

EXCLUSIVE: Official Documents prove Anthony Fauci & Bill Gates are responsible for developing the COVID Virus in a Biolab

BY THE EXPOSÉ ON NOVEMBER 25, 2022

The U.S. government funded Peter Daszak's ECO Health Alliance to the tune of \$61,491,183 to make new coronaviruses that are infectious to humans.

It also directly or indirectly funds the Wuhan Institute of Virology to the tune of \$7.4 million dollars to make new coronaviruses that are infectious to humans.

Meanwhile, the National Institute of Allergy and Infectious Diseases funded Ralph Baric to the tune of \$46,958,414 to make new coronaviruses that are infectious to humans.

This totals more than \$114 million in the period from 2000 to 2020 to make new coronaviruses which are infectious to humans.

There is no question that SARS-CoV 2 (causing Covid-19) is man-made. Here's the evidence...

By a concerned reader

A Brief History of man-made pathogens designed by Baric, Daszak, Zhengli and Moderna and funded by Fauci, the NIH, the NIAID and the US Government to carry out the research necessary for the creation of SARS, MERS and Covid-19 –

1984: Dr. Anthony Fauci is appointed director of the NIAID (National Institute of Allergies and Infectious diseases) – There is an Orwellian Appointment for you.

1986 July1: <https://grantome.com/grant/NIH/R01-AI023946-01> 1986 – 1999 funding amounts not specified and then funded by \$1,987,840 over 7 years from 2000 to 2007 excluding 2002.

NIAID (National Institute of Allergies and Infectious Diseases) funding Ralph Baric at the University of North Carolina, Chapel Hill.

Subject: Mechanism of MHV Replication (Mouse Hepatitis Virus). This was the start of the partnership between Fauci and Baric.

1999: U.S. Dept. of Health & Human Services (HHS) funds research amplifying the infectious character of Coronaviruses.

1999: The Bill and Mellisa Gates Foundation set up GAVI (Global Alliance for Vaccines and Immunization) with a promise of \$750 million in funding. So it was never going to be a bit part actor in the vaccine movie. – <https://www.gavi.org/operating-model/gavis-partnership-model/bill-melinda-gates-foundation>

2000: In May Ralph Baric successfully constructs a 28k base pair Coronavirus from smaller fragments by reverse transcription (making DNA from RNA) He makes an infectious TGE (Transmissible Gastroenteritis Virus – which is a type of Coronavirus) clone from isolated components of the genome. He files for [US Patent 6,593,111 B2](#): *The invention is exemplified by, although not limited to, the assembly of full-length, functional coronavirus genomes.*

The present inventors have successfully assembled a full-length infectious clone of transmissible gastroenteritis virus (TGE). Using a novel approach, six adjoining cDNA subclones that span the entire TGE genome were isolated. Each clone was engineered with unique flanking interconnecting junctions which dictate a precise, systematic assembly with only the correct adjacent cDNA subclones, resulting in an intact TGE cDNA construct of about approximately 28.5 Kb in length –
<https://patents.justia.com/patent/6593111> .

Among other advantages, full-length infectious clones of TGE permit the precise genetic modification of the coronavirus genome – <https://patents.justia.com/patent/6593111>

2001: April 20: NIH Grant GM63228 funds Ralph Baric et al at the University of North Carolina at Chapel Hill to do Chimera research to transform the original SARS Coronavirus which was pathogenic only to animals, existing in nature as a pathogen for dogs and rabbits, to make it pathogenic to mankind targeting lung epithelial cells. The justification for this research was that it was supposedly designed as a vector virus to distribute HIV vaccines. This resulted in US patent application US28531801P

2001 May21: Ralph Baric and Boyd Yount (Assigned to the University of North Carolina at Chapel Hill) file [US Patent 6,593,111 B2](#) Directional Assembly of Large Viral Genomes and Chromosomes The Patent Office search report found: Lai, Michael M.C “The making of infectious viral RNA: No size limit in sight” PNAS Vol 97 No 10, May 9, 2000 pp 5025-5027 And it found: Thiel et al “Infectious RNA transcribed in vitro from a cDNA copy of human coronavirus genome cloud in vaccinia virus 92: 1273-1261 (2001)

2002: April 19: The University of North Carolina files [US patent 7279327](#) for an infectious replication defective coronavirus (to be used as a virus vector for an HIV vaccine), claiming priority from US28531801P. Inventors were: Kristopher M. Curtis, Boyd Yount, Ralph S. Baric
<https://www.webmd.com/lung/news/20030411/sars-timeline-of-outbreak>

2002 November 16: The first case of an atypical pneumonia is reported in the Guangdong province in southern China.

2003 February 25: First cases of unusual pneumonia reported in Hanoi, Vietnam.

2003 February 28: WHO officer Carlo Urbani, MD, examines an American businessman with an unknown form of pneumonia in a French hospital in Hanoi, Vietnam.

