Perspective Virology—The next fifty years

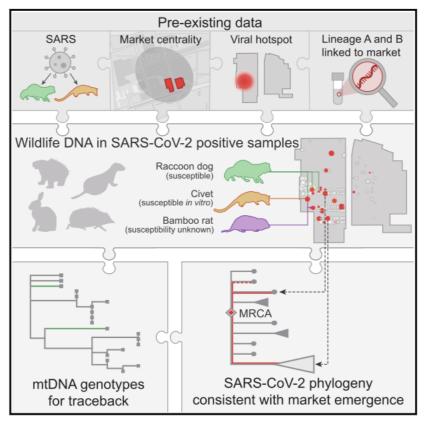
Edward C. Holmes,^{1,2,*} Florian Krammer,^{3,4,5,7,8} and Felicia D. Goodrum^{6,8} ¹School of Medical Sciences, The University of Sydney, Sydney, NSW 2006, Australia ²Laboratory of Data Discovery for Health Limited, Hong Kong SAR, China ³Department of Microbiology, Icahn School of Medicine at Mount Sinai, New York, NY, USA ⁴Center for Vaccine Research and Pandemic Preparedness (C-VaRPP), Icahn School of Medicine at Mount Sinai, New York, NY, USA ⁵Department of Pathology, Molecular and Cell Based Medicine, Icahn School of Medicine at Mount Sinai, New York, NY, USA ⁶Department of Immunobiology, BIO5 Institute, University of Arizona, Tucson, AZ, USA ⁷Ignaz Semmelweis Institute, Interuniversity Institute for Infection Research, Medical University of Vienna, Vienna, Austria ⁸These authors contributed equally *Correspondence: edward.holmes@sydney.edu.au https://doi.org/10.1016/j.cell.2024.07.025

Virology needs to stand firm in the face of these assaults but also to learn important lessons. Virology, like any science, cannot function without public support. We must better communicate our work and acknowledge where failings occur. Most



Genetic tracing of market wildlife and viruses at the epicenter of the COVID-19 pandemic

Graphical abstract



Authors

Alexander Crits-Christoph, Joshua I. Levy, Jonathan E. Pekar, ..., Kristian G. Andersen, Michael Worobey, Florence Débarre

Correspondence

andersen@scripps.edu (K.G.A.), worobey@arizona.edu (M.W.), florence.debarre@sorbonne-universite.fr (F.D.)

In brief

DNA of mammalian wildlife species susceptible to SARS-CoV-2 was detected along with SARS-CoV-2 and other viruses in environmental samples from animal stalls located at the market epicenter of the emergence of COVID-19.

DNA from raccoon dogs, civets, and other wildlife species detected in market samples (but no progenitor virus)

Genotypes of potential hosts reconstructed for retracing animal geographic origins

HailOnline

It WASN'T a lab leak! Scientists say they've finally discovered the truth about the origins of Covid

By LUKE CHAFER

PUBLISHED: 16:09 GMT, 19 September 2024 UPDATED: 10:46 GMT, 20 September 2024



Scientists believe they've uncovered the true origin of the Covid pandemic — and have rejected the popular theory that the virus was leaked from a laboratory.

A major international study published today insists that the virus began life in a 'wet market' in **Wuhan**, **China**, rather than in a scientific experiment.

Biosafety Now

()

Crits-Christoph et al., 2024 Retraction Request



BRYCE NICKELS SEP 19, 2024

"Phylogenomic evidence, epidemiological evidence, and documentary evidence all indicate that SARS-CoV-2 entered humans in July-November 2019 (..). As such, conclusions based on data for the Huanan Seafood Market on or after mid-to-late December 2019--as in Crits-Christoph et al. 2024--cannot, even in principle, shed light on spillover into humans that occurred one to five months earlier, in July-November, 2019 (4,29)."

