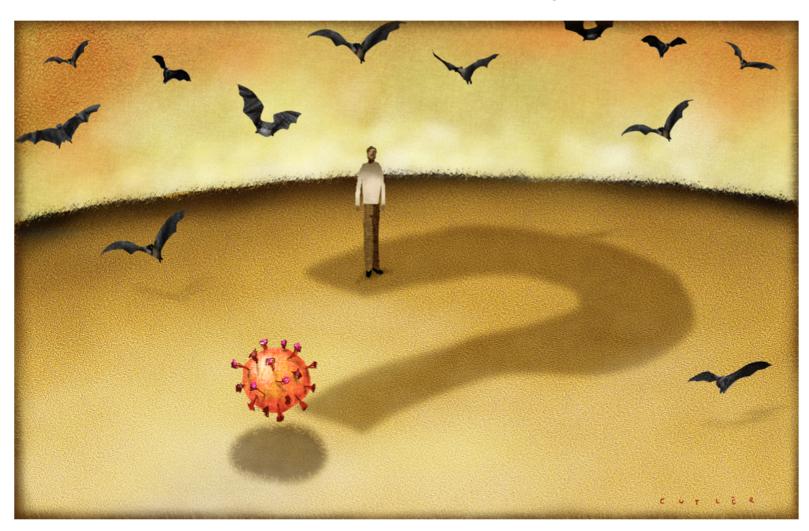




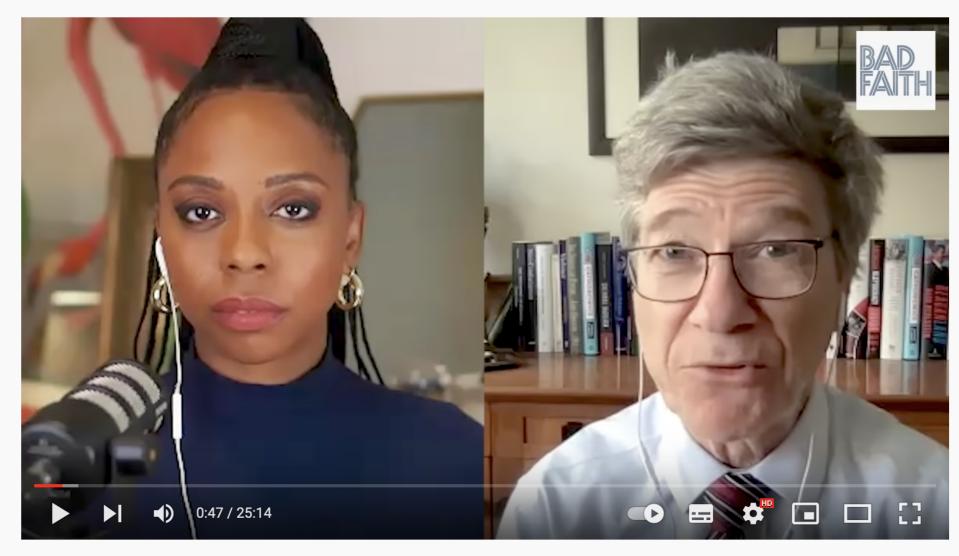
### A call for an independent inquiry into the origin of the SARS-CoV-2 virus

Neil L. Harrison<sup>a,b,1</sup> and Jeffrey D. Sachs<sup>c,1</sup>

19 May







Lab Leak Theory: New Evidence? (w/ Jeffery Sachs)

23 396 vues Sortie le 30 mai 2022 This week, Briahna spoke to Jeffery Sachs, who is heading the

BIOTECHNOLOGY, COMMENTARIES, HEALTH MAY 23, 2022

### Fauci's COVID Origin SWAT Team Versus the Mojiang Miner Passage Theory



by Jonathan Latham, PhD



— US EDITION OF THE WORLD'S OLDEST MAGAZINE

CHINA

### The Harvard connection

Was a Fauci-endorsed Chinese donation part of the lab-leak cover up?

May 26, 2022 | 2:37 pm ■ FROM THE MAGAZINE







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# Mutation signature of SARS-CoV-2 variants raises questions to their natural origins.

n Arakawa, Hiroshi

SARS-CoV-2 variants, such as Omicron, have acquired a number of mutations. These novel mutations have raised concerns regarding the continued efficacy of the antibodies generated in response to vaccination. In this study I show that the evolution of the SARS-CoV-2 variant Omicron, does not appear to follow a Darwinian trajectory. The mutations in the Omicron variant are heavily biased toward nonsynonymous substitutions rather than synonymous substitutions. Interestingly, mutations affecting the spike gene of the Omicron variant are almost exclusively nonsynonymous. Moreover, a nonsynonymous substitution bias within spike gene is a common feature of all of the SARS-CoV-2 variants assessed in this study. This mutational signature is a counter-evidence of neutral evolution, and shows that the spike genes of these SARS-CoV-2 variants have evolved without trial-and-error by mutation and selection. Thus I postulate that the spike genes of these SARS-CoV-2 variants are not the result of natural evolution but rather designed molecules.





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Home / Newsroom / Article / WHO announces proposed members of its Technical Advisory Group on Biosafe

# WHO announces proposed members of its Technical Advisory Group on Biosafety (TAG-B) members

3 June 2022 | Call for consultation



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Wen Wang, Jun-Hua Tian, Xiao Chen, Rui-Xue Hu, Xian-Dan Lin, Yuan-Yuan Pei, Jia-Xin Lv, Jiao-Jiao Zheng, Fa-Hui Dai, Zhi-Gang Song ... Show more

Author Notes

Virus Evolution, veac046, https://doi.org/10.1093/ve/veac046

Published: 04 June 2022 Article history ▼





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# Preliminary Report for the Scientific Advisory Group for the Origins of Novel Pathogens (SAGO)

9 June 2022 | COVID-19: Scientific briefs

"So far neither the virus progenitors nor the natural/intermediate hosts or spill-over event to humans have been identified."

"The SAGO notes that there has not been any new data made available to evaluate the laboratory as a pathway of SARS-CoV-2 into the human population and recommends further investigations into this and all other possible pathways. The SAGO will remain open to any and all scientific evidence that becomes available in the future to allow for comprehensive testing of all reasonable hypotheses."

www.nytimes.com /2022/06/09/science/covid-origins-who-report-china.html

### Mysteries Linger About Covid's Origins, W.H.O. Report Says

Benjamin Mueller, Carl Zimmer ∷ 6/9/2022





The Washington Post Democracy Dies in Darkness

World

# WHO covid origins report says 'lab leak' theory needs further investigation



By Adam Taylor

June 9, 2022 at 10:00 a.m. EDT

#### Foreign Ministry Spokesperson Zhao Lijian's Regular Press Conference on June 10, 2022

2022-06-10 20:59



"China's position on the study of the origins of SARS-CoV-2 is consistent and clear. We always support and participate in science-based global origins-tracing.

"China has shared more data and research findings on SARS-CoV-2 origins study than any other country. This fully demonstrates China's open, transparent and responsible attitude and its support for the work of the WHO and SAGO."