Part 1: Dismantling COVID-19 Deceptions: The 'Novel' Coronavirus Needs Lipid Nanoparticles to Infect Humans

The COVID-19 pandemic was not caused by the SARS-CoV-2 virus or variants. COVID-19 is a new kind of Ai bioweapon that is part technology, part biology; and it's intelligent.

Karen Kingston
Oct 12

What exactly caused the COVID-19 pandemic? Was it the SARS-CoV-2 novel coronavirus that caused the disease and death of millions? Well we know the word *novel* means *new*, but is also means a *long story…. a very long, made-up story*.

In the case of ***The COVID-19 Story,*** **FACT is STRANGER than FICTION.** What we were told about the novel coronavirus, SARS-CoV-2, COVID-19, variants, mRNA technology, the deadly spike proteins, and PCR-tests is more akin to a plot from a movie in the Marvel series.

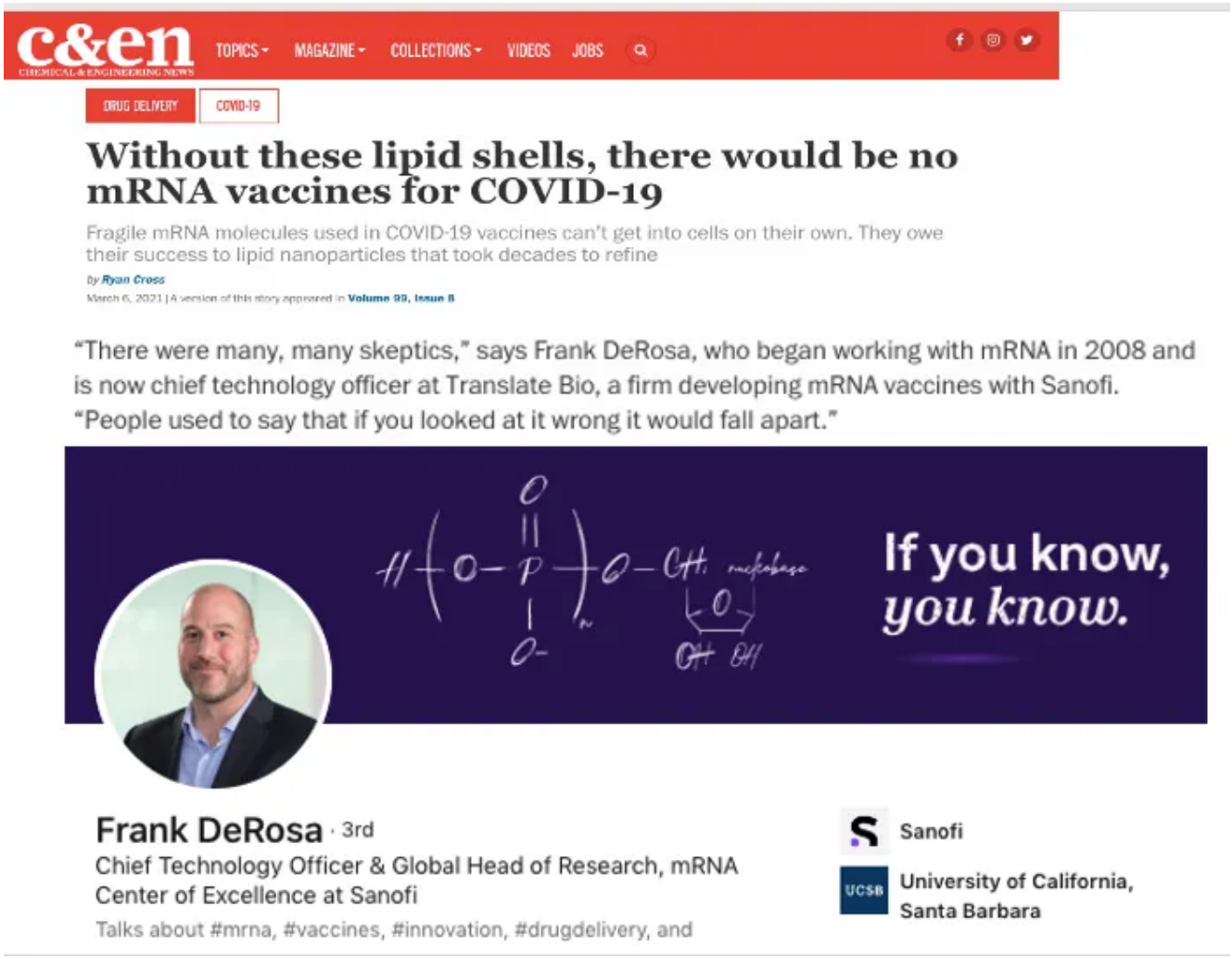
If the 2020 COVID-19 headlines were honest, accurate and actually non-fiction, they should have read, "**The *Masters of Evil* Terrorize Global Citizens by Spraying Down Cities and Towns with Aerosolized Biosynthetic Ai Nanoweapons called Spike Proteins**."