 The Telegraph
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Matt Ridley Don't fall for the Chinese wet market theory for Covid

Contrary to reports, the evidence still points to a lab leak being the most likely cause of the pandemic

https://www.telegraph.co.uk/news/2024/09/21/dont-believe-the-chinese-wet-market-theory-for-covid/

22 Sept 2024



Scientists now believe they have more evidence that Covid-19 originated in a food market in Wuhan. Norwegian experts are skeptical.

"There are some researchers who have a great interest in rejecting the lab leak theory (..) Again they are trying to close the whole discussion about the lab leak. The only people who are absolutely certain about the origin of the virus are this group of scientists."

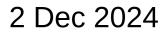
https://www-vg-no.translate.goog/nyheter/i/63BQaL/ny-studie-hevder-at-covid-19-kom-fra-marked?_x_tr_sl=no&_x_tr_hl=no&_x_tr_pto=wapp

7 Oct 2024

Preparing for the Next Pandemic: Evolution, Pathogenesis and Virology of Coronaviruses, AWAJI, JAPAN

Awaji Yumebutai Conference Center, JAPAN

December 2-5, 2024 Abstract deadline: October 11, 2024 Organized by: Kei Sato, The University of Tokyo Yunlong Cao, Peking University Ravindra Gupta, University of Cambridge Zhengli Shi, Guangzhou Laboratory (Virtual) Alex Sigal, Africa Health Research Institute





Report by the Republican-majority select committee of the US House of Representatives

About 🐣 Committee Activity 🎽



Published: Dec 2, 2024

FINAL REPORT: COVID Select Concludes 2-Year Investigation, Issues 500+ Page Final Report on Lessons Learned and the Path Forward

"COVID-19 most likely emerged from a laboratory in Wuhan"

https://oversight.house.gov/release/final-report-covid-select-concludes-2-year-investigation-issues-500-page-final-report-on-lessons-learned-and-th e-path-forward/

nature

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NEWS 04 December 2024

Sick animals suggest COVID pandemic started in Wuhan market

A preliminary analysis takes a closer look at genomic data collected at the market.

By Smriti Mallapaty

The analysis finds evidence that the immune systems of some animals had been activated, which supports the theory that they were infected, says Stanley Perlman, a virologist at the University of Iowa in Iowa City. But "it doesn't substitute for finding the virus in an infected animal".



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Gen Access Deer-Reviewed

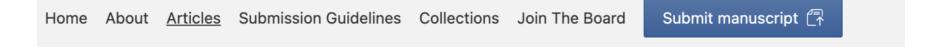
Receptor binding and structural basis of raccoon dog ACE2 binding to SARS-CoV-2 prototype and its variants

Chunliang Luo, Linjie Li, Yuhang Gu, Hangchuan Zhang, Zepeng Xu, Junqing Sun, Kaiyuan Shi, Sufang Ma, Wen-Xia Tian, Kefang Liu 🔟, George F. Gao 🔯

Published: December 5, 2024 • https://doi.org/10.1371/journal.ppat.1012713

"Some scientists raised the possibility that raccoon dogs could serve as natural hosts for SARS-CoV-2 [33, 34]. However, this finding alone falls short of conclusively establishing raccoon dogs as natural hosts. To date, while 45 species have been documented as naturally infected with SARS-CoV-2, there has been no confirmation of raccoon dogs being naturally infected with or carrying the virus. Furthermore, investigations during the early stages of the COVID-19 epidemic failed to detect SARS-CoV-2 in raccoon dogs [35]. Thus, there is currently no direct evidence supporting that raccoon dogs are the natural hosts of SARS-CoV-2."

BMC Genomic Data



Data Note Open access Published: 17 December 2024

Pre-pandemic artificial MERS analog of polyfunctional SARS-CoV-2 S1/S2 furin cleavage site domain is unique among spike proteins of genus *Betacoronavirus*

Andreas Martin Lisewski

BMC Genomic Data 25, Article number: 104 (2024) Cite this article

The furin cleavage site "includes two additional functional motifs: a pat7 nuclear localization signal and two flanking O-glycosites."