2003 March10: Urbani reports an unusual outbreak of the illness, which he calls sudden acute respiratory syndrome or SARS, to the main office of the WHO. He notes that the disease has infected an unusually high number of healthcare workers (22) at the hospital (in Hanoi, Vietnam).

2003 March11: A similar outbreak of a mysterious respiratory disease is reported among healthcare workers in Hong Kong.

2003 March15: WHO issues a heightened global health alert about the mysterious pneumonia with a case definition of SARS after cases in Singapore and Canada are also identified. The alert includes a rare emergency travel advisory to international travelers, healthcare professionals and health authorities.

2003 March17: An international network of 11 leading laboratories is established to determine the cause of SARS and develop potential treatments.

2003 March24: CDC officials present the first evidence that a new strain of a virus most frequently associated with upper respiratory infections and the common cold in humans called the coronavirus might be a likely cause of SARS.

2003 March29: Carlo Urbani, who identified the first cases of SARS, dies as a result of the disease. Researchers later suggest naming the agent that causes the disease after the infectious disease expert.

2003 April4: President George W. Bush adds SARS to the list of quarantinable diseases, which gives the CDC the authority to isolate persons who might have been exposed to the disease.

2003 April12: Canadian researchers announce they have completed the first successful sequencing of the genome of the coronavirus believed to cause SARS.

2003 April14: CDC officials announce their laboratories have sequenced a nearly identical strain of the SARS-related coronavirus. The CDC version includes an additional 15 nucleotides, which provides the important beginning of the sequence.

2003 April16: A new form of a coronavirus never before seen in humans is confirmed as the cause of SARS according to Koch's postulates, which are four specific conditions that must be met for a pathogen to be confirmed as a causal agent of disease.

2003 April 25: CDC Patent filed and ultimately becomes US 7,220,852 (the patent on the RNA sequence) and 7,776,521 (the patent on the testing methodology). These

patents give the U.S. Department of Health and Human Services the ability to control the commercial exploitation of SARS coronavirus.

2003 July2: WHO removes Toronto, Canada from its list of areas with recent local SARS transmission after 20 days passed since the last SARS case was reported and isolated.

2003 January26: Dr. Anthony [Fauci appointed](#) to the Bill and Melinda Gates Foundation's Global Grand Challenges Scientific Advisory Board (served through 2010 – when he joined their Global Vaccine Action Plan).

2006 Chinese researchers combine HCV, HIV-1, SARS-CoV-1 & SARS-CoV-2: Huang Q, Cheng Y, Guo Q, Li Q. Preparation of a Chimeric Armored RNA as a Versatile Calibrator for Multiple Virus Assays. Clinchem 2006; 52(7):1446-1448 and Supplement A.Inform1999200020022003.

2007: NSF Grant IIS-0513650 (Italy, France and Indiana University) study addresses FIRST CRITICAL STEP to control a pandemic – shutdown International Travel. Given this knowledge why did Fauci tell Trump a Travel Ban was unnecessary?

2009 April15: ModernaTX INC is incorporated in Delaware USA

2010 August 6: Moderna (prior to its establishment) files U.S. Patent 9,447,164 which attracted the investment of (and “inventorship” for) venture capitalists at Flagship Ventures. This patent grew out of the work of Dr. Jason P. Schrum of Harvard Medical School supported by National Science Foundation Grant #0434507. While the application claims priority to August 2010, the application didn't get finalized until October, 2015.

2010 September21: Wildlife Trust is [rebranded to become ECO Health Alliance](#) under the directorship of Peter Daszak.

2010: The World Health Organization (WHO), UNICEF, the National Institute of Allergy and Infectious Diseases (NIAID) and the Bill & Melinda Gates Foundation have announced a collaboration to increase coordination across the international vaccine community and create a Global Vaccine Action Plan. <https://www.gatesfoundation.org/ideas/media-center/press-releases/2010/12/global-health-leaders-launch-decade-of-vaccines-collaboration>.

The Leadership Council is comprised of:

Dr. Margaret Chan, Director General of WHO;

Dr. Anthony S. Fauci, Director of NIAID, part of the National Institutes of Health;

Mr. Anthony Lake, Executive Director for UNICEF;

Ms. Joy Phumaphi, Chair of the International Advisory Committee and Executive Secretary, African Leaders Malaria Alliance

Dr. Tachi Yamada, President of Global Health at the Bill & Melinda Gates Foundation

2011 March9: ModernaTX (MODEfied RNA) was [registered in Massachusetts](#)

2012 September22: UK Identifies first case of MERS

2012-April2: Moderna claims priority to US201261618957P in its 5 patents cited below all of which refer to a double CGG codon Furin Cleavage site.