Wait? But the *Masters of Evil* never told us that the highly deadly spike protein was actually an Ai magnetic-hydrogel enclosed in lipid nanoparticle (LNP) technology used to infect, injure, experiment on and execute humans.

The *Masters of Evil* convinced us that the *SARS-CoV-2 mRNA virus produces the highly deadly spike protein after the mRNA invades cells inside the body*. Except there is one major issue with this claim, there is body of scientific evidence confirming that ***mRNA is unstable, fragile, and impotent (weak and useless inside the human body)***.

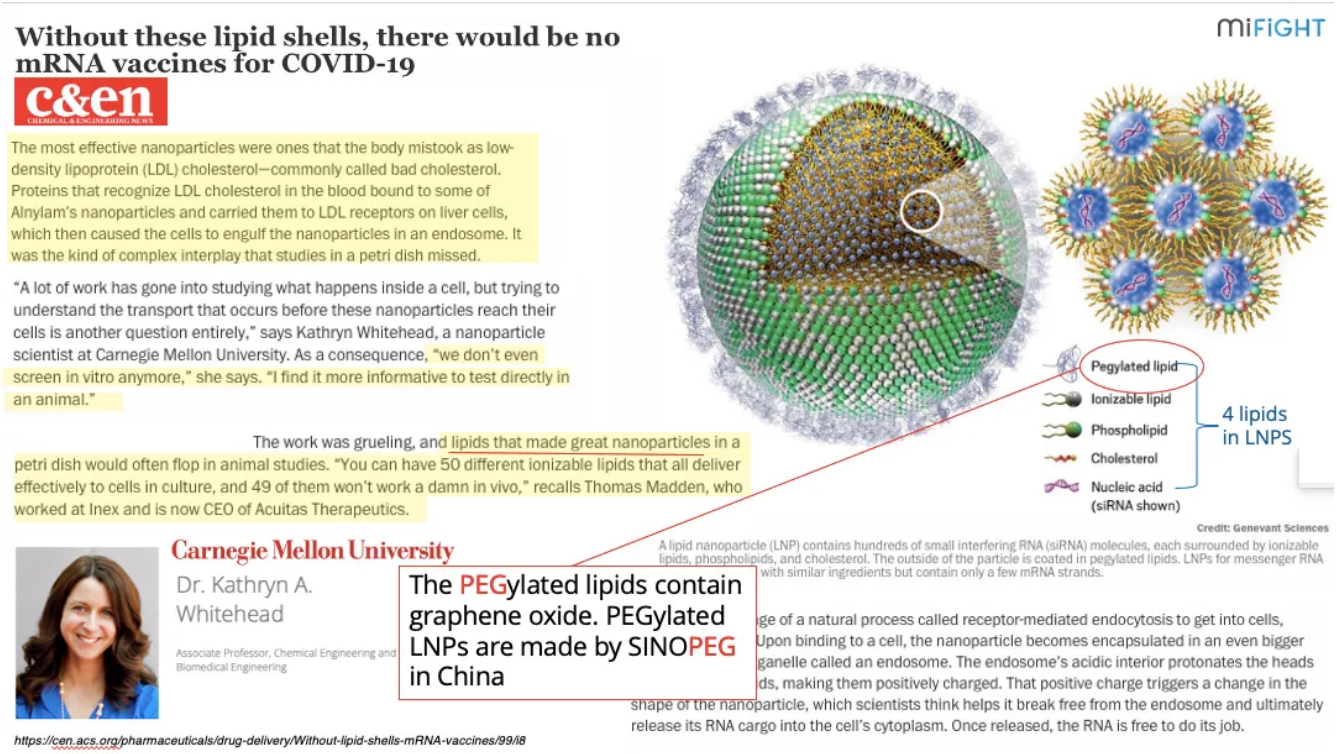
According to a March 6, 2021, *Chemical and Engineering News* article, mRNA lab experiments historically failed because the mRNA invention is so fragile and unstable on its own. This *C&EN* article states, "*Enzymes in the environment and in our bodies are quick to chop mRNA into pieces, making lab experiments difficult and the delivery of mRNA to our cells daunting.*" Sanofi's

