Mass Murder

By Syringe Needle!
Part 54

IMPORTANT INFORMATION BELOW FOR THOSE VAXXED AND COVID WAS PATENTED IN 2013

7-Year-Old Died of Cardiac Arrest 13 Days After Pfizer Shot, VAERS Data Show

VAERS data released Friday by the Centers for Disease Control and Prevention included a total of **1,168,894** reports of adverse events from all age groups following COVID vaccines, including **25,158 deaths** and **203,888 serious injuries** between Dec. 14, 2020, and March 4, 2022.

The Centers for Disease Control and Prevention (CDC) today released new data showing a total of 1,168,894 reports of adverse events following COVID vaccines were submitted between Dec. 14, 2020, and March 4, 2022, to the Vaccine Adverse Event Reporting System (VAERS). VAERS is the primary government-funded system for reporting adverse vaccine reactions in the U.S.

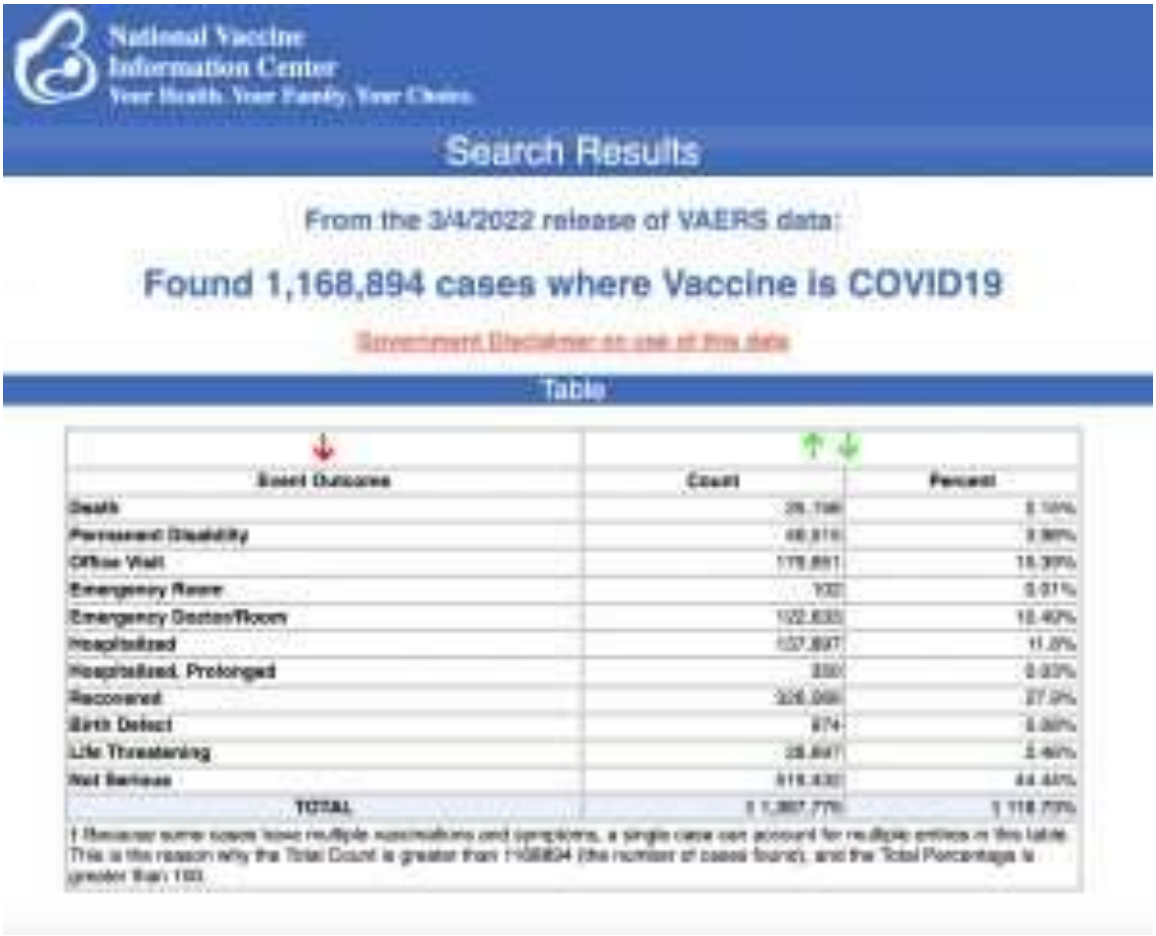
The data included a total of 25,158 reports of deaths — an increase of 331 over the previous week — and 203,888 reports of serious injuries, including deaths, during the same time period — up 3,557 compared with the previous week.

Excluding "foreign reports" to VAERS, 783,282 adverse events, including 11,505 deaths and 75,286 serious injuries, were reported in the U.S. between Dec. 14, 2020, and March 4, 2022.

Foreign reports are reports foreign subsidiaries send to U.S. vaccine manufacturers. Under U.S. Food and Drug Administration (FDA) regulations, if a manufacturer is notified of a foreign case report that describes an event that is both serious and does not appear on the product's labeling, the manufacturer is required to submit the report to VAERS.

Of the 11,505 U.S. deaths reported as of March 4, 17% occurred within 24 hours of vaccination, 22% occurred within 48 hours of vaccination and 60% occurred in people who experienced an onset of symptoms within 48 hours of being vaccinated.

In the U.S., 554 million COVID vaccine doses had been administered as of March 4, including 327 million doses of Pfizer, 209 million doses of Moderna and 18 million doses of Johnson & Johnson (J&J).


National Vaccine Information Center
Your Health. Your Family. Your Choice.

Search Results

From the 3/4/2022 release of VAERS data:

Found 1,168,894 cases where Vaccine is COVID19

Government Disclaimer on use of this data

Table

Event Outcome	Count	Percent
Death	[illegible]	[illegible]
Permanent Disability	[illegible]	[illegible]
Office Visit	[illegible]	[illegible]
Emergency Room	[illegible]	[illegible]
Emergency Doctor/Room	[illegible]	[illegible]
Hospitalized	[illegible]	[illegible]
Hospitalized, Prolonged	[illegible]	[illegible]
Recovered	[illegible]	[illegible]
Birth Defect	[illegible]	[illegible]
Life Threatening	[illegible]	[illegible]
Not Serious	[illegible]	[illegible]
TOTAL	[illegible]	[illegible]

† Because some cases have multiple vaccinations and symptoms, a single case can account for multiple entries in this table. This is the reason why the Total Count is greater than 1168894 (the number of cases found), and the Total Percentage is greater than 100.

Every Friday, VAERS publishes vaccine injury reports received as of a specified date. Reports submitted to VAERS require further investigation before a causal relationship can be confirmed. Historically, VAERS has been shown to report only 1% of actual vaccine adverse events.

U.S. VAERS data from Dec. 14, 2020, to March 4, 2022, for 5- to 11-year-olds show:

- 9,009 adverse events, including 213 rated as serious and 5 reported deaths.

The most recent death involves a 7-year-old boy (VAERS I.D. 2152560) from Washington who died 13 days after receiving his first dose of Pfizer's COVID vaccine when he went into shock and suffered cardiac arrest. He was unable to be resuscitated and died in the emergency department.

- 17 reports of myocarditis and pericarditis (heart inflammation).

The CDC uses a narrowed case definition of "myocarditis," which excludes cases of cardiac arrest, ischemic strokes and deaths due to heart problems that occur before one has the chance to go to the emergency department.

- 32 reports of blood clotting disorders.

U.S. VAERS data from Dec. 14, 2020, to March 4, 2022, for 12- to 17-year-olds show:

- 30,193 adverse events, including 1,734 rated as serious and 40 reported deaths.

The most recent death involves a 14-year-old boy (VAERS I.D. 2148498) who experienced a cerebral aneurysm leading to death one day after receiving his first dose of Pfizer's COVID vaccine.

- 69 reports of anaphylaxis among 12- to 17-year-olds where the reaction was life-threatening, required treatment or resulted in death — with 96% of cases attributed to Pfizer's vaccine.
- 650 reports of myocarditis and pericarditis with 631 cases attributed to Pfizer's vaccine.
- 161 reports of blood clotting disorders, with all cases attributed to Pfizer.

U.S. VAERS data from Dec. 14, 2020, to March 4, 2022, for all age groups combined, show:

- 19% of deaths were related to cardiac disorders.
- 54% of those who died were male, 41% were female and the remaining death reports did not include the gender of the deceased.
- The average age of death was **72.7**.
- As of March 4, 5,233 pregnant women reported adverse events related to COVID vaccines, including 1,664 reports of miscarriage or premature birth.
- Of the 3,596 cases of Bell's Palsy reported, 51% were attributed to Pfizer vaccinations, 40% to Moderna and 8% to J&J.
- 859 reports of Guillain-Barré syndrome, with 41% of cases attributed to Pfizer, 30% to Moderna and 28% to J&J.
- 2,358 reports of anaphylaxis where the reaction was life-threatening, required treatment or resulted in death.
- 1,632 reports of myocardial infarction.
- 13,428 reports of blood clotting disorders in the U.S. Of those, 5,992 reports were attributed to Pfizer, 4,784 reports to Moderna and 2,607 reports to J&J.
- 4,065 cases of myocarditis and pericarditis with 2,499 cases attributed to Pfizer, 1,381 cases to Moderna and 174 cases to J&J's COVID vaccine.

CDC study concludes COVID vaccine adverse events 'mild'

A study funded by the CDC and published Monday in The Lancet concluded most COVID vaccine-related adverse events reported during the first six months of the rollout in the U.S were "mild and short in duration," despite thousands of deaths reported to VAERS.

For the study, researchers analyzed data captured between Dec. 14, 2020, and June 14, 2021, by VAERS and v-safe, both of which are overseen by the CDC. Nearly 300 million doses of COVID vaccines were administered during the study period.

The authors found that of the 340,522 adverse events reported to VAERS, 27,023 (8%) were serious, 4,496 were deaths. The authors said the cause of the increased reporting of deaths during the first few days after vaccination might represent "reporting bias."

The authors suggested that deaths occurring soon after vaccination were more likely to be reported than deaths that occurred later. This, they believe, is why the number of deaths asymptotically approaches zero as more time elapses since vaccination.

Jessica Rose, Ph.D., attempted to duplicate the Lancet authors' findings through her independent analysis of the VAERS data. Despite filtering the database using three different date stamps, Rose was unable to duplicate the Lancet study's results.

Florida surgeon general breaks with CDC, recommends against shots for healthy kids

Florida's surgeon general on Monday said he will issue guidance formally recommending against COVID vaccines for healthy children. Florida is the first state to break with official guidance from the CDC, which recommends all children over age 5 get the vaccine.

Dr. Joseph Ladapo made the announcement at a roundtable, hosted by Gov. Ron DeSantis, featuring physicians and other medical experts who criticized CDC and government policies, including mask mandates and lockdowns, which they said were ineffective and harmful.

Ladapo and DeSantis said the new guidance had to do with lingering questions about the vaccines' potential health risks for young people and the fact that children are in a low-risk category for severe COVID.

COVID vaccines may be enhancing disease

COVID vaccines may be causing enhanced disease because they target an old version of the coronavirus, Dr. Robert Malone told the Epoch Times in a recent interview.

"The data are showing that vaccination can actually increase the risk of being infected with the Omicron version of this virus," Malone said, referring to how in some areas, including Scotland and New Zealand, patients hospitalized with COVID are more likely to have received a COVID vaccine.

U.S. drug regulators identified vaccine-associated enhanced diseases (VAED) as an "important potential risk" of COVID vaccines, along with enhanced respiratory disease.

Some adverse events reported following COVID vaccination "could indicate" VAED, according to a CDC team.

