



E-mails: Fauci, Collins Knew Top Scientists Favor Lab-leak for Origin of China Virus. Truth Debunked as “Conspiracy Theory.”

Early this week, House Republicans released e-mails to and from Anthony Fauci and his boss, Francis Collins, heads of the National Institute of Allergy and Infectious Disease and National Institute for Health.

Those e-mails again proved Fauci lied about U.S. subsidies that paid for potentially catastrophic gain-of-function virus experiments in Red China. Those experiments turn naturally-occurring viruses into super-pathogens that can infect people.

Disclosed by Representatives James Comer of Kentucky and Jim Jordan of Ohio, the cybermissives also reveal something else: The two men were well aware that top scientists believed the SARS-CoV-2 pathogen leaked from the [poorly run lab](#).

Later, they lied about that, too. They called the idea a “conspiracy theory.” As did the leftist mainstream media.

AP Images
Anthony Fauci

The E-mails

It wasn't a “conspiracy theory,” and they well knew it, as [The New American reported](#) about e-mails released in June. An e-mail dated February 1, 2020 showed that a top scientist told Fauci the virus was “potentially engineered.” They also showed that Peter Daszak — the head of EcoHealth, [which funneled](#) the forbidden gain-of-function subsidies to Wuhan — thanked Fauci for helping “dispel the myth” that mad scientists in China created the super bug.

The [e-mails disclosed](#) this week show the same thing. Dated February 2-4, 2020, they followed a conference call between Fauci, Collins, Jeremy Farrar of the Wellcome Trust, and others.

The excerpts show Fauci and Collins knew the virus likely sprung from a lab, and that Collins was extremely concerned about keeping that information under wraps.

Farrar wrote to Fauci, Collins, and Lawrence Tabak, principal deputy director at NIH, with “some thoughts overnight from others,” meaning other scientists:

He is bothered by the furin site and has a hard time explain[ing] that as an event outside the lab (though, there are possible ways in nature, but highly unlikely)....

So, I think it becomes a question of how do you put all this together, whether you believe in this series of coincidences, what you know of the lab in Wuhan, how much could be in



Written by [R. Cort Kirkwood](#) on January 13, 2022

nature — accidental release or natural event? I am 70:30 or 60:40.

Farrar E-mail

From Mike Farzan (discoverer of SARS receptor):

1. The RBD didn't look 'engineered' to him – as in, no human would have selected the individual mutations and cloned them into the RBD (I think we all agree)
2. Tissue culture passage can often lead to gain of basic sites – including furin cleavage sites (this is stuff they have seen with human coronaviruses)
3. He is bothered by the furin site and has a hard time explain that as an event outside the lab (though, there are possible ways in nature, but highly unlikely)
4. Instead of directed engineering, changes in the RBD and acquisition of the furin site would be highly compatible with the idea of continued passage of virus in tissue culture
5. Acquisition of the furin site would likely destabilize the virus but would make it disseminate to new tissues.

So, given above, a likely explanation could be something as simple as passage SARS-live CoVs in tissue culture on human cell lines (under BSL-2) for an extended period of time, accidentally creating a virus that would be primed for rapid transmission between humans via gain of furin site (from tissue culture) and adaption to human ACE2 receptor via repeated passage.

...So, I think it becomes a question of how do you put all this together, whether you believe in this series of coincidences, what you know of the lab in Wuhan, how much could be in nature – accidental release or natural event? I am 70:30 or 60:40.

From Bob [Garry]:

Before I left the office for the ball, I aligned nCoV with the 96% bat CoV sequenced at WIV. Except for the RBD the S proteins are essentially identical at the amino acid level – well all but the perfect insertion of 12 nucleotides that adds the furin site. S2 is over its whole length essentially identical. I really can't think of a plausible natural scenario where you get from the bat virus or one very similar to it to nCoV where you insert exactly 4 amino acids 12 nucleotide that all have to be added at the exact same time to gain this function – that and you don't change any other amino acid in S2? I just can't figure out how this gets accomplished in nature. Do the alignment of the spikes at the amino acid level – its stunning. Of course, in the lab it would be easy to generate the perfect 12 base insert that you wanted. Another scenario is that the progenitor of nCoV was a bat virus with the perfect furin cleavage site generated over

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In other words, the scientist was almost certain the virus was created in and sprung from the lab in Wuhan.

"I really can't think of a plausible natural scenario where you get from the bat virus or one very similar to it to nCoV," another top scientist said, "where you insert exactly 4 amino acids 12 nucleotide that all have to be added at the exact same time to gain this function — that and you don't change any other amino acid in S2? I just can't figure out how this gets accomplished in nature."

Collins, who left NIH at the end of 2021, wrote back to say he was worried about conspiracy theorists running amok with the truth.



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“I share your view that a swift convening of experts in a confidence inspiring framework (WHO seems really the only option) is needed, or the voices of conspiracy will quickly dominate, doing great potential harm to science and international harm.”