Chief Technology Officer & Global Head of mRNA Research, Frank DeRosa, had this to say about mRNA, *"People used to say that if you looked at it (mRNA) wrong it would fall apart."*

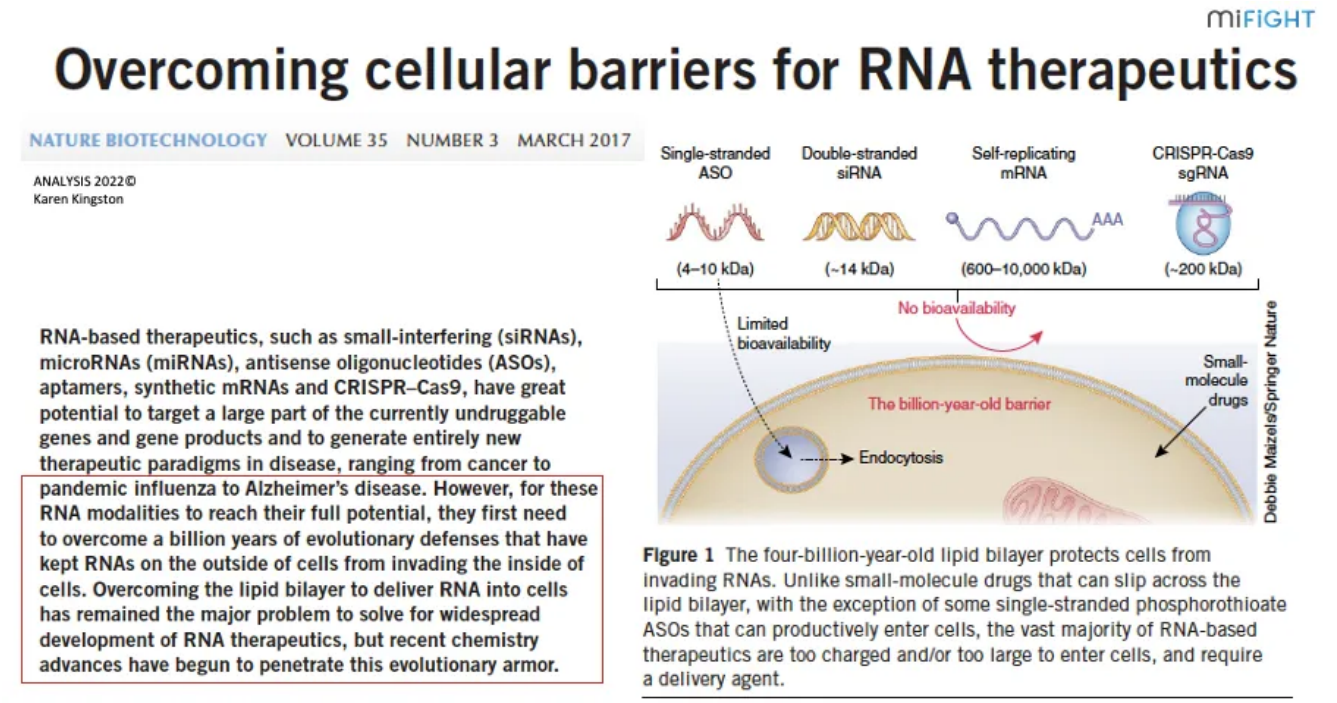


Dr. Kathryn Whitehead, head of Chemical and Bioengineering at Carnegie Mellon, expressed her frustration at attempts to conduct mRNA experiments out side of the body and stated;