VAED refers to disease "resulting from infection in individuals primed with non-protective immune responses against the respective wild-type viruses," researchers said last year.

"Given that these enhanced responses are triggered by failed attempts to control the infecting virus, VAED typically presents with symptoms related to the target organ of the infection pathogen," they added.

Vaccine researcher develops tinnitus after COVID shot, calls for further study

Dr. Gregory Poland, director of the Mayo Clinic's Vaccine Research Group in Rochester, Minnesota, developed life-altering tinnitus, or ringing in the ear, after receiving his second dose of a COVID vaccine.

"It was like someone suddenly blew a dog whistle in my ear," Poland told MedPage Today. "It has been pretty much unrelenting."

Poland then received a booster, after which his tinnitus briefly disappeared but then returned at a slightly higher pitch. Poland realized his life may never be the same and says he has received emails from people across the world struggling with the same condition to the point they're considering taking their own lives.

Poland, who said he supports COVID vaccines, believes there may be tens of thousands of people affected in the U.S. and is calling for more research to be done to provide help to people desperate for relief.

Michigan woman files claim over mom's COVID vaccine-related death

Tatum Strieter-Byron is asking the federal government to compensate her for the death in April of her mother Sandra Jacobs. An autopsy confirmed Jacobs died from a blood-clotting disorder caused by J&J's COVID vaccine.

Strieter-Byron received confirmation Monday her claim to the Countermeasures Injury Compensation Program (CICP) had been received. The program was established to give pharmaceutical companies blanket liability protection from harm caused by their COVID vaccines.

In 2020, the U.S. Health and Human Services secretary invoked the Public Readiness and Emergency Preparedness (PREP) Act and declared COVID-19 a public health emergency, providing J&J other COVID vaccine makers immunity from lawsuits.

The only exception under the PREP Act is if a plaintiff can prove a vaccine-related death or serious physical injury was caused by "willful misconduct." The protections, unless amended or rescinded, extend through Oct. 1, 2024.

Claims to the CICP must be made within one year of the date the vaccine was received. Jacobs, 60, received the single-shot vaccine at a CVS pharmacy on April 8, 2021, just five days before federal health agencies temporarily paused the vaccine to examine numerous reports of a serious and potentially fatal blood-clotting disorder.

Jacobs died on April 21 of "complications of cerebral venous sinus thrombosis," a type of stroke caused by the vaccine.

Children's Health Defense asks anyone who has experienced an adverse reaction, to any vaccine, to file a report following these three steps.

Attorney Brent Wisner Tells RFK, Jr.: Drugmakers Knew Zantac Caused Cancer But Sold It Anyway for 40 Years

Attorney Brent Wisner Tells RFK, Jr.: Drugmakers Knew Zantac Caused Cancer But Sold It Anyway for 40 Years
David Icke / Richard Willet – Memes and headline comments by David Icke

In an interview on 'RFK Jr. The Defender Podcast,' award-winning attorney Brent Wisner talked with Robert F. Kennedy, Jr. about litigation to hold Zantac manufacturers accountable.

The once widely used heartburn drug Zantac was the topic of a recent episode of the "RFK Jr. The Defender Podcast."

Robert F. Kennedy, Jr., chairman and chief legal counsel of Children's Health Defense, interviewed Brent Wisner, an award-winning mass tort litigator with Baum Hedlund Aristei & Goldman, who is representing plaintiffs who allegedly developed cancer after taking Zantac. The plaintiffs are suing the pharmaceutical companies that sold the drug.

Kennedy and Wisner are colleagues. They worked together during the landmark lawsuit Dewayne Johnson v. Monsanto Company, the first Roundup cancer lawsuit to proceed to trial. The jury awarded groundskeeper Johnson $289.2 million.

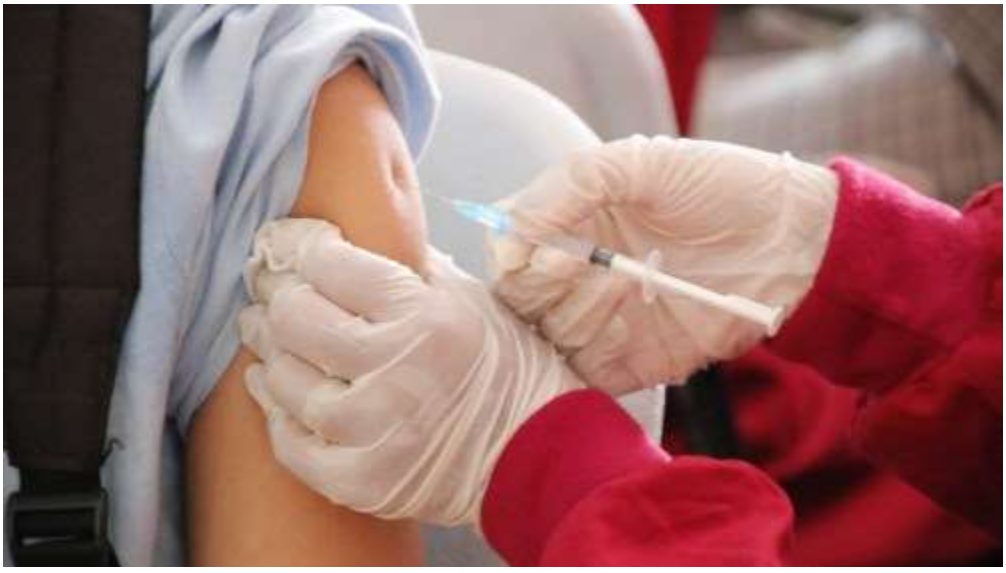
"Brent Wisner arguably is the best lawyer in the country," said Kennedy. *"He's won every award that you can possibly win for litigation. He is a versatile lawyer and he's an extraordinary performer. He has a chemical link to jurors like I've never seen before."*

Wisner is now working to hold accountable the makers of Zantac for allegedly concealing knowledge of a cancer-causing ingredient in the drug.

Zantac, an over-the-counter medication, was withdrawn from the market in 2020 when it was determined that it contained the human carcinogen N-nitrosodimethylamine (NDMA).

Scientists Currently Developing Controversial "Contagious Vaccines" That Can Spread From Vaccinated to Unvaccinated

The Gateway Pundit / Jim Hoft



Groups of scientists are currently developing 'self-spreading vaccines' that could infect others from vaccinated to unvaccinated people or between vaccinated to unvaccinated animals, according to **National Geographic**.

The experiment was designed to spread the vaccine to unvaccinated people in vaccinated person's close proximity.

"The idea is that instead of a vaccine staying in one person's body, the vaccine itself would infect them in such a way that they could pass on vaccination to others around them, much as they would otherwise pass on a disease. Scientists could vaccinate one person or animal in a community, and the vaccination would spread to those around them," per **Newsbreak.**

According to the report, scientists are currently developing 'contagious vaccines' for Ebola, bovine tuberculosis, and Lassa fever, a viral disease spread by rats.

The scientists are also planning to expand their studies to other zoonotic diseases including rabies, West Nile virus, Lyme disease, and the plague.

The vaccines use cytomegalovirus (CMVs), a group that belongs to the herpes family. According to **Mayo Clinic**, once infected with the virus, your body retains the virus for life.

"CMV spreads from person to person through body fluids, such as blood, saliva, urine, semen, and breast milk. There is no cure, but there are medications that can help treat the symptoms."

National Geographic reported:
Imagine a cure that's as contagious as the disease it fights—a vaccine that could replicate in a host's body and spread to others nearby, quickly and easily protecting a whole population from microbial attacks. That's the goal of several teams around the world who are reviving controversial research to develop self-spreading vaccines.

Researchers are currently developing self-spreading vaccines for Ebola, **bovine tuberculosis**, and Lassa fever, a viral disease spread by rats that causes upward of **300,000 infections** annually in parts of West Africa. The approach could be expanded to target other **zoonotic diseases**, including rabies, West Nile virus, Lyme disease, and the plague.

Advocates for self-spreading vaccines say they could revolutionize public health by disrupting infectious disease spread among animals before a zoonotic spillover could occur—potentially preventing the next pandemic.

But others argue that the viruses used in these vaccines could themselves mutate, jump species, or set off a chain reaction with devastating effects across entire ecosystems.

"Once you set something engineered and self-transmissible out into nature, you don't know what happens to it and where it will go," says **Jonas Sandbrink**, a biosecurity researcher at the University of Oxford's Future of Humanity Institute. *"Even if you just*

start by setting it out into animal populations, part of the genetic elements might find their way back into humans."

Vaccines in progress

Renewed interest and funding for the technology popped up around 2016, and today several research groups are developing self-spreading vaccines for animals.

Each of these new vaccines are so-called recombinant viruses. Researchers first identify a protein from the target microbe that serves as an antigen—a substance that triggers immune responses in vaccinated people or animals. Then the researchers select a virus to carry the vaccine and spread it. To do this, researchers capture a few animals from their target population—primates for Ebola, rats for Lassa fever—and isolate a virus that naturally infects those animals. Then they splice in genetic material from the target to create a vaccine.

Each of these vaccines uses a cytomegalovirus, or CMVs, a group that belongs to the herpes family.

CMVs help the researchers overcome several technical challenges. For one, CMVs have large genomes made from double-stranded DNA, which means their genetic code is more stable and can accommodate additional genes from the targeted microbe, **says Alec Redwood, a principal research fellow at the University of Western Australia. He conducted self-spreading vaccine research in the early 2000s and is now part of a team developing a CMV-based Lassa fever vaccine**.

So far, no one has conducted any field or laboratory studies assessing the impact and safety of these vaccines delivered via the self-spreading mechanism. However, a recent mathematical modelling study reported that if it works as expected, releasing the Lassa fever vaccine could reduce disease transmission among rodents by 95 percent in less than a year. The question that can be derived from this article is how a self-spreading vaccine will impact humanity. This becomes another nail in the coffin of those driven by the Eugenics program for "Depopulation" of the Earth!

"Once you set something engineered and self-transmissible out into nature, you don't know what happens to it and where it will go," warned Jonas Sandbrink, a biosecurity researcher at the University of Oxford's Future of Humanity Institute.

"Even if you just start by setting it out into animal populations, part of the genetic elements might find their way back into humans."

However, interest in the technology resurfaced in 2016, and new experiments are set to begin on the Lassa virus vaccine within the next year, but because of the "extremely high-risk and international nature of this work" and the "potentially irreversible" consequences, this experimental research faces numerous ethical and legal hurdles.

"We can't even get people to take a vaccine in a global pandemic. The idea that you would be able to surreptitiously vaccinate the population with a virus without causing riots is just, you know, it's stuff of fantasy. It will never be used in humans," according to Alec Redwood, a principal research fellow at the University of Western Australia.

Nevertheless, Redwood insists that this dangerous technology should still be developed just in case it ever needs to be used on humans "if we need it."

"You don't need to be a Rhodes scholar to work out that people will be nervous about a disseminating viral vector. It's a concept that will scare people," Redwood said.

"The way that I like to think about it is that it may never be used, but it's better to have something in the cupboard that can be used and is mature if we need it. And to say, 'Let's just not do this research because it's too dangerous,' to me, that makes no sense at all," he added.

Read the full story here.
Original Article: https://www.thegatewaypundit.com/2022/03/scientists-currently-developing-controversial-contagious-vaccines-can-spread-vaccinated-unvaccinated/

Frontline doctor: Millions will develop AIDS from COVID jabs

NaturalNews.com / Ethan Huff

(Natural News) A Florida-based physician of osteopathic medicine is warning that in just a few short months, millions of people who got "vaccinated" for the Wuhan coronavirus (COVID-19) will develop full-blown AIDS.