2013 December 16: Moderna files 4 patents referring to a double CGG codon Furin Cleavage site.

US9149506B2: Modified polynucleotides encoding septin-4 –

<https://patents.google.com/patent/US9149506B2/en>

US9216205B2: Modified polynucleotides encoding granulysin –

<https://patents.google.com/patent/US9216205B2/en>

US9255129B2: Modified polynucleotides encoding SIAH E3 ubiquitin protein ligase 1 –

<https://patents.google.com/patent/US9255129B2/en>

US9301993B2: Modified polynucleotides encoding apoptosis inducing factor 1 –

<https://patents.google.com/patent/US9301993B2/en>

2013: Baric and Chinese scientists isolate 3 coronaviruses from bats with HKU4 spike protein – unable to infect human cells. Yang Y...Baric RS, et al. Receptor usage and cell entry of bat coronavirus HKU4 provide insight into bat-to-human transmission of MERS coronavirus. PNAS 2014;111(34):12516-12521. Funded with NIH grants RO1AI089728 & R21AI109094.

2014 October 17: Obama Administration halts Gain-of-Function Research

2015 Dr. Zhengli et al “reengineered HKU4 spike aiming to build its capacity to infect human cells.” “To this end, we introduced two single mutations...mutations in these motifs in coronavirus spikes have demonstrated dramatic effects on viral entry into human cells.”

2015: Baric and Zhengli announce they can make a more dangerous, virulent and infectious virus. Two Mutations Were Critical for Bat-to-Human Transmission of Middle East Respiratory Syndrome Coronavirus (MERS). J Virol.2015;89(17):9199-9123. Funded by NIH grants RO1AI089728, RO1AI110700.

2015 July: South Korea had its last case of MERS. There were 2,494 confirmed laboratory cases and a total of 858 deaths in 27 countries; 80% of these cases came from Saudi Arabia. So the mortality rate for MERS was 34.4%. But it was not very transmissible. Infections were limited to family members, other hospital patients and health care workers who encountered MERS patients.

2016 June2: Moderna files US9587003B2 referring to a double CGG codon Furin Cleavage site: Modified polynucleotides for the production of oncology-related proteins and peptides – <https://patents.google.com/patent/US9587003B2/en>

2017: Gain-of-Function Research Ban Lifted

2018: US Patent 7279327 for the chimeric adaption of the naturally occurring animal SARS Coronavirus to become infectious to humans targeting lung epithelial cells is transferred from University of North Carolina to the US National Institute for Health (NIH) – who funded it in the first place.

2018 March 27: Peter Daszac and the Eco Health Alliance in 2018 submitted a proposal to DARPA to do gain of function research on SARS coronaviruses to insert a furin cleavage site (such as is found in HIV1) in an optimal position into the coronavirus gene. COVID-19 has a furin cleavage site optimally positioned to infect humans. <https://theintercept.com/2021/09/23/coronavirus-research-grant-darpa/>

Let's look at the big picture: A novel SARS coronavirus emerges in Wuhan with a novel cleavage site in it. We now have evidence that, in early 2018, they had pitched inserting novel cleavage sites into novel SARS-related viruses in their lab," said Chan. "This definitely tips the scales for me. And I think it should do that for many other scientists too."

<https://www.documentcloud.org/documents/21066966-defuse-proposal> – this is the Eco health alliance DARPA funding proposal.

Oh dear. That really nails it. DARPA confirms that Peter Daszac of the Eco Health Alliance submitted a proposal on 2018March27 to add an HIV1 type furin cleavage site to SARS1. SARS CoV 2 has that furin cleavage site. In fact the double CGG codon Furin cleavage site is the main difference between Covid-19 and the Bat Coronavirus RaGT13. Fauci funded Wuhan through the Eco Health Alliance. Here is Peter Daszak admitting that he and Ralph Baric made new forms of SARS in the lab –

The [Daily Mail said](#) of this proposal – *The most alarming aspect of the research plan revealed in the documents is a plan to search viral genetic databases for new types of 'furin cleavage sites' which help a virus attack a host. According to the proposal, 'high-risk' versions of these sites, once identified, would then be grafted onto SARS-like bat coronaviruses via genetic engineering. This revelation is alarming because SARS-CoV-2, the virus which causes COVID-19, has just such a furin cleavage site, which increases its power and deadliness, but a similar feature has never been observed in other SARS-like coronaviruses in nature.*

Now it seems that indictment and prosecution is in due order and for criminal action to proceed to punish the guilty partners. Get the Nuremberg gallows ready for the execution of the named guilty!

Blessings,

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