



Evidence of Covid Pandemic and Bioweapon Cover-Ups all leads back to Metabiota, Peter Daszak and Hunter Biden

Description

As evidence of a potential bioweapons cover-up has started emerging, a company called Metabiota is gaining prominence. The links between Metabiota and several key players in the COVID pandemic and/or the Ukraine labs story are manifold, so there's no really simple way to unravel it in a logical sequence. That said, let's start with what Metabiota does and the connections of its founder, and expand from there.

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Metabiota's Mission

Metabiota's mission is to make the world more resilient to epidemics by providing "data, analytics, advice and training to prepare for global health threats and mitigate their impacts."¹

Through data analysis, they help "decision makers across government and industry" to estimate and mitigate pandemic risks. But they also claim to support "sustainable development," which seems to have little to do with pandemic risk management.

That term, "sustainable development," is one promoted by Klaus Schwab, founder of the World Economic Forum (WEF). It's part and parcel of Schwab's plan for a global Great Reset and transhumanist revolution (aka, the Fourth Industrial Revolution).

It's not surprising, then, to find out that the founder of Metabiota, Nathan Wolfe, not only has close ties to the WEF, but is also a rising star there. He's a WEF Young Global Leader graduate and was awarded the WEF's Technology Pioneer award in 2021.

Metabiota and the Search for Pandemic Viruses

Metabiota was a core partner of a United States Agency for International Development's (USAID) Pandemic Threat Program called PREDICT, which sought to identify viruses with pandemic potential.

Contractors funded through this program have included the EcoHealth Alliance, headed by Peter Daszak. The PREDICT program, directed by Dennis Carroll, appears to have served as a proof of concept for the Global Virome Project that Carroll founded.

According to a recent investigation by U.S. Right to Know (USRTK),² Carroll appears to have diverted government funds from the PREDICT program while he was still running it, to fund this personal side project, which was set up with the intention to collect, identify and catalogue 1 million viruses from wildlife in an effort to predict which ones might cause a human epidemic.

Metabiota's Funding

Metabiota receives funding from several interconnected organizations and agencies, including:³

- Pilot Growth Management, cofounded by Neil Callahan. Callahan is also a cofounder of Rosemont Seneca Technology Partners, and he sits on Metabiota's board of advisers
- The Global Virome Project, which reportedly paid (or was planning to pay) Metabiota \$341,000 to conduct a cost-benefit analysis⁴
- In-Q-Tel, a CIA venture capital firm that specializes in high-tech investments that support or benefit the intelligence capacity of U.S. intelligence agencies
- The U.S. Department of Defense's Threat Reduction Agency (DTRA).⁵ Specifically, in 2014, DTRA

awarded Metabiota \$18.4 million in federal contracts for scientific and technical consulting services to the DTRA's labs in Ukraine and Georgia⁶

By outsourcing work to private companies, DTRA is able to circumvent Congressional oversight. Russia is now accusing the U.S. of funding secret and illegal bioweapons research in these Ukraine labs, and claims this was the real reason behind its invasion

- Rosemont Seneca,⁷ an investment fund co-managed by Hunter Biden.⁸ If Russia's accusations turn out to be true, this tie may prove deeply problematic for the White House, as this means the Biden family was more or less directly involved in the funding of that research

Wolfe has also received more than \$20 million in research grants from Google, the NIH and the Bill & Melinda Gates Foundation, just to name a few, and was a friend of now-deceased Jeffrey Epstein. In his 2012 book, "The Viral Storm," Wolfe thanked friends for their support, including Epstein and Boris Nikolic. Nikolic, a biotech venture capitalist, was named "back-up executor" in Epstein's will.⁹

Epstein, who besides being a convicted pedophile and accused child sex trafficker, had a robust interest in eugenics. It's now well-known that he dreamed of creating a "superhuman" race of his own by impregnating dozens of women at a time at his New Mexico ranch.¹⁰ Epstein also managed to secure meetings with Bill Gates,¹¹ whose family history is also marked by an interest in eugenics and population control.

Metabiota's Founder Tied to Suspect in COVID Pandemic

In addition to having close ties to the WEF and its Great Reset agenda, Wolfe, the founder of Metabiota, has also served on the EcoHealth Alliance's editorial board since 2004. In 2017, he even co-wrote a study on coronaviruses in bats together with EcoHealth Alliance president, Peter Daszak.