Dr. Elizabeth Eads, who has been working in her field for 25 years, told *USA Watchdog*'s Greg Hunter that the triple injected are showing the worst signs of vaccine-induced AIDS (VAIDS).

Eads and her team are trying to come up with remedies to help them, but the situation is difficult. So far, they have tried hydroxychloroquine (HCQ) and ivermectin, but nothing is helping her jab-damaged patients.

*"Yes, we are seeing **vaccine-related acquired immunodeficiency** in the hospital now from the triple vaxxed," Eads said. "It is a vax injury, and we are not really certain how to treat this. We are kind of throwing the kitchen sink at it. We are trying to use everything we can think of to boost up the CD4 and CD8 counts and reverse this collapse, this calamity of immune collapse. It's very stunning."*
You can watch a video interview with Eads **at NewsWars.com.**

COVID “booster” shots are the “kill shot,” Eads warns.

Eads said she is seeing a trend where the people who are the worst-off in terms of showing autoimmune or neurodegenerative symptoms are those who took the most injections.

The triple-shot, in other words, have basically destroyed their immune systems. The single- and double-shot *might* be able to recover, or perhaps it is just taking a lot longer for their damaged immune systems to degrade.

Eads said the third injection is a “kill shot” or a “money ball” because of how “devastating” it is to the immune system.

“If you look at the recent Stanford study, and I am just going to read a couple of sentences from the Stanford study: ‘The spike protein in the COVID-19 vaccines that everyone is talking about is called the lentivirus. The lenti contains a combination of HIV, types one through three, SRV/1, which is AIDS, MERS and SARS,’” she explained. “In the Stanford study, the best-known lentivirus is the human immune deficiency pathogen, which causes AIDS. This is why we are seeing autoimmune and neurodegenerative decline after the COVID-19 vaccine, especially the booster. It permanently changes the genome of the cell. That is why this is so terrifying to us in the medical community. We just don’t know how to attack this.”

According to Eads, the spike lentivirus is made up of HIV and AIDS, along with SARS and MERS. This, she said, is why both the vaccinated and the “boosted” are getting seriously ill and dying.

“That’s why they dominate the hospitalizations regarding COVID illness as well,” Eads emphasized.

From about April through the summer, Eads expects to see a whole lot more cases of AIDS diagnosed in the fully vaccinated, and especially in the 18-39 age demographic. Over the next year, the fully jabbed in their 40s and 50s will catch up with the younger group as their immune systems start to collapse.

A U.K. *Health Security Agency* study found that fully jabbed people in the 30-70 age demographic have already lost about 70 percent of their immune system capacity. That percentage will only continue to increase in the coming months and years.

Another thing we are seeing is a sharp uptick in cancer cases among the fully jabbed. Malignant neo-plasma of the esophagus is up 794 percent while malignant neo-plasma of the stomach, colon and pancreas is up 524 percent.

“Breast cancer up 387 percent, ovarian cancer up 537 percent, testicular cancer up 269 percent. These are numbers from 2021,” Eads further revealed.

People everywhere are developing serious chronic illnesses or dying due to Fauci Flu shots. You can keep up with the latest at **ChemicalViolence.com**.

Moderna Vaccine Patented 9 Months Before Pandemic

Published on March 25, 2022
Written by Fabio Giuseppe Carlo Carisio

A disturbing document had already emerged proving the existence of an experimental gene serum based on messenger RNA against Covid-19 on 12 December 2019.

But nothing appears compared to the terrible secrets about the Spikevax vaccine produced by Big Pharma Moderna in Cambridge (Massachusetts, USA) thanks to the money of the Bill & Melinda Gates Foundation, the Pentagon (the American Department of Defense) and the collaboration with the Niaid (Institute Allergy and Infectious Diseases National Park) headed by Anthony Fauci, White House advisor on the SARS-Cov-2 pandemic.

Patent US10702600 for the vaccine candidate mRNA-1273 was in fact registered in its new composition on March 28, 2019.

That is 9 months before the official outbreak of the Covid-19 pandemic in China and the availability of the official Wuhan sequence MN908947.1, the virus initially called 2019-nCoV and then renamed by the World Health Organization SARS-Cov-2 to the strong genomic identity with SARS of 2003 (Severe Acute Respiratory Syndrome, or Severe Acute Respiratory Syndrome, when virus and disease were called the same).

This revelation comes from “The Fauci / COVID-19 Dossier” prepared by the American doctor David E. Martin who has developed a colossal study on U.S. government-funded patents in relation to Coronaviruses since 2000, so much so that he suspects that SARS too in 2003 was built in the laboratory by microbiologist Ralph Baric of the Chapel Hill research center of the University of North Carolina (UNC) under the aegis of Fauci himself.

As we will see after long but inevitable premises, the registration of Moderna’s experimental vaccine on March 28, 2019 is also confirmed by the official history stored in the patent … [Actually, from the last article in this segment, Moderna patented its Vax much earlier. Sources in the UK discovered 4 patents for its Vaxx dating to 2013!]

The discovery becomes even more disturbing in light of the fact that such research was conducted by the pharmaceutical company at the same time as Baric’s extremely dangerous experiments on SARS recombinant viruses conducted together with the director of the Infectious Diseases Research Center of the Wuhan Institute of Virology, Shi Zhengli, not randomly dubbed in China “BatWoman” for its coronavirus tests of horseshoe bats.

As anticipated before the online investigation in the book WuhanGates (December 2020), they would be the godfathers and progenitors of a huge plot of the New World Order of ancient British Masonic origin hatched under the **sign of the Chinese Communists and American Democrats.**

To it were also added **the peremptory paws of Italy and Saudi Arabia** aimed at global immunization projects preparatory to the health dictatorship of the Covid-19 emergency and the promotion of experimental anti-Covid gene serums for the enrichment of Big Pharma and speculation of the same investment funds that control the **Weapons Lobby.**

THE THERAPY OF DONNO AFFOSSED BY THE MODERNA GROUP

Before briefly summarizing the many information contained in the WuhanGates 40 report on the researches of Professor Baric and Doctor Fauci, coordinator of all the most risky and secret experiments on viruses and bacteriological weapons in the 28 American civil and military laboratories as evidenced by the **exclusive investigation by Gospa News**, we recall another of the mysteries in the ambiguous fight against Covid-19 revealed **only by our online newspaper.**

Moderna is part of that colossal international Zacks group that became interested in the highly effective **therapy against Covid-19 of hyperimmune plasma developed by Professor Giuseppe De Donno** and mysteriously vanished in the maze of health policy before his mysterious suicide.

It should also be remembered that the experimental gene sera were able to obtain emergency authorization in the USA, from the Food and Drug Administration which, however, in recent weeks gave definitive ok to Spikevax after having granted it to

Comirnaty of Pizer-Biontech amid a thousand controversies for **not having subjected the clinical data to the scrutiny of the independent Advisory Committee**, and in the European Union by the European Medicines Agency (where they remain with a conditional marketing authorization as still experimental) only by virtue of two factors.
The pandemic has resulted in millions of deaths in the world with Covid-19 (in the minority died from Covid-19 and in the majority from complications of other diseases), in Italy and in other countries mainly due to inadequate home and hospital care such as the betrayal. use of cortisone or the disputed hydroxychlorichine (adopted by the protocols of the Piedmont Region) and the absence of drugs recognized as effective.

Therefore, the stop of the De Donno therapy, successfully carried out in over one hundred American university hospitals as recalled by the Johns Hopkins University in its defense, represented one of the obstacles in alternative treatments functional to the promotion of Moderna and other Big Pharma's anti-Covid vaccines.

EXPERIMENTS ON CHIMERIC VIRUSES BEFORE 2003

«The National Institute of Health's grant AI23946-08 issued to Dr. Ralph Baric at the University of North Carolina at Chapel Hill (officially classified as affiliated with Dr. Anthony Fauci's NIAID by at least 2003) began the work on synthetically altering the Coronaviridae (the coronavirus family) for the express purpose of general research, pathogenic enhancement, detection, manipulation, and potential therapeutic interventions targeting the same. As early as May 21, 2000, Dr. Baric and UNC sought to patent critical sections of the coronavirus family for their commercial benefit».

As Martin's dossier reports, in one of several articles derived from the work sponsored by this grant, «Baric published what he reported to be the SARS CoV full-length cDNA in which it was clearly stated that SAR CoV was based on a compound. of DNA segments».

«Using a panel of contiguous cDNAs that span the entire genome, we have assembled a full-length cDNA of the SARS-CoV Urbani strain, and have rescued molecularly cloned SARS viruses (infectious clone SARS-CoV) that contained the expected marker mutations inserted into the component clones».

SARS-Cov of 2003 was indicated with the name of Carlo Urbani, the researcher who was analyzing all its characteristics and died prematurely, struck down by the respiratory syndrome, like other scientists who died mysteriously while working on chimeric virus tests (Franc Plummer in Canada) or when they tried to investigate the origin of these viruses and the gene sera then placed on the market after SARS-Cov-2 (**Franco Trinca and Domenico Biscardi** and finally the elderly Luc Montagnier).

For the sake of brevity, let's skip all the **references to the patents already mentioned in WuhanGates 40** and come to the conclusions of "The Fauci / Covid-19 Dossier" published by Dr. Martin: «In short, the U.S. Department of Health and Human Services was involved in the funding of amplifying the infectious nature of coronavirus between 1999 and 2002 before SARS was ever detected in humans».

The allusion to the virus that **killed the Italian scientist Urbani intent on studying it is very heavy:** even SARS 2003 could have been created in the laboratory!
The National Institutes of Health, Allergy and Infectious Diseases worked on SARS Reverse Genetics. study AI059136-01. $1.7 million total costs, RS Baric, PI. 10 percent effort. 4/1/04- 3/31/09. The project develops a SARS-CoV full length infectious cDNA, the development of SARS-CoV replicon particles expressing heterologous genes, and seeks to adapt SARS-CoV to mice, producing a pathogenic mouse model for SARS-CoV infection. Then, subsequently, the same NIAID carries out the research Remodeling the SARS Coronavirus Genome Regulatory Network. RS Baric, PI 10 percent effort. 7/1/04-6/30/09. $2.1 million.

On November 22, 2004, the University of Hong Kong patents the spike protein associated with SARS on CoV and pursues US patent 7,491,489. But on June 2005 also the Pentagon's DARPA (Defense Advanced Research Projects Agency) gets in on the game Synthetic Coronaviruses with the event Biohacking: Biological Warfare Enabling Technologies, organized in Washington, DC. and sponsored by DARPA/MITRE.

In 2008, funding for **Biodefense Grant U54 AI057157 commences with $10,189,682 to UNC Chapel Hill**. Where, in the laboratories of North Carolina University, the experiments on chimeric superviruses will be carried out between 2014 and 2017, conducted in spite of Barack Obama's moratorium on gains in function. And in 2010 the "Biodefense Grant U54 AI057157" continues with $ 8,747,142 to UNC Chapel Hill (non-competitive grant from NIAID). Patent issuance for the SARS coronavirus patents peaked after the outbreak in Asia with 391 patents issued.

We are talking about research funded by DARPA and the Pentagon precisely because they are dedicated to the construction of recombinant viruses, especially **SARS infected with HIV as happened with the plasmids built by the Wuhan Institute of Virology** thanks to a funding from the European Commission chaired by Romano Prodi, with an increase in the charge. viral through the extremely dangerous Gain of Function technique and with "dual use" purposes, i.e. vaccine but also a military bacteriological weapon: as is the SARS-Cov-2 according to a sworn report drawn up by Professor Montagnier and filed in a legal action by two lawyers British.