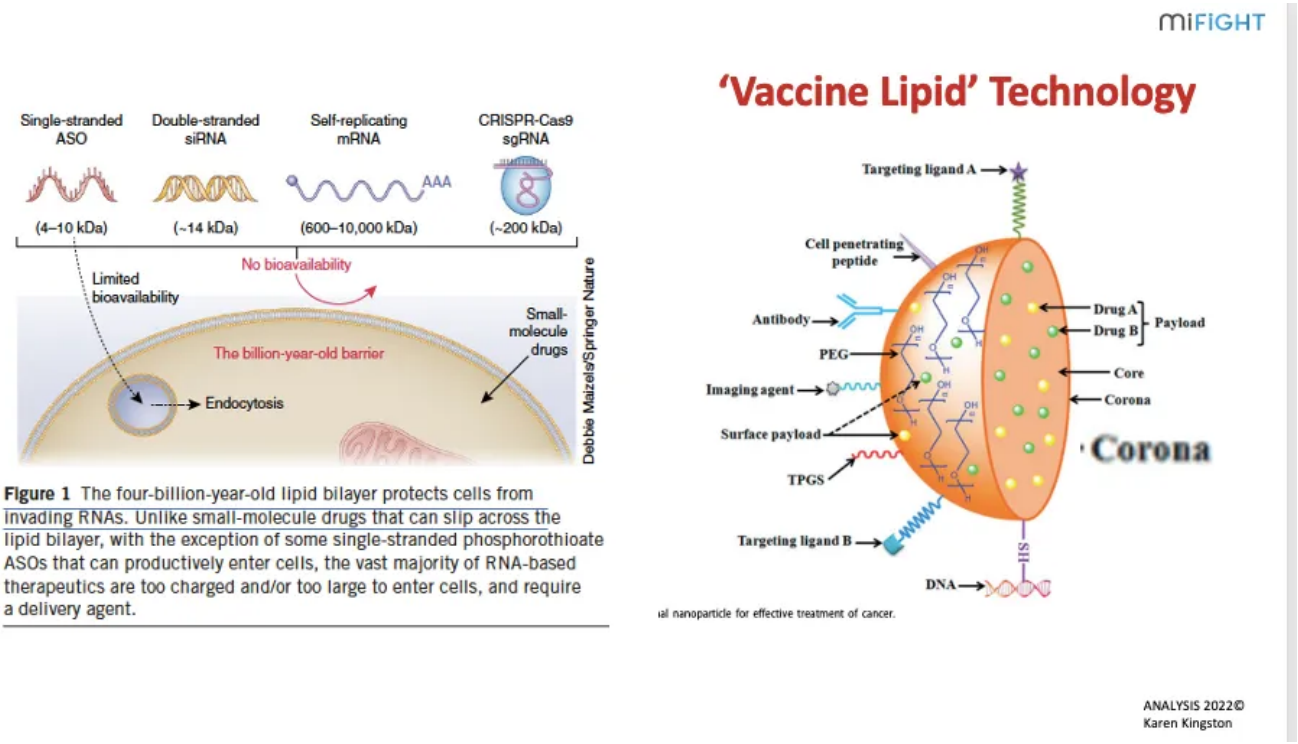
Dr. Kathryn Whitehead, Carnegie Mellon, "We don't even screen (mRNA) in vitro anymore. I find it more informative to test (mRNA) directly inside the animal."