As you may recall, EcoHealth Alliance, a nonprofit organization focused on pandemic prevention, worked closely with the Wuhan Institute of Virology (WIV) in China, where SARS-CoV-2 is suspected of having originated.¹²

Daszak — who received funding for coronavirus research from the National Institute of Allergy and Infectious Diseases (NIAID), led by Dr. Anthony Fauci, and the U.S. State Department¹³ — subcontracted some of that work to Shi Zheng-li at the WIV. He was also the coauthor on research projects at the WIV.

Once rumors of SARS-CoV-2 being man-made first began, Daszak played a central role in the plot to obscure the lab origin by crafting a scientific statement condemning such inquiries as "conspiracy theory."^{14,15} This manufactured "consensus" was then relied on by the media to counter anyone presenting theories and evidence to the contrary.

This, despite the fact that he, in 2015, warned that a global pandemic might occur from a laboratory incident — and that "the risks were greater with the sort of virus manipulation research being carried out in Wuhan!"¹⁶

In 2021, two investigations into the origins of the COVID pandemic were opened, one by the World Health Organization¹⁷ and another by The Lancet,¹⁸ and Daszak somehow managed to end up on

both of these committees, despite having openly and repeatedly dismissed the possibility of the pandemic being the result of a lab leak.¹⁹

Editor's note: The WHO reference has been scrubbed from both the agency's website and internet archives, but several news stories like this one from NPR,²⁰ published after the investigation was launched, are still live and accessible.

Interestingly, one of EcoHealth Alliance's policy advisers is a former Fort Detrick commander named David Franz. Fort Detrick is the principal U.S. government-run "biodefense" facility, although Franz himself has publicly admitted that "in biology ... everything is dual use — the people, the facilities and the equipment."²¹

Metabiota and the DTRA

In late May 2016, Metabiota hired Andrew C. Weber,²² a member of the Council on Foreign Relations, to head up its Global Partnerships.²³ Between 2009 and 2014, Weber served as assistant secretary of defense for Nuclear, Chemical and Biological Defense under then-president Obama.

Weber is credited with creating the Defense Threat Reduction Agency (DTRA) — a combat support agency within the U.S. DoD, specializing in countering weapons of mass destruction, including biological weapons^{24,25} — and as mentioned earlier, the DTRA has reportedly funded Metabiota to operate U.S.-funded biological research labs in Ukraine.

The DTRA has also issued a number of grants to the EcoHealth Alliance, totaling at least \$37.5 million,^{26,27} including a 2017 grant for \$6.5 million to "understand the risk of bat-borne zoonotic disease emergence in Western Asia."²⁸

According to a December 2020 report by The Defender,²⁹ EcoHealth Alliance had tried to hide most of the Pentagon funding that it had received between 2013 and 2020, most of which came from the DTRA.

Metabiota's Bungled Ebola Response

In 2016, CBS News published a scathing critique of Metabiota's response to the 2014 Ebola epidemic in West Africa.³⁰ Metabiota had been hired by the WHO and the local government of Sierra Leone to monitor the spread of the epidemic, but according to an investigation by The Associated Press, "some of the company's actions made an already chaotic situation worse."

In a July 17, 2014, email obtained by AP, Dr. Eric Bertherat, medical officer at the WHO's Department of Epidemic and Pandemic Alert and Response, complained about misdiagnoses and "total confusion" at the small laboratory Metabiota shared with Tulane University in Kenema, Sierra Leone.

According to Bertherat, there was "no tracking of the samples" and "absolutely no control on what is being done." "This is a situation that WHO can no longer endorse," he wrote. Similarly, Sylvia Blyden, special executive assistant to the president of Sierra Leone, told AP Metabiota's response was a disaster.³¹

“‘They messed up the entire region,’ she said. She called Metabiota’s attempt to claim credit for its Ebola work ‘an insult for the memories of thousands of Africans who have died.’”

U.S. health official Austin Demby, who evaluated Metabiota’s and Tulane’s lab work at the request of the U.S. Centers for Disease Control and Prevention and the government of Sierra Leone, was also critical.

In one email, Demby noted used needles were left out and there was no ultraviolet light for decontamination. The space was also too small to safely process blood samples. “The cross-contamination potential is huge and quite frankly unacceptable,” he wrote.