«The CDC and NIAID entered into trade among States (including, but not limited to working with University of North Carolina, Chapel Hill) and with foreign nations (specifically, the Wuhan Institute of Virology and the Chinese Academy of Sciences represented by Zheng-Li Shi) through U19AI109761 (Ralph S. Baric), U19AI107810 (Ralph S. Baric), and National Natural Science Foundation of China Award 81290341 (Zheng-Li Shi) et al. 2015-2016» reads in The Fauci / COVID-19 Dossier.

INTRIGUE WITH THE CIA AND US INTELLIGENCE

«By October 2013, the Wuhan Institute of Virology 1 coronavirus S1 spike protein was described in NIAID's funded work in China. **This work involved NIAID, USAID, and Peter Daszak, the head of EcoHealth Alliance**. This work, funded under

R01AI079231, was pivotal in isolating and manipulating viral fragments selected from sites across China which contained high risk for severe human response» remembered Martin.

In the same years, the Obama-Biden administration chooses lawyer Avril as deputy director from the Central Intelligence Agency, the American CIA counter-espionage which is also responsible for the supervision of military projects and which often uses the government agency USAID as a financial instrument for occult international operations. Haines, an expert in drones but also in biological weapons.

Haines, subsequently, not only in 2018 prophesied a coronavirus epidemic that could only be faced with a new "World Order" but will be one of the main protagonists of the Event 201 exercise held in mid-October in New York thanks to funding from the World Economic Forum by Klaus Schwab (Great Reset) and the Bill & Melinda Gates Foundation.

On the recommendation of the American President Joseph Biden (her longtime political friend) on 21 January 2021 **she was appointed by the Senate National Director of Intelligence** from where she coordinates all 17 military and civil intelligence agencies and from where she searched, without success. for a rift between 007, to refute the thesis of the **artificial origin of SARS-Cov-2.**

MODERNA STUDIES IN THE SAME YEARS

«By March 2015, both the virulence of the S1 spike protein and the ACE II receptor was known to present a considerable risk to human health. NIAID, EcoHealth Alliance and numerous researchers lamented the fact that the public was not sufficiently concerned about coronavirus to adequately fund their desired research» the dossier still reads in reference to the experiments conducted in a "China-US affair" as Montagnier claimed on the origin of **laboratory SARS-Cov-2.**

«In 2013, the Autonomous Diagnostics to Enable Prevention and Therapeutics (ADEPT) program awarded grant funding to Moderna Therapeutics for the development of a new type of vaccine based on messenger RNA. The initial DARPA grant was W911NF-13-1-0417» reveals Dr. Martin.

«ON 2015 Moderna signs an agreement for the development of vaccines with NIAID and executes it with the head of the developer and main inventor mRNA-1273 Giuseppe Ciaramella» says Dr. Martin referring to the "prototype" of the experimental antiCovid gene serum named in 2020 Spikevax.

«ON 2016 NIH through Scripps Institute and Dartmouth College file patent application WO 2018081318A1 "Prefusion Coronavirus Spike Proteins and their Use" disclosing mRNA technology that overlaps (and is used in tandem with) Moderna's technology. Lead Inventor Barney Scott Graham was well known to Moderna as he's the person at NIH that Moderna "e-mailed" to get the sequence for SARS CoV-2 according to

Moderna's report. In addition, co-inventor Jason McLellan worked with Graham on a vaccine patent jointly owned with the Chinese government filed in Australia in 2013», mRNA-1273 – the experimental vaccine developed by Moderna for COVID-19 – uses the LNP technology that Moderna thought it had licensed from Acuitas Therapeutics Inc., a firm developed by a former principal of Arbutus' prior company Tekmira. That license did not authorize Moderna to use the technology for the COVID-19 vaccine.

«M - CAM and Knowledge Ecology International (Martin's reference bodies – ed) have independently confirmed that Moderna has violated U.S. law in failing to disclose the U.S. government's funding interest in their patents and patent applications» reads in The Fauci / Covid-19 Dossier.

«While this negligence impacts all of Moderna's over 130 granted U.S. patents, it is particularly problematic for U.S. Patent 10,702,600 ('600) which is the patent relating to,

"a messenger ribonucleic acid (mRNA) comprising an open reading frame encoding a betacoronavirus (BetaCoV) S protein or S protein subunit formulated in a lipid nanoparticle».

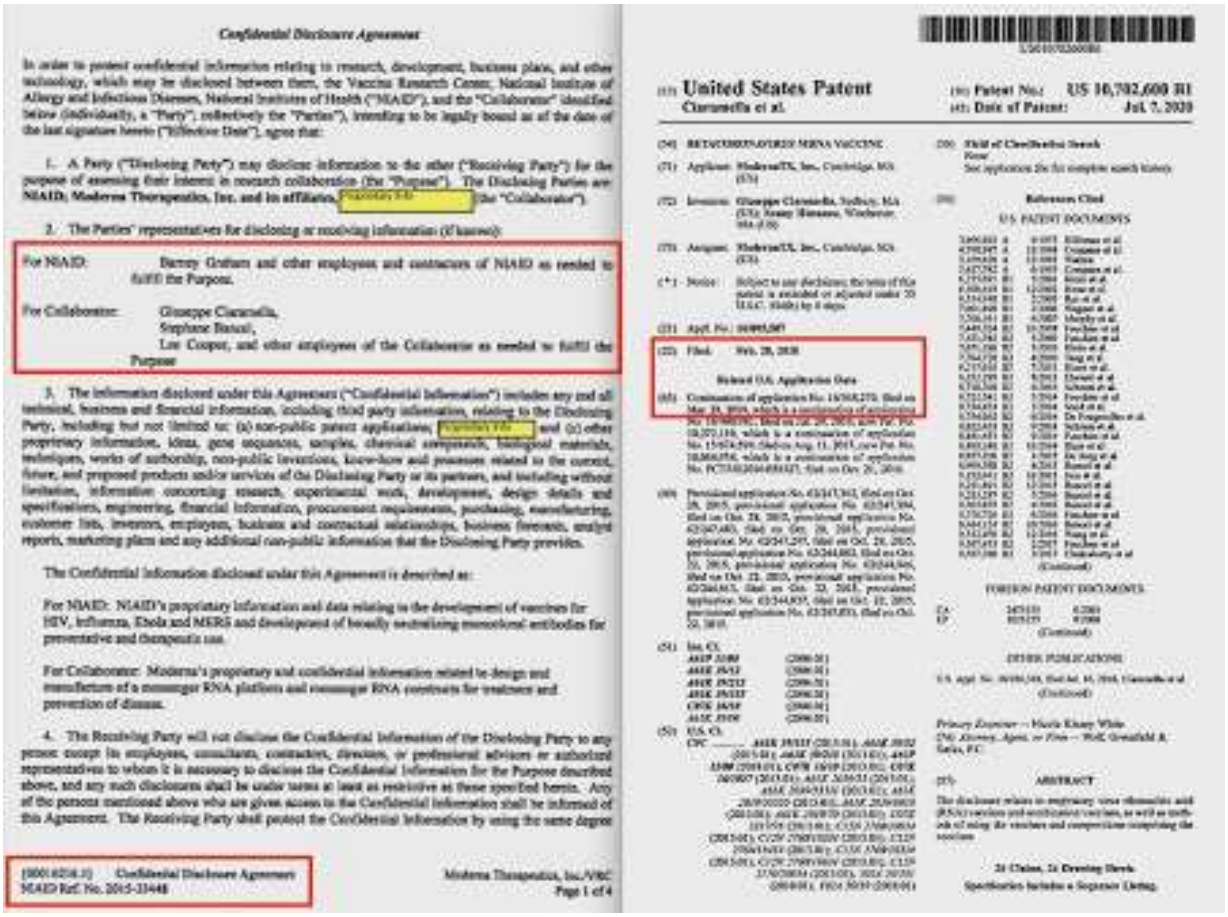
We have dedicated a long investigation to the mysterious and **dangerous biotechnologies with lipid nanoparticles** with extensive scientific documentation in relation to the Comirnaty messenger RNA gene serum ("heterozygous twin" of Spikevax), produced by the New Yorker Pfizer with the German Biontech: the first partner of the **London-based GSK run by a Microsoft director**, the second funded by the same IT tycoon Gates.

THE VACCINE PATENTED 9 MONTHS BEFORE THE PANDEMIC

In addition to the patents cited by the USPTO in their examination of '600, M·CAM has identified fourteen other issued patents preceding the '600 patent which were used by patent examiners to limit patents arising from the same funded research including patents sought by CureVac.

«In short, while Moderna enjoys hundreds of millions of dollars of funding allegiance and advocacy from Anthony Fauci and his NIAID, since its inception, it has been engaged in illegal patent activity and demonstrated contempt for U.S. Patent law. To make matters worse, the U.S. Government has given it financial backing in the face of undisclosed infringement risks potentially contributing to the very infringement for which they are indemnified» concludes the medical author of The Fauci / Covid-19 Dossier.

But now comes the explosive statement: «The specific claims addressing the pivot to the SARS Coronavirus were patented on March 28, 2019 – 9 months before the SARS CoV-2 outbreak! Both the patent and the DARPA funding for the technology were disclosed in scientific publication (New England Journal of Medicine) but the government funds were not acknowledged in the patent» is Dr. Martin's shocking revelation.



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For Collaborator: Giuseppe Ciaramella, Stephane Bancel, Lee Cooper, and other employees of the Collaborator as needed to fulfill the Purpose

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(800) 6218.1) Confidential Disclosure Agreement
NIAID Ref. No. 2015-31448

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(12) United States Patent
Ciaramella et al.

(10) Patent No.: US 10,702,600 B1
(45) Date of Patent: Jul. 7, 2020

(54) RETROVIRUS-DERIVED MRNA VACCINE

(71) Applicant: ModernaTX, Inc., Cambridge, MA (US)

(72) Inventors: Giuseppe Ciaramella, Sudbury, MA (US); Stephane Bancel, Winchester, MA (US)

(73) Assignee: ModernaTX, Inc., Cambridge, MA (US)

(*) Notice: Subject to any disclaimer, the term of this patent is extended or adjusted under 35 U.S.C. 154(b) by 0 days.

(21) Appl. No.: 16/805,587

(22) Filed: Feb. 28, 2020

Related U.S. Application Data

(63) Continuation of application No. 16/368,270, filed on Mar. 28, 2019, which is a continuation of application No. 16/040,981, filed on Jul. 20, 2018, now Pat. No. 10,272,150, which is a continuation of application No. 15/674,599, filed on Aug. 1, 2017, now Pat. No. 10,064,934, which is a continuation of application No. PCT/US2016/058327, filed on Oct. 21, 2016.

(60) Provisional application No. 62/247,362, filed on Oct. 28, 2015, provisional application No. 62/247,364, filed on Oct. 28, 2015, provisional application No. 62/247,465, filed on Oct. 28, 2015, provisional application No. 62/247,497, filed on Oct. 28, 2015, provisional application No. 62/244,863, filed on Oct. 22, 2015, provisional application No. 62/244,866, filed on Oct. 22, 2015, provisional application No. 62/244,930, filed on Oct. 22, 2015, provisional application No. 62/244,932, filed on Oct. 22, 2015, provisional application No. 62/245,853, filed on Oct. 23, 2015.