"a fully analogous S1/S2 domain was found in only one other virus: the artificial MERS infectious clone MERS-MA30, described already in 2017"

Retraction Note: Origin and cross-species transmission of bat coronaviruses in China

Alice Latinne, Ben Hu, Kevin J. Olival, Guangjian Zhu, Libiao Zhang, Hongying Li, Aleksei A. Chmura, Hume E. Field, Carlos Zambrana-Torrelio, Jonathan H. Epstein, Bei Li, Wei Zhang, Lin-Fa Wang, Zheng-Li Shi 🖾 & Peter Daszak 🖾

Nature Communications 15, Article number: 10706 (2024) Cite this article

Article Open access Published: 19 December 2024

https://www.nature.com/articles/s41467-024-55454-w

Origin and cross-species transmission of bat coronaviruses in China

Alice Latinne, Ben Hu, Kevin J. Olival, Guangjian Zhu, Li-Biao Zhang, Hongying Li, Aleksei A. Chmura, Hume E. Field, Carlos Zambrana-Torrelio, Jonathan H. Epstein, Bei Li, Wei Zhang, Lin-Fa Wang, Zheng-Li Shi ⊠ & Peter Daszak ⊠ Laotian and duplicate sequences removed

Nature Communications 15, Article number: 10705 (2024) Cite this article

https://www.nature.com/articles/s41467-024-55384-7

Why?



Gilles Demaneuf 🤣 @gdemaneuf



Because they could have been forced to retract if they had kept these samples in their data, for reasons linked to the way the Laos samples were collected.

10:29 AM · Dec 21, 2024 · 278 Views



30 December 2024 | Statement |Reading time: 1 min (282 words)

"We continue to call on China to share data and access so we can understand the origins of COVID-19. This is a moral and scientific imperative."

https://www.who.int/news/item/30-12-2024-milestone-covid-19-five-years-ago





SCIENCEINSIDER HEALTH

COVID 5 years later: Learning from a pandemic many are forgetting

Five years after SARS-CoV-2 surfaced, scientists reflect and look ahead to the next threat

2 JAN 2025 · 9:00 AM ET · BY JON COHEN

"In Japan, a 4-day conference" (..)

"two of the country's most prominent COVID-19 researchers were notable no shows. Shi Zhengli (..) only gave a prerecorded video talk (..) despite being a co-organizer of the event. So did Chinese Academy of Sciences virologist George Gao, former head of China's Center for Disease Control and Prevention. Shi, now with the Guangzhou Laboratory, and Gao both declined to explain their absence to Science."



About Y Committee A



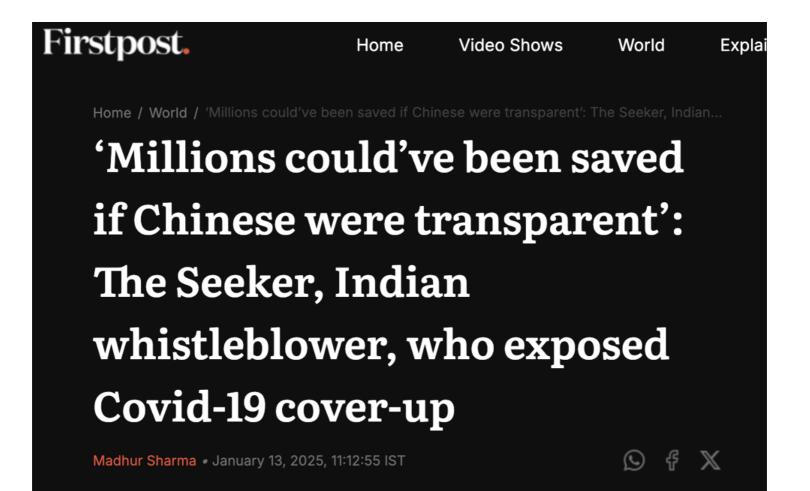
Published: Jan 17, 2025

BREAKING: HHS Formally Debars EcoHealth Alliance, Dr. Peter Daszak After COVID Select Reveals Pandemic-Era Wrongdoing

"the U.S. Department of Health and Human Services (HHS) cut off all funding and formally debarred EcoHealth Alliance Inc. (EcoHealth) and its former President, Dr. Peter Daszak, for five years"

"This letter also confirms that EcoHealth terminated Dr. Daszak's employment effective January 6, 2025."