Anja Wolz, an emergency coordinator with Doctors Without Borders, told AP she witnessed Metabiota workers entering homes of suspected Ebola patients without protective gear, and leaving high-risk areas without performing any kind of decontamination procedure. She also accused Metabiota of miscalculating the severity of the outbreak, while insisting that they had the situation under control when clearly, they didn’t.

Tulane microbiology professor Bob Garry was also critical of Metabiota’s choice to have Dr. Jean-Paul Gonzalez run the operation, as Gonzalez, in 1994, had accidentally gotten infected with a rare hemorrhagic fever while working in a Yale University lab.

He failed to notify anyone about the exposure for more than a week, a delay that put more than 100 other people at risk. Gonzalez was ordered to take a remedial safety course, but according to Garry, such carelessness was a red flag, and he didn’t think Gonzalez was the right man to teach Sierra Leoneans about Ebola.

“Do you really want the person who infected himself with hemorrhagic fever going around explaining to people how to be safe?” Garry asked in an email to a Metabiota media representative. Wolfe defended his company, saying there was no evidence they’d done anything wrong. Some of the problems he blamed on misunderstandings, and others on commercial rivalry.

Lab Accident ‘Most Likely,’ yet Least Probed Cause of COVID

In a March 28, 2022, report,³² U.S. Right to Know (USRTK) revealed the contents of a 2020 State Department memo³³ obtained by the group. USRTK writes:³⁴

“‘Origin of the outbreak: The Wuhan labs remained the most likely but least probed,’ reads the topline. The memo is written as a BLUF — ‘bottom line up front’ — a style of communication used in the military. The identity of the author or authors is unknown ...

‘BLUF: There is no direct, smoking gun evidence to prove that a leak from Wuhan labs caused the pandemic, but there is circumstantial evidence to suggest such is the case,’ the memo reads. Apparently drafted in spring 2020, the memo details circumstantial evidence for the ‘lab leak’ theory — the idea that COVID-19 originated at one of the labs in Wuhan, China, the pandemic’s epicenter.

The memo raises concerns about the ‘massive amount’ of research on novel coronaviruses apparently conducted at the Wuhan Institute of Virology and the nearby Wuhan Center for Disease Control lab ... The memo also flags biosafety lapses at both labs, calling the Wuhan Institute of Virology’s ‘management of deadly viruses and virus-carrying lab animals ... appallingly poor and negligent.’

The memo provides an extraordinary window into behind-the-scenes concerns about a lab accident among U.S. foreign policy leaders, even as this line of inquiry was deemed a conspiracy theory by international virologists, some of whom had undisclosed conflicts of interest.

The memo also calls into question these virologists’ impartiality. Shi Zhengli, a Wuhan Institute of Virology coronavirus researcher nicknamed the ‘Bat Woman,’ has forged wide-reaching international collaborations, including with prestigious Western virologists, the memo notes.

‘Suspicion lingers that Shi holds an important and powerful position in the field in China and has extensive cooperation with many [international] virologists who might be doing her a favor,’ it reads ...

The memo laments that ‘the most logical place to investigate the virus origin has been completely sealed off from inquiry by the [Chinese Communist Party]’ ... The memo even suggests that other hypotheses may have served as a distraction from a probe of the city’s extensive research on novel coronaviruses. ‘All other theories are likely to be a decoy to prevent an inquiry [into] the WCDC and WIV,’ it states ...

The memo cites a 2015 paper³⁵ coauthored by Shi titled ‘A SARS-like cluster of circulating bat coronaviruses shows potential for human emergence’ that described creating a ‘chimera,’ or engineered virus, with the spike protein of a coronavirus from a Chinese horseshoe bat.

Editors at Nature Medicine added a note in March 2020 cautioning that the article was ‘being used as the basis for unverified theories that the novel coronavirus causing COVID-19 was engineered’ ... But the memo shows that the State Department indeed considered the paper relevant to the pandemic’s origins.”

NIH Retracted Gene Sequence at WIV Researcher’s Request

While we’ve yet to obtain bulletproof evidence that SARS-CoV-2 was developed as a bioweapon, there’s plenty of circumstantial evidence that points in that direction. Disturbingly, as time goes on, more and more of this circumstantial evidence seems to highlight the United States’ involvement. If one proverbial finger is pointing at China, four others are pointing back at us.