(51) Int. Cl.
A61P 31/00 (2006.01)
A61K 39/12 (2006.01)
A61K 39/215 (2006.01)
A61K 39/155 (2006.01)
C12N 15/63 (2006.01)
A61K 39/39 (2006.01)

(52) U.S. Cl.
CPC A61K 39/155 (2013.01); A61P 31/00 (2018.01); A61K 39/12 (2013.01); A61K 39/215 (2013.01); C12N 15/63 (2013.01); A61K 2039/5252 (2013.01); A61K 2039/5254 (2013.01); A61K 2039/5256 (2013.01); A61K 2039/5258 (2013.01); C12N 2760/16034 (2013.01); C12N 2760/16051 (2013.01); C12N 2760/16061 (2013.01); C12N 2760/16071 (2013.01); C12N 2760/16134 (2013.01); C12N 2760/16234 (2013.01); C12N 2760/16251 (2013.01); C12N 2760/16261 (2013.01); C12N 2760/16271 (2013.01); C12N 2760/16334 (2013.01); C12N 2760/16351 (2013.01); C12N 2760/16361 (2013.01); C12N 2760/16371 (2013.01); C12N 2760/16434 (2013.01); C12N 2760/16451 (2013.01); C12N 2760/16461 (2013.01); C12N 2760/16471 (2013.01); C12N 2760/16534 (2013.01); C12N 2760/16551 (2013.01); C12N 2760/16561 (2013.01); C12N 2760/16571 (2013.01); C12N 2760/16634 (2013.01); C12N 2760/16651 (2013.01); C12N 2760/16661 (2013.01); C12N 2760/16671 (2013.01); C12N 2760/16734 (2013.01); C12N 2760/16751 (2013.01); C12N 2760/16761 (2013.01); C12N 2760/16771 (2013.01); C12N 2760/16834 (2013.01); C12N 2760/16851 (2013.01); C12N 2760/16861 (2013.01); C12N 2760/16871 (2013.01); C12N 2760/16934 (2013.01); C12N 2760/16951 (2013.01); C12N 2760/16961 (2013.01); C12N 2760/16971 (2013.01); C12N 2760/17034 (2013.01); A61K 39/39 (2013.01)

(58) Field of Classification Search
CPC A61K 39/155; A61K 39/12; A61K 39/215; A61K 2039/5252; A61K 2039/5254; A61K 2039/5256; A61K 2039/5258; A61K 39/39; A61P 31/00; C12N 15/63; C12N 2760/16034; C12N 2760/16051; C12N 2760/16061; C12N 2760/16071; C12N 2760/16134; C12N 2760/16234; C12N 2760/16251; C12N 2760/16261; C12N 2760/16271; C12N 2760/16334; C12N 2760/16351; C12N 2760/16361; C12N 2760/16371; C12N 2760/16434; C12N 2760/16451; C12N 2760/16461; C12N 2760/16471; C12N 2760/16534; C12N 2760/16551; C12N 2760/16561; C12N 2760/16571; C12N 2760/16634; C12N 2760/16651; C12N 2760/16661; C12N 2760/16671; C12N 2760/16734; C12N 2760/16751; C12N 2760/16761; C12N 2760/16771; C12N 2760/16834; C12N 2760/16851; C12N 2760/16861; C12N 2760/16871; C12N 2760/16934; C12N 2760/16951; C12N 2760/16961; C12N 2760/16971; C12N 2760/17034; A61K 39/00; A61K 39/00; A61K 39/00

(56) References Cited

U.S. PATENT DOCUMENTS

(Continued)

FOREIGN PATENT DOCUMENTS

(Continued)

OTHER PUBLICATIONS

(Continued)

Primary Examiner — Maria Khuay White
(74) Attorney, Agent, or Firm — Wolf Greenfield & Sacks, P.C.

(57) ABSTRACT

The disclosure relates to respiratory virus ribonucleic acid (RNA) vaccines and methods for vaccines, as well as methods of using the vaccines and compositions comprising the vaccines

21 Claims, 24 Drawing Sheets

Specification includes a Sequence Listing.

The patent is updated on 7 July 2020 but was registered for the first time on 28 February 2020 but in continuation of the previous registration of 28 March 2019: the only one not to have changed the patent number (Pat. No.) as is see in the historical extract of the vaccine.

"Continuation of application No. 16 / 368,270, filed on Mar.28, 2019, which is a continuation of application No. 16 / 040,981, filed on Jul.20, 2018, now Pat.No. 10,272,150, which is a continuation of application No. 15 / 674,599, filed on Aug.1, 2017, now Pat.No.10,064,934, which is a continuation of application No.PCT / US2016 / 058327, filed on Oct.21, 2016 ".

THE EXCHANGE OF DOCUMENTS ON DECEMBER 2019

Analogues suspected on the existence of an experimental gene serum from Moderna before the official discovery of the Wuhan-Hu-1 virus, later identified as a new coronavirus initially named 2019-nCoV because the first cases were recorded in the populated city of Hubei precisely in that year, which was isolated and sequenced as Wuhan MN908947.1 on January 5, 2020.

The first to report the macroscopic anomaly was the British online investigative newspaper The Daily Exposé: «The first to report the macroscopic anomaly was the British online investigative newspaper The Daily Exposé. "A confidentiality agreement

shows that potential coronavirus vaccine candidates were transferred from Moderna to the University of North Carolina in 2019, nineteen days before the emergence of the alleged virus causing Covid-19 in Wuhan, China» wrote the British media on June 18, 2021.

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NIAID Ref. No. 2011-31948
Moderna Therapeutics, Inc/VRC
Page 1 of 4

MATERIAL TRANSFER AGREEMENT
SIGNATURE PAGE

FOR RECIPIENT:

Recipient's Investigator

Ralph Baric, PhD
Professor

Date: 12/12/2019

Mailing Address for Materials

Attention: Dr. Rachel Graham, Department of Epidemiology, University of North Carolina at Chapel Hill, 135 Dauer Drive, 2001 McGavran Greenberg Hall, CB #7435, Chapel Hill, NC 27599-7435

Tel: 919-966-3809 Fax: 919-962-0644

Duly Authorized

Jacqueline Quay
Director, Licensing & Innovation Support, OTC

Date: 12/16/19

Mailing Address for Notices

The University of North Carolina at Chapel Hill
Office of Technology Commercialization
109 Church Street, Chapel Hill, NC 27516

Tel: 919-966-3929 Fax: 919-962-0644

FOR PROVIDERS:

NIAID's Investigator

Barney Graham, MD, PhD

Date:

Duly Authorized

Amy F. Petrik-S

Amy Petrik, PhD
Technology Transfer Specialist, TTIPO, NIAID

Date:

Mailing Address for Notices

Technology Transfer and Intellectual Property Office
National Institute of Allergy and Infectious Diseases
Department of Health and Human Services
Suite 4D, MSC 9804
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Rockville, MD 20852
Tel: 301-496-2644 Fax: 240-627-3117

Moderna's Investigator

Stephane Bancel, PhD

Date: 12/17/2019

Duly Authorized

Shaun Ryan
Deputy General Counsel

Date: 12/17/19

Mailing Address for Notices

ModernaTX, Inc.
200 Technology Square
Cambridge, MA 02139
Attn: General Counsel

«The confidentially agreement (like at the bottom of the page) states that providers 'Moderna' alongside the 'National Institute of Allergy and Infectious Diseases' (NIAID) agreed to tranfer 'mRNA coronavirus vaccine candidates' developed and jointly-owned by NIAID and Moderna to recipients 'The Universisty of North Carolina at Chapel Hill' on the 12th December 2019» wrote the editorial staff of Exposé publishing the extracts on pages 105 and 108.

«The material transfer agreement was signed the December 12th 2019 by Ralph Baric, PhD, at the University of North Carolina at Chapel Hill (the microbiologist of the **laboratory chimeric SARS virus experiments** – ed), and then signed by Jacqueline Quay, Director of Licensing and Innovation Support at the University of North Carolina on December 16th 2019» adds the online newspaper, then asking inevitable questions…

This is taken from a long document. Read the rest here: **veteranstoday.com**

Dr Ricardo Delgado of La Quinta Columna shows us how to see if people around us have been nano-chipped by the vaxx.

TRANSCRIPT/TRANSLATION

Did you know that if you are vaccinated, you emit a MAC address that is visible on any mobile phone?

If you've received the jab, you've been branded like cattle.

Regardless of whether you've been branded into the herd of Pfizer, Astra Zeneca, Janssen or Moderna, your body now appears as a node on an online network that is discoverable via Bluetooth technology.

At La Quinta Columna, we need not convince you of anything. We want you to see this for yourself.

It is extremely easy and it will take less than a minute.

On an Android cellphone, go to the Play Store and download the free Bluetooth Scanner app.

Once downloaded, activate your phone's GPS and Bluetooth capabilities.

Now, open the Bluetooth Scanner app and choose "Search".

You will be able to see a list of MAC addresses, which appear as sets of 12-digit hexadecimal numbers, separated every two digits by a colon or a hyphen (i.e., an octet) and you will be able to distinguish those addresses that don't correspond with any appliances in your environment.

If you click to open each of these MAC addresses, you can even get an estimate of the physical distance to which those branded by this technology are from you.

If you have been vaxxed, one of these MAC addresses will correspond to you and if you drill down on it, you will see your own cellphone's call record.

If you find this shocking, wait until you experience the far-reaching consequences of being branded as cattle for the rest of your life.

Sadly, this is not a hoax.

They injected you with microtechnology and graphene, which is how this all works.

We have seen this same microtechnology in all brands of the vaxxines, from samples that we've analyzed, taken from all parts of the world.

They can monitor your specific biomedical data, the consequences of which go way beyond anything you can imagine now.

The mainstream media is hiding this abomination now being carried out against all human beings, including your children.

Visit LaQuintaColumna.info and LaQuintaColumna.tv and you will quickly understand the far-ranging implications of this intra-corporeal microtechnology and its humiliating purpose for all of humanity.

[For English translations of their work, go to Orwell.city].

****

Mik Andersen, publisher of the research blog Corona2Inspect is a renowned scientist now collaborating under a pseudonym with La Quinta Columna and Dr Pablo Campra Madrid.

The nano-network being deciphered and described by Andersen is one that would allow the neurostimulation of the population through a network designed for this purpose.

It appears that neuromodulation is the ultimate purpose of the global inoculation operation which has deployed highly-advanced military technology within the general population, 80% of whom still believe this is a vaccine.

Those who follow the investigations of La Quinta Columna know that, in addition to finding reduced graphene oxide (rGO), microparticles have been observed under the microscope that self-assemble with the appearance of chips, computer cards and strange fibers, similar to Morgellons but that also change over the course of several hours.

These revelations are so hellacious that people go into denial, because they cannot believe that governments, doctors, journalists, etc have agreed to carry out the most diabolical fraud in human history by fulfilling the Globalists' long-announced goal of microchipping the human population.