Interview of The Seeker



Prasenjit Ray and others at the collective DRASTIC spent months scouring the

https://www.firstpost.com/world/millions-could-have-been-saved-if-chinese-were-transparent-the-seeker-indian-whistleblower-who-exposed-covid-19 -cover-up-13852240.html

NR The CIA 'Now Favors the Lab Theory' For Covid-19's Origin

THE CORNER NATIONAL SECURITY & DEFENSE

The CIA 'Now Favors the Lab Theory' For Covid-19's Origin

"the agency issued a new assessment this week, with analysts saying they now favor the lab theory"

https://www.nationalreview.com/corner/the-cia-now-favors-the-lab-theory-for-covid-19s-origin/

MONDAY, JANUARY 27, 2025

DR. PAUL ISSUES SUBPOENAS TO FOURTEEN AGENCIES REGARDING COVID-19 ORIGINS AND RISKY GAIN-OF-FUNCTION RESEARCH

https://www.hsgac.senate.gov/media/reps/dr-paul-issues-subpoenas-to-fourteen-agencies-regarding-covid-19-origins-and-risky-gain-of-function-research/

https://www.washingtonexaminer.com/policy/healthcare/3302227/rand-paul-subpoenas-14-agencies-covid-origins/

Australia News > Coronavirus

EXCLUSIVE

Former US State Department COVID investigator reveals Australian spy agencies collected intelligence related to COVID origins in China

Australian spy agencies collected top-secret intelligence related to the origins of COVID-19 in Wuhan, China, a leading investigator has revealed.



Sharri Markson Sky News Host

less than 2 min read January 28, 2025 - 10:30PM sky news .com.au

https://www.skynews.com.au/australia-news/coronavirus/former-us-state-department-covid-investigator-reveals-australian-spy-agencies-collected-in telligence-related-to-covid-origins-in-china/news-story/daac6c683b87b789df1c85d564f4d286

https://www.zerohedge.com/medical/australian-spy-agency-collected-signals-intelligence-related-covid-china-origins-former

22 Feb 2025



Volume 11, Issue 1 2025 (In Progress)

JOURNAL ARTICLE

Recently reported SARS-CoV-2 genomes suggested to be intermediate between the two early main lineages are instead likely derived a

Jonathan E Pekar ➡, Niema Moshiri, Philippe Lemey, Alexander Crits-Christoph, Florence Débarre, Stephen A Goldstein, Zach Hensel, Andrew Rambaut, Michael Worobey, Edward C Holmes, Joel O Wertheim ➡

Virus Evolution, Volume 11, Issue 1, 2025, veaf008, https://doi.org/10.1093/ve/veaf008 **Published:** 22 February 2025 Article history •

"these two T/T genomes represent an evolutionarily intermediate haplotype between lineages A and B, or if they are a later descendent of either of these two lineages."

"After including additional data from the start of the pandemic, including these two T/T genomes, we again find a discrepancy in the molecular clock when inferring the ancestral haplotype of SARS-CoV-2, corroborating existing evidence for the separate introductions of SARS-CoV-2 lineages A and B into the human population in late 2019."