This is profoundly bad news, but it really ought to strengthen our resolve to get to the bottom of it. None of us are safe until the mad scientists responsible for this pandemic are brought to justice. It doesn’t matter who they are. In all likelihood, we’ll find that blame cannot be pinned on a single nation. At bare minimum, the U.S. and China appear to be covering for each other.

As just one example, there are the deletions of information that have occurred both at the National Institutes of Health and the WIV, either at the other’s request, or as what appears to be a favor.

As reported by Just the News,³⁶ NIH deleted a genetic sequencing submission of SARS-CoV-2 from

its Sequence Read Archive (SRA) at the request of a researcher at the WIV. Emails³⁷ obtained via FOIA request to the NIH by Empower Oversight show a WIV researcher who had submitted two genetic sequences to the SRA, one in March 2020, and a second in June 2020, asked to have the last one retracted.

NIH initially stated that it would be better to edit or replace the submission rather than retracting it, but the researcher insisted it be removed, which they did. To be fair, the NIH also states it has retracted at least eight SRA submissions in total, most from American researchers, at their request. However, emails also show the NIH directed reporters on how to provide more favorable and less sensationalized coverage of the deletion of the Chinese sequence. Just the News writes:³⁸

“[Empower Oversight] says one of the most disconcerting elements of the emails is evidence showing the NIH has refused to participate in a transparent process to examine data on the deleted sequences.

‘Most importantly, why has NIH refused to examine archival copies of deleted sequences in an open scientific process to determine whether any of that information might be able to shed light on the origins of the COVID-19 pandemic?’ the group asked.

However, that argument was dismissed by NIH official Steve Sherry. Although sequences are never fully deleted, according to the agency, Sherry told a researcher who asked for transparency, ‘As you know, when data sets are withdrawn from the database, that status does not permit use for further analyses.’”

WIV Deleted Mentions of US Collaborators

The WIV has also deleted information in what appears to be an effort to shield the NIH. Shortly after Fauci testified in a Senate hearing in March 2021,³⁹ the WIV quietly deleted all mentions of its collaboration with Fauci’s NIAID, the NIH and other American research partners from its website. As reported May 15, 2021, by The National Pulse:⁴⁰

“March 21st, 2021, the lab’s website listed six U.S.-based research partners: University of Alabama, University of North Texas, EcoHealth Alliance, Harvard University, the National Institutes of Health (NIH), the United States, and the National Wildlife Federation.”⁴¹

One day later, the page was revised to contain just two research partners — EcoHealth Alliance and the University of Alabama.⁴² By March 23rd, EcoHealth Alliance was the sole partner remaining.⁴³

EcoHealth Alliance is run by long-standing Chinese Communist Party-partner Dr. Peter Daszak, who National Pulse Editor-in-Chief Raheem Kassam has repeatedly claimed will be the first ‘fall guy’ of the Wuhan lab debacle ...

Beyond establishing a working relationship between the NIH and the Wuhan Institute of Virology, now-deleted posts⁴⁴ from the site also detail studies bearing the hallmarks of gain-of-function research conducted with the Wuhan-based lab.”

Indeed, a now-deleted WIV web page titled “Will SARS Come Back?” stated that:⁴⁵

“Prof. Zhengli Shi and Xingyi Ge from WIV, in cooperation with researchers from University of North Carolina, Harvard Medical School, Bellinzona Institute of Microbiology ... examine the disease potential of a SARS-like virus, SHC014-CoV, which is currently circulating in Chinese horseshoe bat

populations.

Using the SARS-CoV reverse genetics system, the scientists generated and characterized a chimeric virus expressing the spike of bat coronavirus SHC014 in a mouse-adapted SARS-CoV backbone.

The results indicate that group 2b viruses encoding the SHC014 spike in a wild-type backbone can efficiently use multiple orthologs of the SARS receptor human angiotensin converting enzyme II (ACE2), replicate efficiently in primary human airway cells and achieve in vitro titers equivalent to epidemic strains of SARS-CoV.

Evaluation of available SARS-based immune-therapeutic and prophylactic modalities revealed poor efficacy; both monoclonal antibody and vaccine approaches failed to neutralize and protect from infection with CoVs using the novel spike protein.

On the basis of these findings, they synthetically re-derived an infectious full-length SHC014 recombinant virus and demonstrate robust viral replication both in vitro and in vivo ...”

By Dr Mercola

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