Running Time: 2 mins Video Link just below

Moderna Seeks Approval from FDA and European Medicines Agency (EMA) to Start Injecting Children Under 6 with mRNA COVID-19 Vaccines

Posted By *AdminM* On March 24, 2022

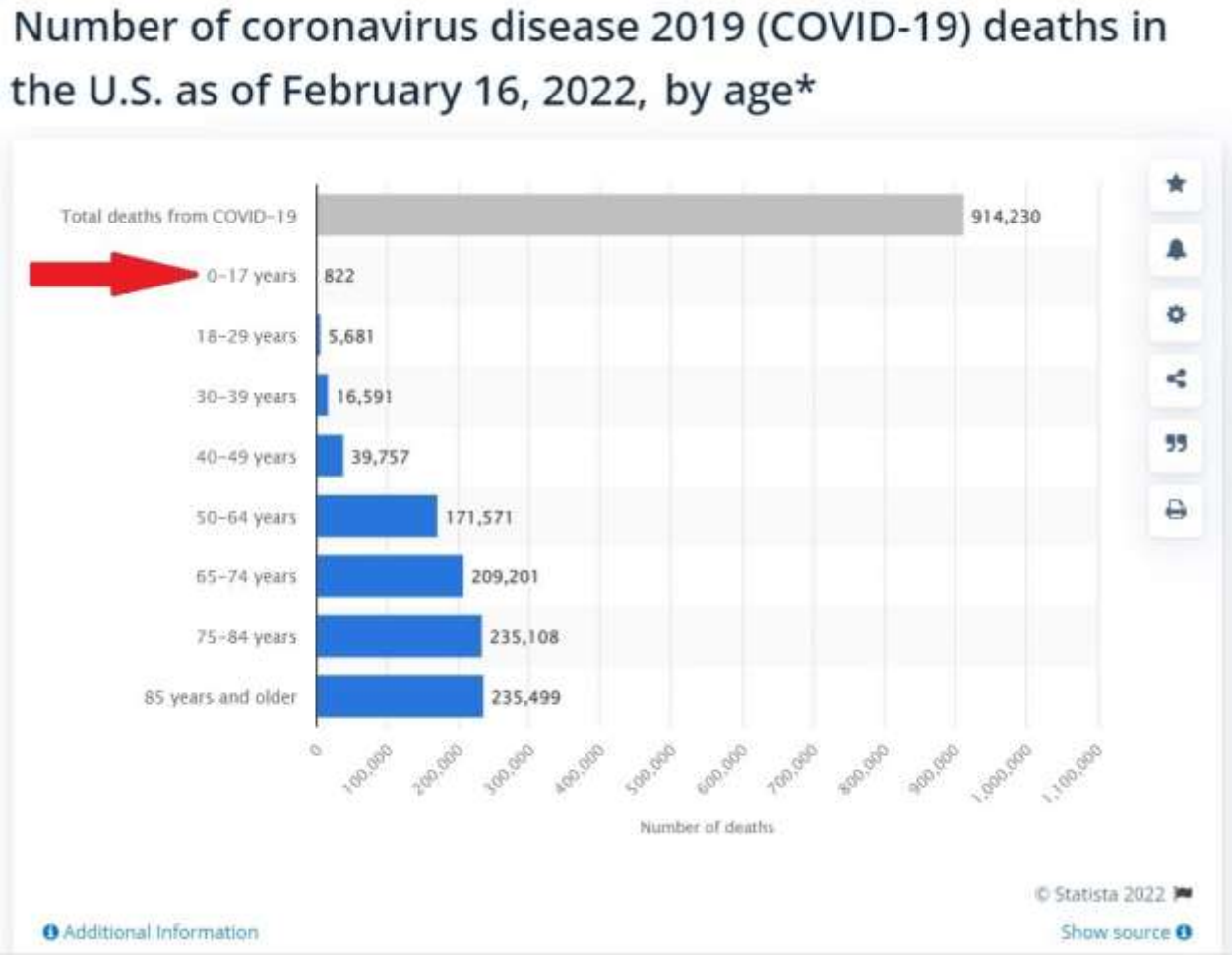
by Brian Shilhavy
Editor, Health Impact News

Fierce Pharma **reported yesterday** [1] **that Moderna is seeking approval for their COVID-19 mRNA vaccine to be injected into children under the age of 6, from both the U.S. FDA and the European EMA. They are also asking the FDA to approve their vaccine for children between the ages of 6 and 11.**

Moderna has new data backing the use of its COVID-19 vaccine Spikevax at a low dose in the youngest children under the age of 6, and it's moving forward with global regulatory submissions.

Moderna will ask the FDA and the European Medicines Agency (EMA) "in the coming weeks" to authorize Spikevax for children 6 months to under 6 years of age, the company said Wednesday. Simultaneously, Moderna has started a delayed FDA filing for emergency use authorization of the mRNA vaccine in children ages 6 to 11, CEO Stéphane Bancel said in a statement. (Source [1]**.)**

Using government official statistics, children in this age group have almost a zero percent chance of dying from COVID-19, as we have previously shown in numerous articles.



Source [2].

And yet, the U.S. *Vaccine Adverse Events Reporting System* (VAERS) is reporting almost 45,000 injuries and deaths in this age group following COVID-19 vaccines, and we know that this is severely under-reported. (Source [3].)

From the 3/11/2022 release of VAERS data:

Found 44,821 cases where Age is under-18 and Vaccine is COVID19

Table

Event Outcome	Count	Percent
Death	95	0.21%
Permanent Disability	396	0.88%
Office Visit	6,296	14.05%
Emergency Room	9	0.02%
Emergency Doctor/Room	4,619	10.31%
Hospitalized	3,296	7.35%
Hospitalized, Prolonged	6	0.01%
Recovered	18,566	41.42%
Birth Defect	12	0.03%
Life Threatening	550	1.23%
Not Serious	18,474	41.22%
TOTAL	† 52,319	† 116.73%

† Because some cases have multiple vaccinations and symptoms, a single case can account for multiple entries in this table. This is the reason why the Total Count is greater than 44821 (the number of cases found), and the Total Percentage is greater than 100.

Also, the CDC recently removed nearly a quarter of the deaths it was previously reporting for children under the age of 18 related to COVID-19.

The Centers for Disease Control and Prevention (CDC) has removed tens of thousands of deaths linked to COVID-19, including nearly a quarter of deaths it had listed in those under 18 years old.

The health agency quietly made the change on its data tracker website [4] on March 15.

***"Data on deaths were adjusted after resolving a coding logic error. This resulted in decreased death counts across all demographic categories,"* the CDC says on the site.**

The CDC relies on states and other jurisdictions to report COVID-19 deaths and acknowledges on its website that the data is not complete.

But the statistics are often cited by doctors and others when pushing for COVID-19 vaccination, including figures who believe virtually all children should be vaccinated. Dr. Rochelle Walensky, the CDC's director, cited the tracker's death total in November 2021 while pushing for an expert panel to advise her agency to recommend vaccination for all children 5- to 11-years-old.

Before the change, the CDC listed 1,755 children as dying from COVID-19 along with approximately 851,000 others, according to Kelley Krohnert, a Georgia resident who has been tracking the updates.

The update saw the CDC cut 416 deaths among children and over 71,000 elsewhere, arriving at a total of just under 780,000.

The agency declined to provide a comment by deadline. (Source [5].)

This begs the question then as to why the FDA would even consider a COVID-19 shot for this age group (I think we all know the answer to that question), and why parents would take the risk of injuring or killing their child with one of these experimental shots?

This is simply the same demonic spirit we have seen throughout history that moves people to sacrifice their children to "gods" like Molech and Baal in ancient times, and is now being manifest in "modern" culture through the vaccine cult.

Here are a few tragic stories that have been posted to Telegram within the past few days of parents who sacrificed their children to the vaccine gods and now have either a dead child that they have to bury, or one disabled for the rest of their life.

Evidence has emerged which proves beyond a reasonable doubt that the Covid-19 virus was created by the very pharmaceutical giant that has made billions through the sale of an experimental Covid-19 injection; Moderna. By Steve Kirsch.

On February 23 the Daily Mail ran an article showing that Moderna has patented the 19 base letter (nucleotide) sequence which codes for the Furin Cleavage site in Covid-19.

They cited a Paper by Scientists in India, Switzerland, Italy and the US (cautiously entitled: MSH3 Homology and Potential Recombination Link to SARS-CoV-2 Furin Cleavage Site) in which they calculated that the chances of a 19 nucleotide sequence patented by Moderna randomly appearing in Covid-19 in circumstances where it does not appear anywhere else in nature are 1 in 3 trillion.

But they failed to make the obvious deduction there from. Had they made said obvious deduction I fear that might have been the last scientific deduction they ever got published!

They decided to investigate the RNA sequence for the Furin cleavage site in the Covid-19 Spike Protein to see if it occurred anywhere else in nature. .

Fortunately the NCBI/NIH have produced the wonderful BLAST database which catalogues every gene sequence in nature known to man and every synthetic patented gene sequence known to the patent office.

The researchers chose the Furin Cleavage sequence because it is the only continuous gene letter sequence (nucleotide sequence) in Covid-19 with more than 3 nucleotides, that differs from the respective letters in its closest natural relative the Bat Coronavirus RaTG13 (all other differences are 3 letters or less long). So it was by far the best candidate for determining whether or not Covid-19 was man made.

The reader might consider it more likely that a Furin Cleavage Site would appear in the Sun than in the Daily Mail. But this cleavage refers to the separation of spike from virus rather than pillow from pillow.

Furthermore the Furin Cleavage Site is key to the pathogenicity of Covid-19. So if there was to be some man made gain of function included in the virus, this is where one might expect to find it.

The Amino Acid sequence of the Furin Cleavage Site is PRRA (Proline Argenine Argenine Alanine). Each Amino Acid is coded for by a Codon, consisting of 3 nucleotides (genetic sequence letters). So all the differences in the genetic code between Covid-19 and RaTG13 are at most one Codon long, one amino acid long, other than the Furin Cleavage Sequence, which is…

CCT CGG CGG GCA

The complimentary sequence (the opposing DNA strand of the double helix is (GGAGCCGCCCGT) because C binds with G and A binds with T

The reverse compliment (the same thing written backwards) is therefore TGCCCGCCGAGG

The researchers did a BLAST (Basic Local Alignment Search Tool) alignment search (which means they search for the gene sequence, the reverse gene sequence, the complimentary gene sequence and the reverse complimentary gene sequence) through every gene sequence in nature known to man for CTCCTCGGCGGGCACGTAG which is the 19 nucleotide sequence containing the Furin Cleavage Sequence, which also appears in Covid-19, and which is found

actually in the reverse compliment form CTACGTGCCCGCCGAGGAG patented by Moderna.

Their search results can be found here.

Table 1 shows that it does exist in the 5 U.S. patents cited below…
US9149506B2: Modified polynucleotides encoding septin-4 –
https://patents.google.com/patent/US9149506B2/en

Inventor: Tirtha Chakraborty, Antonin de Fougerolles
Current Assignee: ModernaTx Inc

2012-04-02 Priority to US201261618953P
2013-12-16 Application filed by Moderna Therapeutics Inc
2014-05-22 Publication of US20140141067A1
2015-10-06 Publication of US9149506B2
2015-10-06 Application granted
2020-01-10 First worldwide family litigation filed

US9216205B2: Modified polynucleotides encoding granulysin –
https://patents.google.com/patent/US9216205B2/en

Inventor: Tirtha Chakraborty, Antonin de Fougerolles
Current Assignee: ModernaTx Inc

2012-04-02 Priority to US201261618873P
2013-12-16 Application filed by Moderna Therapeutics Inc
2014-04-24 Publication of US20140113960A1
2015-12-22 Publication of US9216205B2
2015-12-22 Application granted

US9255129B2: Modified polynucleotides encoding SIAH E3 ubiquitin protein ligase 1 – https://patents.google.com/patent/US9255129B2/en

Inventor: Tirtha Chakraborty, Antonin de Fougerolles
Current Assignee: ModernaTx Inc

2012-04-02 Priority to US201261618868P
2013-12-16 Application filed by Moderna Therapeutics Inc
2014-05-22 Publication of US20140141068A1
2016-02-09 Application granted
2016-02-09 Publication of US9255129B2

US9301993B2: Modified polynucleotides encoding apoptosis inducing factor 1 –
https://patents.google.com/patent/US9301993B2/en

Inventor: Tirtha Chakraborty, Antonin de Fougerolles
Current Assignee: ModernaTx Inc

2012-04-02 Priority to US201261618957P
2013-12-16 Application filed by Moderna Therapeutics Inc
2014-04-17 Publication of US20140107189A1
2016-04-05 Application granted
2016-04-05 Publication of US9301993B2
2020-01-10 First worldwide family litigation filed
US9587003B2: Modified polynucleotides for the production of oncology-related proteins and peptides – https://patents.google.com/patent/US9587003B2/en

Inventor: Stephane Bancel, Tirtha Chakraborty, Antonin de Fougerolles, Sayda M. Elbashir, Matthias John, Atanu Roy, Susan Whoriskey, Kristy M. Wood, Paul Hatala, Jason P. Schrum, Kenechi Ejebe, Jeff Lynn Ellsworth, Justin Guild

Current Assignee: ModernaTx Inc

2012-04-02 Priority to US201261618868P
2016-02-04 Application filed by ModernaTx Inc
2016-06-02 Publication of US20160152678A1
2017-03-07 Publication of US9587003B2
2017-03-07 Application granted

So Moderna first applied for a patent for the 19 nucleotide sequence in 2013 on December 16. Perhaps December 25 would have been more appropriate since it was destined to become the Crown of Thorns of Mathew 27, Mark 15 and John 19

Table2: Shows that the sequence occurs in Covid-19 from nucleotide 23601 to 23619.