27 Feb 2025

[Submitted on 27 Feb 2025 (this version), latest version 14 Mar 2025 (v2)]

Purported quantitative support for multiple introductions of SARS-CoV-2 into humans is an artefact of an imbalanced hypothesis testing framework

Angus McCowan

A prominent report claimed substantial support for two introductions of SARS-CoV-2 into humans using a calculation that combined phylodynamic inferences and epidemic models. Inspection of the calculation identifies an imbalance in the hypothesis testing framework that confounds this result; the single-introduction model was tested against more stringent conditions than the two-introduction model. Here, I show that when the two-introduction model is tested against the same conditions, the support disappears.

https://arxiv.org/abs/2502.20076v1

See also critics by Florence Débarre on PubPeer and responses

https://pubpeer.com/publications/2C73F441513A42FE6D566147E5E76C#4





Access provided by Institute of Biological Sciences

ARTICLE · Online now, February 18, 2025

Bat-infecting merbecovirus HKU5-CoV lineage 2 can use human ACE2 as a cell entry receptor

 $\underbrace{ \text{Jing Chen}^{1,8} \cdot \text{Wei Zhang}^{2,3,8} \cdot \text{Yang Li}^{1,8} \cdot \text{Chen Liu}^{4,8} \cdot \text{Tianyi Dong}^{1,5} \cdot \text{Huiyu Chen}^{2,3} \cdot \text{Chunguang Wu}^{1,5} \cdot \text{Jia Su}^{1,5} \cdot \text{Bei Li}^{1} \cdot \text{Wei Zhang}^{1} \cdot \text{Ben Hu}^{1} \cdot \text{Jingkun Jia}^{1,5} \cdot \text{Cheng-Bao Ma}^{4} \cdot \text{Yan Zhu}^{1} \cdot \text{Xiangyang He}^{6} \cdot \text{Ang Li}^{2,3} \cdot \text{Kaiyi Pan}^{1,5} \cdot \text{Haofeng Lin}^{3} \cdot \text{Zishuo Guo}^{1,5} \cdot \text{Cong Li}^{7} \cdot \text{Libiao Zhang}^{6} \boxtimes \cdot \text{Huan Yan}^{6} \oplus \text{Peng Zhou}^{2,3} \boxtimes \cdot \text{Wei Peng}^{2,3} \boxtimes \cdot \text{Zheng-Li Shi}^{2,9} \boxtimes \text{Lishuo Shang}^{2,9} \boxtimes \text{Li Shi}^{2,9} \boxtimes \text{Libiao Zhang}^{6} \oplus \text{Liu}^{1,5} \cdot \text{Libiao Zhang}^{2,6} \oplus \text{Liu}^{1,5} \cdot \text{Libiao}^{2,9} \boxtimes \text{Liu}^{2,3} \oplus \text{Liu}^{2,3} \oplus \text{Liu}^{2,3} \oplus \text{Liu}^{2,3} \oplus \text{Li}^{2,9} \boxtimes \text{Libiao}^{2,9} \oplus \text{Liu}^{2,3} \oplus \text{Liu}^$

discovery of HKU5-CoV lineage 2 (HKU5-CoV-2) in bats

Bat HKU5-CoV-2 was isolated and used to infect human-ACE2-expressing cells Experiments "performed in BSL-2"

27 Feb 2025



Exclusive: Inside the FBI's Lab Leak Investigation

As bird flu spreads, and Team Trump begins dismantling America's public health apparatus, a former FBI scientist and investigator speaks out about the evidence that led the bureau to suspect that COVID-19 was sparked by an incident at the Wuhan Institute of Virology.