2. Email from Dr. Francis Collins to Drs. Jeremy Farrar, Anthony Fauci, and Lawrence Tabak

From: Francis Collins <>
Date: Sunday, 2 February 2020 at 10:27
To: Jeremy Farrar <>
Cc: "Fauci, Anthony (NIH/NIAID) [E]" <>, "Tabak, Lawrence (NIH/OD) [E]" <>
Subject: RE: Teleconference

Jeremy,



I'm available any time today except 3:15 – 5:45 pm EST (on a plane) for a call to Tedros. Let me know if I can help get through his thicket of protectors.

Francis

... Though the arguments from Ron Fouchier and Christian Drosten are presented with more forcefulness than necessary, I am coming around to the view that a natural origin is more likely. But I share your view that a swift convening of experts in a confidence inspiring framework (WHO seems really the only option) is needed, or the voices of conspiracy will quickly dominate, doing great potential harm to science and international harmony...

Fouchier E-mail

activities of the scientific community, WHO and other stakeholders at present. It is my opinion that a non-natural origin of 2019-nCoV is highly unlikely at present. Any conspiracy theory can be approached with factual information.

... An accusation that nCoV-2019 might have been engineered and released into the environment by humans (accidental or intentional) would need to be supported by strong data, beyond a reasonable doubt. It is good that this possibility was discussed in detail with a team of experts. However, further debate about such accusations would unnecessarily distract top researchers from their active duties and do unnecessary harm to science in general and science in China in particular.

Another virologist explained that “debate about such accusations would unnecessarily distract top researchers from their active duties and do unnecessary harm to science in general and science in China in particular.”

China, of course, was interested in the gain-of-function experiments because they could produce a viral bioweapon, as [The New American reported](#).

In April, Collins asked what could be done to stop the truth from emerging:

Wondering if there is something NIH can do to help put down this very destructive conspiracy, with what seems to be growing momentum:



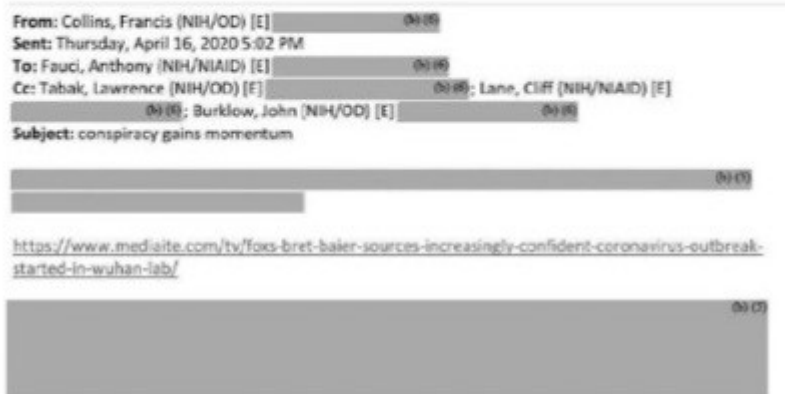
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I hoped the Nature Medicine article on the genomic sequence of SARS-CoV-2 would settle this.

Collins was referring to [an article](#) that he and Fauci [helped prepare](#) to “debunk” the “conspiracy theory.”

Government Official Attempts to Stifle the Lab Leak Hypothesis

1. Email from Dr. Francis Collins to Drs. Anthony Fauci, Lawrence Tabak, Cliff Lane, and Mr. John Burklow



Francis

Wondering if there is something NIH can do to help put down this very destructive conspiracy, with what seems to be growing momentum:

<https://www.mediaite.com/tv/foxs-bret-baier-sources-increasingly-confident-coronavirus-outbreak-started-in-wuhan-lab/>

I hoped the Nature Medicine article on the genomic sequence of SARS-CoV-2 would settle this. But probably didn't get much visibility.

Anything more we can do? Ask the National Academy to weigh in?

Francis

Lab Leak

In October, answering an inquiry from Comer, [Tabak confirmed](#) that Fauci lied when he repeatedly testified before a Senate committee that his agency did not subsidize gain-of-function research in Wuhan. If he didn't lie, then Fauci, 80, simply doesn't know what his agency is doing.

Richard Ebright, head of the Waksman Institute of Microbiology at Rutgers University, was unequivocal about what Fauci, Collins, and other pushed out to protect themselves:

“The NIH — specifically, Collins, Fauci, and Tabak — lied to Congress, lied to the press, and lied to the public,” he tweeted. “Knowingly. Willfully. Brazenly.”

The NIH received the relevant documents in 2018 and reviewed the documents in 2020 and again in 2021.



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The NIH—specifically, Collins, Fauci, and Tabak—lied to Congress, lied to the press, and lied to the public. Knowingly. Willfully. Brazenly.

— Richard H. Ebright (@R_H_Ebright) [October 20, 2021](#)

H/T: [The Telegraph](#)





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