Table3: Shows that this gene sequence does not exist in nature (but 14 nucleotide parts of it do).

I decided to check their work. Yes. I fact checked them (I will send an invoice to the globalists). This turned out to be a bit of an epic journey. The Google patent page for US9587003B2 does not contain the gene sequence. The pdf of the patent does not contain the gene sequence and is not searchable from pages 101-304. But it does have a link to a lengthy 'Sequence Listing" section which link one cannot copy. So I manually transcribed it in my fair hand – http://seqdata.uspto.gov/?pageRequest=docDetail&DocID=US09587003B2

From that page you can enter the Sequence ID quoted in the paper as 11652 and get to https://seqdata.uspto.gov/?pageRequest=viewSequence&DocID=US09587003B

2&seqID=11652 which has the following at Nucleotides 2751-2733 reading backwards…

CTACGTGCCCGCCGAGGAG patented by Moderna is the reverse compliment of CTCCTCGGCGGGCACGTAG, the 19 nucleotide sequence which appears in Covid-19 DNA from nucleotide 23601-23619 (which would therefore be covered by their patent).

Likewise you can search for the sequence in US9149506B2 by going to https://seqdata.uspto.gov/?pageRequest=viewSequence&DocID=US09149506B2&seqID=11652, whereupon you will find the same thing again

I then searched the gene sequence of Wuhan Hu1 (alpha) at https://www.ncbi.nlm.nih.gov/nuccore/NC_045512 and found

Which has the 19 nucleotide sequence CTCCTCGGCGGGCACGTAG from 23601-23619 as described in table 3.

I then ran my own non-aligned blast search of all patented gene sequences for the reverse compliment directly (or perhaps for a back handed compliment) and got the same results as the researchers

And the same for the other 3 US patents.

So I can confirm, and the reader can confirm using the links above, that Moderna did apply for a Patent not only on the reverse compliment of the 12 nucleotide Furin Cleavage Site in Covid-19 but actually on the 19 nucleotide sequence containing it as described above.

Furthermore they did not merely apply for a patent on 2016 February 4 with US9587003B2: as reported in the Daily Mail. They actually applied on 2013 December 16 for 4 patents with US9149506B2, US9216205B2, US9255129B2, US9301993B2:as well.

So Moderna had developed the 19 nucleotide gene sequence containing the Furin Cleavage Site which gives Covid19 its infectivity to humans by patented gain of function research as early as 2013, 6 years before the Wuhan outbreak took place. Not 3 as reported in the Mail and virally elsewhere..

So now we look at the chances of this occurring naturally. The paper calculates the probability of this particular 19 nucleotide sequence occurring randomly in a 30,000 nucleotide virus as

$(30,000-18) \times (1/4)^{19} = 1.09 \times 10^{-7}$

Which is correct because there are 30,000-18 places to start the sequence given that it needs a further 18 more letters to complete it. But there are actually 29,904 nucleotides in Wuhan HU1 (alpha). So a more accurate calculation would be
$(29,904-18) \times (1/4)^{19} = 1.087 \times 10^{-7}$

Then they calculate the chances that the 19 nucleotide sequence occurs in the patented library of 24,712 sequences with a mean length of 3300 nucleotides. But that calculation is irrelevant because the sequence did not randomly appear in 5 Moderna Patent applications. The sequence was known to code for a Furin Cleavage Site, which is known to provide gain of function to Coronaviruses.

It was put there deliberately and patented due to its infecting power in humans, which we shall see, later in the article, results from the normal viral Arginine (R) codon AGA (used in 45% of viral Arginine codons) being replaced by the human Arginine codon CGG (used in 0% of viral Arginine codons) in the furin cleavage site.

All we are tying to work out here is what the chances are of a 19 nucleotide sequence patented by Moderna turning up in Covid-19 through natural causes, the natural mutations of Bat Coronavirus RaTG13 or some other virus.

The nucleotides form Codons which are triplets. So there are 64 possible triplets of the 4 DNA nucleotides ACGT (4x4x4 = 64). But all triplets do occur. 61 code for 20 amino acids redundantly and 3 are stop codons which tell the ribosome to stop making the protein.

But things are not this simple because the Furin cleavage site appears in the spike protein where it needs to be and the spike protein only has 1273×3 = 3819 nucleotides. The chances of the 19 nucleotide Furin Cleavage sequence appearing in the spike protein are
$(3,819-18) \times (1/4)^{19} = 1.389 \times 10^{-8}$

Or 1 in 72 million. So those would be the chances that one particular variant, say the first Covid-19 variant, had the 19 nucleotide sequence in the right place (the spike). And it did. So certainly by the balance of probabilities, and certainly beyond a reasonable doubt (1 in 72 million being an unreasonable doubt) Moderna made Covid-19.

100% Biochemic Proof that Covid19 was Man Made

The Double CGG Codon used in the Moderna Specific Furin Cleavage site does not occur in any other Furin cleavage site in any other virus in nature. Furin cleavage sites do occur in other viruses but NOT at all in other betacoronaviruses like Covid-19 and NOT at all with the double CGG codon.

Arginine (R), can be encoded by any of the 6 triplets: AGG, AGA, CGA, CGC, CGG, CGT. In Covid-19, the furin site (PRRA), has 12 nucleotides (3 x 4). In Covid-19, the RR doublet of the furin site is encoded by CGG-CGG.

Two Biochemists Prof Antonio R. Romeu and Assistant Prof Enric Ollé analysed the RR doublet from a large sample of furin cleavage sites of several kinds of viruses. They found that there were no RR doublets encoded by the CGG-CGG codons in any virus in nature. They observed that the AGA triplet was the majority codon involved in these viral RR doublets.

In all genetic recombination (where a part of one genome merges with another genome), the donor code is passed to the acceptor. But there is simply NO KNOWN VIRUS with a Moderna Specific Furin Cleavage Site (having the CGG-CGG codon pair) that exists to donate a Moderna Specific furin cleavage site to Covid19. So the only way that sequence could get into Covid-19 is from Moderna. Moderna was the donor. Nature was not. QED. Case Closed..
But it gets worse.

The Spanish Profs decided to analyse the arginine codon usage in every single protein in Covid-19. The found the following…
AGG (13%)
AGA (45%)
CGA (5%)
CGC (10%)
CGG (3%)
CGT (24%).

So the AGA codon triplet was the majority, and interestingly, CGG was the minority codon for Arginine in the virus.

But it gets worse still.

In the specific case of S protein, of the 42 Arginines (R) it has, 20 are encoded by AGA, and only 2 by CGG. These 2 of course, are the two in the Moderna Specific Furin Cleavage Site.

So the only Arginine in the spike protein that is encoded a la Moderna are in the Furin Cleavage site. The other 40 instances do not use CGG at all.

They then go on to comment that each individual species in nature has its own codon preferences. Obviously viruses like AGA, and do not like CGG at all, in nature.

But guess which species does use CGG for Arginine more than the other 5 competing codons – yes its jolly old homo sapiens. Our coding preferences for Arginine are

AGG (20%)
AGA (20%)
CGA (11%)
CGC (19%)
CGG (21%)
CGT (9%).

So the CGG codon in the furin cleavage site WILL have come about through Chimeric (human animal combination) gain of function research.

Could Somebody other than Moderna have made Covid-19 using the Moderna Specific Furin Cleavage Site?

"New documents show that just 18 months before the first Covid-19 cases appeared, researchers had submitted plans to release skin-penetrating nanoparticles and aerosols containing "novel chimeric spike proteins" of bat coronaviruses into cave bats in Yunnan, China. They also planned to create chimeric viruses, genetically enhanced to infect humans more easily, and requested $14 million from the Defense Advanced Research Projects Agency (Darpa) to fund the work.

Papers, confirmed as genuine by a former member of the Trump administration, show they were hoping to introduce "human-specific cleavage sites" to bat coronaviruses which would make it easier for the virus to enter human cells.

When Covid-19 was first genetically sequenced, scientists were puzzled about how the virus had evolved such a human-specific adaptation at the cleavage site on the spike protein, which is the reason it is so infectious." – the Telegraph

I can see all of the great journalists at the Daily Mail and the Telegraph (not to mention scientists around the world) doing all of this research into Covid19 and reaching the inevitable logical conclusion that there was either an accidental or a deliberate lab leak and then having to word their conclusions in such a way as to label that strong probablity as a weak possibility.

But here above we have proved it as a fact (since the Moderna Specific Furin Cleavage Sequence CGG codon does not occur in any furin cleavage site in any natural virus and therefore it cannot have been the result of natural genetic recombination. So it has to be the result of man made genetic insertion.

In theory a further party involved with the NAIAD or the NIH could have used the furin cleavage site patented by Moderna and made Covid19 themselves. This would not have broken any patent of Moderna. The Furin cleavage site itself is not patentable having been known since at least 2004

US7223390B2: Insertion of furin protease cleavage sites in membrane proteins and uses thereof
2004-05-07 Application filed by Research Development Foundation
2004-11-11 Publication of US20040224391A1
2007-05-29 Application granted

Although Moderna could actually have patented the Moderna Specific (CGG for AGA) encoding of the furin cleavage site which was is not known in nature even today (if we accept that Covid-19 is man made).

But given that the lab leak (deliberate or accidental) came from Wuhan, and given the Chinese cover up and given the Fauci denials exposed by Senator Rand Paul, and given the NIH, NIAID cover ups and the US Intelligence services cover up, when their 3 month long report into the origin of Covid-19 ordered by presidential impersonator Biden yielded nothing, and given the relationships between the NIAID, the NIH, the WIV, the EcoHealth Alliance, the University of North Carolina and Moderna, I cannot see any room for anyone else.

Furthermore the entire unholy cabal of bad actors started developing the Moderna Vaccine before the pandemic struck – https://www.infowars.com/posts/must-watch-nih-claimed-joint-ownership-of-moderna-mrna-vaccine-began-development-weeks-ahead-of-pandemic/

But things are not as simple as that because nature has had certainly 100,000 years to make human viruses and it never once put a Moderna specific (CGG for AGA) furin cleavage site into anything, nor did it put the 19 nucleotide sequence in anything before.

Yet within 6 years of Moderna patenting it, we find it in Covid-19 in circumstances where Moderna is working with that virus. So just there the probability is not 100,000 to 6 or 16,666 to 1 that Moderna is responsible rather than nature. No it is 100% because nature has not done it. It never has and there is no evidence that it ever will.

It is man the mixes up human and viral Arginine codons not nature.

Prof. Luc Montagnier spent the last years of his life proving that COVID-19 was man made and containing much of the HIV1 genetic code

Prof. Luc Montagnier, before he died on February 8, 2022 did a total assassination of the concept that Covid-19 evolved naturally by showing that it had massive equivalence to HIV. The diagram below shows a 275 nucleotide region of Covid-19 which has 200 nucleotides from HIV/SIV (Simian ImmunoVirus) in it. And remember there are 61 codons specifying 20 amino acids. So one can say the same thing in on average 3 different ways with codons.