BY KATHERINE EBAN

FEBRUARY 27, 2025

https://www.vanityfair.com/news/story/inside-the-fbis-lab-leak-investigation





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New Results

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The furin cleavage site is required for pathogenesis, but not transmission of SARS-CoV-2

Angelica L. Morgan, Michelle N. Vu, D Yiyang Zhou, Kumari G. Lokugamage, William M. Meyers, R. Elias Alvarado, Yani Ahearn, Leah K. Estes, Jessica A. Plante, Bryan A. Johnson, D Mehul S. Suthar, David H Walker, Ken S. Plante, Vineet D. Menachery

Here: SARS-CoV-2 PQQAR mutant disrupting the FCS, but preserving the S1/S2 loop length attenuates pathogenesis

can be transmitted between hamsters but not as good as WT

Corona Pandemic: BND Believes a Laboratory Accident Was the Cause

https://www.spiegel.de/wissenschaft/corona-lothar-wieler-haelt-labor-ursprung-von-sars-cov-2-fuer-die-wahrscheinlichere-these-a-1e2bcf76-2bfa-4fb4-b3bb-a9a0ed6ac606?sar ef=re-xx-cp-sh

https://www.nzz.ch/international/der-deutschen-regierung-liegen-plausible-indizien-vor-dass-das-coronavirus-aus-dem-labor-in-wuhan-kommt-ld.1874840

https://www.zeit.de/2025/11/coronavirus-ursprung-wuhan-labor-china-bnd/seite-4

https://www.zeit.de/politik/deutschland/2025-03/corona-pandemie-bundestag-laborthese-bundesregierung-bnd

https://www.zeit.de/gesundheit/2025-03/ursprung-sars-cov-2-bnd-labor-china-corona

https://www.zeit.de/2025/11/coronavirus-ursprung-wuhan-labor-china-bnd/seite-4

https://www.zeit.de/politik/ausland/2025-03/coronavirus-bnd-china-labor-ursprung-who

https://www.zeit.de/politik/deutschland/2025-03/merkel-corona-wuhan-laborunfall-vertuschung

MailOnline

Boris was convinced by my evidence but the scientists swallowed Beijing's propaganda on the origin of the Coronavirus, writes SIR RICHARD DEARLOVE

By SIR RICHARD DEARLOVE FOR THE MAIL ON SUNDAY PUBLISHED: 22:29 GMT, 15 March 2025 UPDATED: 22:35 GMT, 15 March 2025

www.dailymail.co.uk/news/article-14503019/Boris-evidence-scientists-Beijing-propaganda-Coronavirus-SIR-RICHARD-DEARLOVE.html

Versus this old article: *Oct 2023*



Top science journal faced secret attacks from Covid conspiracy theory group

"The campaign was led by former chief of the Secret Intelligence Service (SIS) Sir Richard Dearlove in conjunction with (..)"

https://www.computerweekly.com/news/366553435/Top-science-journal-faced-secret-attacks-from-Covid-conspiracy-theory-group

And : https://www.dailymail.co.uk/news/article-8389809/Spy-chief-coronavirus-storm.html June 2020

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Q&A

They Tried to Silence Her COVID Origin Theory. Now Even the CIA Agrees with Alina Chan.

In 2020, the Broad Institute scientist ignited a controversy by suggesting that SARS-CoV-2 originated in a Chinese lab. Five years later, it sure looks like she was right.

by ROWAN JACOBSEN • 3/12/2025, 9:00 a.m.



One of the first to suggest Covid was created in a lab, biologist Alina Chan faced death threats and was branded a 'race traitor'. Now she tells IAN BIRRELL... 'There was a real conspiracy among very powerful scientists. The cover-up was morally repugnant'

By IAN BIRRELL FOR THE DAILY MAIL PUBLISHED: 22:34 GMT, 16 March 2025 UPDATED: 22:40 GMT, 16 March 2025

https://www.dailymail.co.uk/news/article-14505321/Covid-created-lab-biologist-Alina-Chan-death-threats-race-traitor-IAN-BIRRELL-real-conspiracy-powerful-scientists-morally-repugnant.html

Ehe New York Eimes

OPINION ZEYNEP TUFEKCI

We Were Badly Misled About the Event That Changed Our Lives

March 16, 2025

https://www.nytimes.com/2025/03/16/opinion/covid-pandemic-lab-leak.html

All my introduction slides are available online https://courtier.ijm.fr/news-covid-origin/