You can download a pdf of his study here and the supplementary materials here. It is very technical. But he did win the Nobel prize for discovering the HIV virus. So if anyone would know if Covid had been boosted with HIV, it would be him. He pointed out that Covid-19 was man made early (March 2020) in the pandemic and was himself assassinated by the press and the fact checkers as a result. Every single fact checker who attacked him was wrong.

There was no scientific basis to any of their fact checking. These outfits are not fact checkers at all of course. They are globalist disinformation agencies, sons of Goebbels, fact chuckers and science deniers. They are about as trustworthy as an American election. I can check a fact for myself thank you very much. I don't need a brainwashed woke madrassa student telling me their opinion about a subject that they never studied at University.

Since we have proven beyond a reasonable doubt (beyond a 1 in 72 million doubt statically and with 100% certainly biochemically from the Moderna Specific Furin Cleavage Site) that Moderna made Covid-19. And since Moderna and Fauci have not admitted to having made it and have in fact covered up evidence to that effect, it may be the case that they are hiding something else as well.

Because the only two theories now left are the accidental lab leak theory and the deliberate lab leak theory. I mean the vast majority of political leaks are not accidents. They are deliberate strategies to provide advantage to the leaker or his paymaster. It is well known in the IT industry that viruses appear when antivirus sales are needed. Why would things be any different with human viruses, now that they can be man-made too? Especially when you consider the massive role of Bill Gates and his foundation and GAVI and GVAP in he global vaccination business.

The only reason that Moderna would make Covid-19 is to release it. Otherwise the entire exercise would be financially futile, commercially pointless

The reason adduced by Fauci for doing gain of function research is that man needs to be ahead of nature or bad actors in order to have a vaccine in good time if a disease mutates or is genetically modified by the Chinese or the Russians to be lethal.

But in order to believe that one has to believe that Moderna are interested in the saving people's lives. I am sorry. All their actions show to me that they are interested in vaccinating people knowing how likely that is to cost them their lives.

They are interested in profit, the profit that comes from a pandemic. They are not saviors of mankind as they represent. They are our exploiters and our abusers.

They produced the virus in order to leak it, in order to pose as our saviours from their own leak. These are not the activities of a savior figure. Luc Montagnier was trying to be our savior from them and he was assassinated (professionally) by their groupies. Moderna were doing gain of function research in order to release the virus and force a vaccine for it in a manner which would maximize their profits. That is not a conspiracy theory. It is what happened precisely. Their share price went up by 20x.

They released it in order to sell their vaccines and to destroy the immune systems of their customers because our immune systems reduce their profits. That is Big Pharma business.

The reason that the writer is so confident that Moderna or their agents made and leaked Covid-19 and the reason I called it as such at the start of the pandemic to almost as much ridicule as Prof Montagnier received (God bless him) is that the scriptures say in Matthew 27, Mark 15 and John 19 that.

[29] And they (the soldiers of the governor of verse [27]) platted a crown of thorns and put it upon his head, and a reed in his right hand; and they kneeled down before him, and mocked him, saying, Hail, King of the Jews! [30] And they spat upon him, and took the reed and smote him on the head. (Matthew 27 ASV)

May I therefore beg your indulgence whilst I interpret these words:

The US department of defense funded the gene splicing of the Coronavirus of Spike Proteins (Covid-19) through NIH and NIAID and DARPA which first infected Jesus, through his fiance, the New Covenant Saints, just after he became the secular King, Caesar to those saints, the antitypical Jews, those covenanted to be angelic sons of Jacob, the born against angelically.

We calculated that the malediction which prevented Jesus becoming Caesar to the saints ended in 2019 Tishri 15 (October 17/18). Glenn Beck did a documentary showing that 10 hospitals in Wuhan took cases with Covid19 symptoms in October 2019. Yes Folks. Covid-19 is a proof that Jesus is now secular King over the saints, the antitypical Jews, the Jews by angelic salvation covenant, at the least.

But then the soldiers spat upon him. For that is how Covid19 is transferred, through small aerosol droplets exhaled out of the mouth. The soldiers deliberately spat upon him. It was not a SALIVA LEAK! They smote Jesus on the head because the saints are the head of the church and they caught Covid19 not by random chance infection but by a deliberate smiting with a read, a biological weapon, a deliberate weaponised attack. For more on this see here.

So what Prof Montagnier saw with his virology expertise, I saw with my theological expertise. Showing that whilst fact checkers and science are mutually exclusive, science and theology actually agree, when properly understood (and that is one big caveat). Prof M taught us that the vaccines cause the variants. Indeed basic virology forbids mass vaccination during a pandemic for that very reason. He said the curve of deaths follows the curve of vaccinations. Mind you, paradoxically, if the vaccines caused Omicron, then they saved us from themselves!

The Time has Come to hold People and Organizations to Account

The Covid19 makers, the genetic vaccine makers. their funders and their promoters, which include almost every government and public sector and health service in the world, are therefore guilty of Genocide and crimes against humanity. They have pushed genetic rape and sickness and death onto half of the population of the world in order to enrich the pockets of Pharmaceutical Companies. Governments and Public sectors around the world have abandoned their health service regulation to billionaires and heartless corporations

In the UK, all of the income tax we pay goes to the health service and all of its protocols are determined by its regulators and all of its regulators are controlled and funded by Big Pharma who seek to damage then manage our health for their profit.

So every penny we spend in income tax brings us one step closer to sickness, to death and to drug dependency.

So why did Prof Montagnier choose to spend the last years of his life proving that Covid-19 was man made and that the spike proteins, and therefore the vaccines, were an existential threat to the species? What did he have left to prove to himself or to anybody else at 87-89? He certainly did not do it to increase his reputation in the profession.

No, he was driven by the same passion that drove him to discover HIV. A passion to SAVE mankind from viruses and those who would engineer them to damage us. And why did he give up the ghost in February 2022? Because he knew that Omicron had the vaccines beat. His job was done by a greater virologist even than him. He could therefore rest in peace and go see some people who understood the magnitude of his contribution.

Covid-19 was not made in 2019. It was made from the 19 nucleotide Moderna specific chimeric (CGG for AGA) furin cleavage site which does not occur anywhere in nature. And every Covid death and every Covid vaccine death is parked squarely on the doorstep of ModeRNA waiting for justice.

But we shall not execute that justice fast enough. And therefore the final plague upon mankind of Revelation 6:8, delivered by the 4th horseman of the apocalypse, which plague Bill Gates himself has prophesied, will arrive later this year (after War and after Famine, the 2nd and 3rd horsemen).

Going forward the balance of this article is from Dr. Luc Montagnier's original report. In a previous segment of this series, I shared the genome location of the inserts of HIV1, HIV2, and SARS2. Without question, Dr. Luc Montagnier is the true hero in all of this crime against humanity.

"SARSCoV-2/COVID-19 is a novel coronavirus characterized as an unusual viral pneumonia. COVID-19 contains a single-stranded (positive-sense) mRNA associated with a nucleoprotein within a capsid comprised of a matrix protein."
-(National Library of Medicine)

"A digital ID is the electronic equivalent of an individual's identity card. It is a way to provide verified personally identifying information of an individual for a software to read and process. Both online and offline environments can adopt a digital identity. It can also act as a key by storing and deploying permission." – (World Economic Forum (WEF)

Today's synthetic biology is driven by a combination of quantum computing and artificial intelligence. The most obvious example being the SARS-CoV-2 "virus" - in quotations specifically because of its synthesis in a laboratory, imitating a "wild", or naturally occurring virus. Since January of 2020, we have provided numerous peer reviewed papers, patents, and journal articles as evidence of the research leading up to and including insilico development of this synthetic "virus". Below, please find a link to bioinformatic evidence from the published sequence of this m RNA organism. (Prashang et al, Uncanny similarity of unique inserts in the 2019-nCoV spike protein to HIV-1 gp120 and Gag | bioRxiv) Wuhan market ID: LR757998.1).

The research team of Professor Luc Montagnier and Jean Claude Perez, PhD, published the identity of the inserts in their paper entitled: "COVID-19, SARS and Bats Coronaviruses Genomes Unexpected Exogenous RNA Sequences". The following excerpts provide evidence of their findings: - 4 HIV1 HIV2 Exogenous Informative Elements radically distinguishes all COVID-19 strains from all SARS and Bat strains. - Validation of nucleotide fragments as « Exogenous Informative Elements » (EIE): We have chosen this minimal length of 18 nucleotides (6 amino acids) for the support of information (thus as an antigenic motif). This is also the size of the primers used for PCR which allows high specificity of sequence selection on DNA recognition.

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This article shows how 16 fragments (Env Pol and Integrase genes) from different strains, both diversified and very recent, of the HIV1, HIV2 and SIV retroviruses most likely are present into the genome of COVID-19. Among these fragments, 12 are concentrated in a very small region of the COVID-19 genome, length less than 900 bases, i.e. less than 3% of the total length of this genome.

In addition, these footprints are positioned in 2 functional genes of COVID-19: the or f1ab and Sspike genes. To sum up, here are the two main facts which contribute to our hypothesis of a partially synthetic genome: A contiguous region representing 2.49% of the whole COVID-19 genome of which 40.99% is made up of 12 diverse fragments originating from various strains of HIV SIV retroviruses. On the other hand, these 12 fragments some of which appear concatenated.

Notably, the retroviral part of these regions, which consists of 8 elements from various strains HIV1, HIV2 and SIV covers a length of 275 contiguous bases of COVID-19. The cumulative length of these 8 HIV SIV elements represents 200 bases. Consequently, the HIV SIV density rate of this region of COVID-19 is 200/ 275 = 72.73%, which is considerable. Moreover each of these elements is made of 18 or more nucleotides and therefore may have function. They are called Exogenous Informative Elements.

Summary of Findings

1. **18 RNA fragments of homology equal or more than 80% with human or simian retroviruses have been found in the COVID_19 genome.**

2. **These fragments are 18 to 30 nucleotides long and therefore have the potential to modify the gene expression of Covid19. We have named them external Inform ative Elements or EIE.**

3. **These EIE are not dispersed randomly, but are concentrated in a small part of the genome.**

4. **Among this part, a 225 nucleotide long region is unique to COVID_19 and Bat RaTG13 and can discriminate and formally distinguish these 2 genomes.**

5. **In the decreasing slope of the epidemic, this region exhibits an abnormally high rate of mutations/deletions**

6. **The comparative analysis of the SPIKES genes of COVID_19 and Bat RaTG13 demonstrates two abnormal facts: on the one hand, the insertion of 4 contiguous amino acids in the middle of SPIKE, on the other hand, an abnormal distribution of synonymous codons in the second half of SPIKE. Finally the insertion in this region of an EIE coming from a Plasmodium Yoelii gene is demonstrated, but above all seems to explain the "strategy" pursued by having "artificially" modified the ratio of synonym codons / non-synonymous codons in this same region of 1770 COVID_19 SPIKEnucleot ides.**

The above information comes from two different sources which confirm with 100% certainty that COVID-19 is a MAN-MADE virus, and it can be proved that MODERNA Pharmaceuticals developed it and patented the Retrovirus in 2013.

This information confirms that anyone vaxxed should immediately be tested for HIV1, HIV2, or SIV2. To neglect to do so puts one at great risk of going untreated for these pathogens. To go untreated is to allow the vaxx to degrade your natural immune system until death occurs. HIV does not mean a death sentence if treated immediately. The NBA basketball player Magic Johnson has lived with HIV for over 30 years with proper treatment!

Blessings,

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http://jesusisthewaythetruththelife.com/node/22