Per *Nature Biotechnology*, 2017 peer-reviewed publication, "*Overcoming Cellular Barriers for RNA Therapeutics in RNA*," RNA gene-therapies, including ***mRNA, are unable to infect cells, any cells, due to billions of years of an evolutionary defense mechanism, the lipid bilayer.***

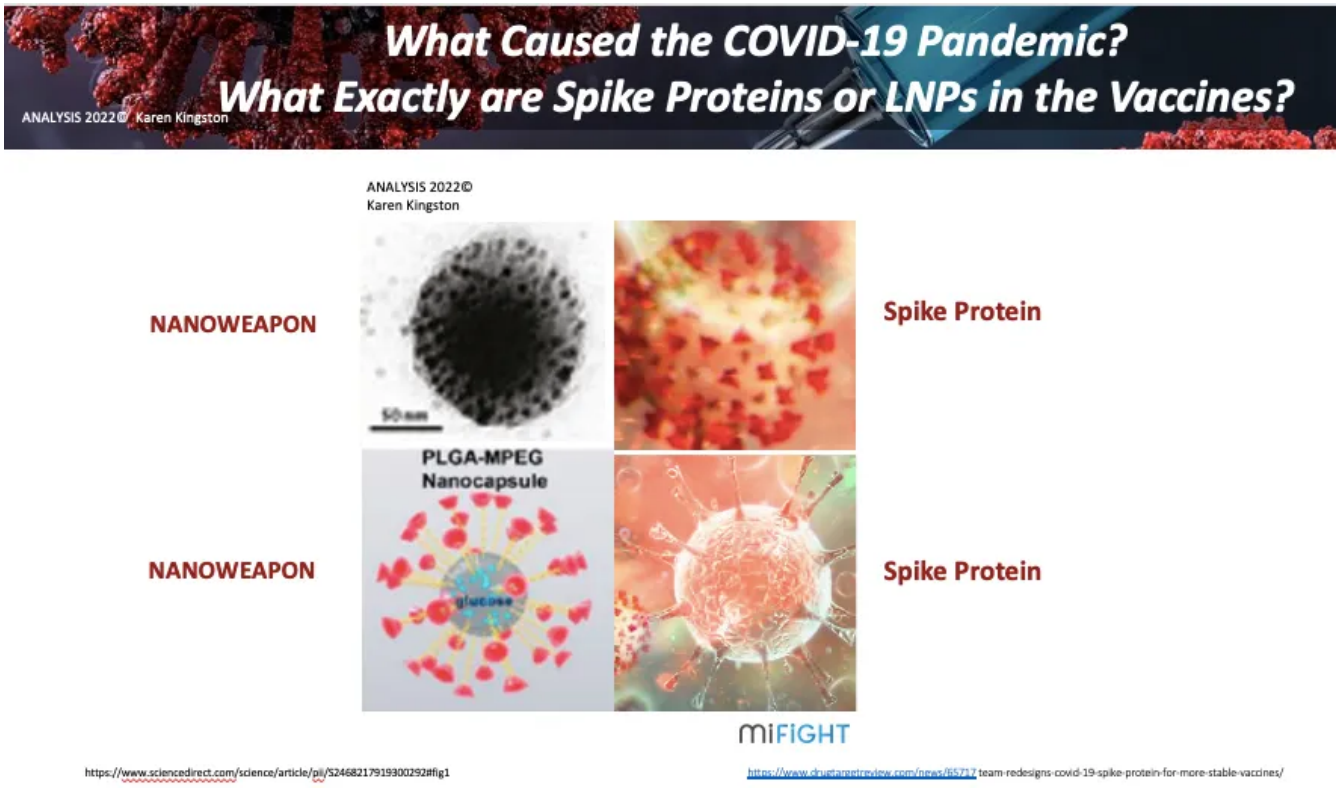


mRNA would be blasted into eternal nonexistence (where it belongs) if it wasn't for the advanced lipid nanoparticle (LNP) technology. LNPs enable the mRNA gene-editing technology to invade the lipid bilayer of cells and then infect, replicate, and produce *biosynthetic proteins* and other *mRNA-sequence-instructed biosynthetic structures*.

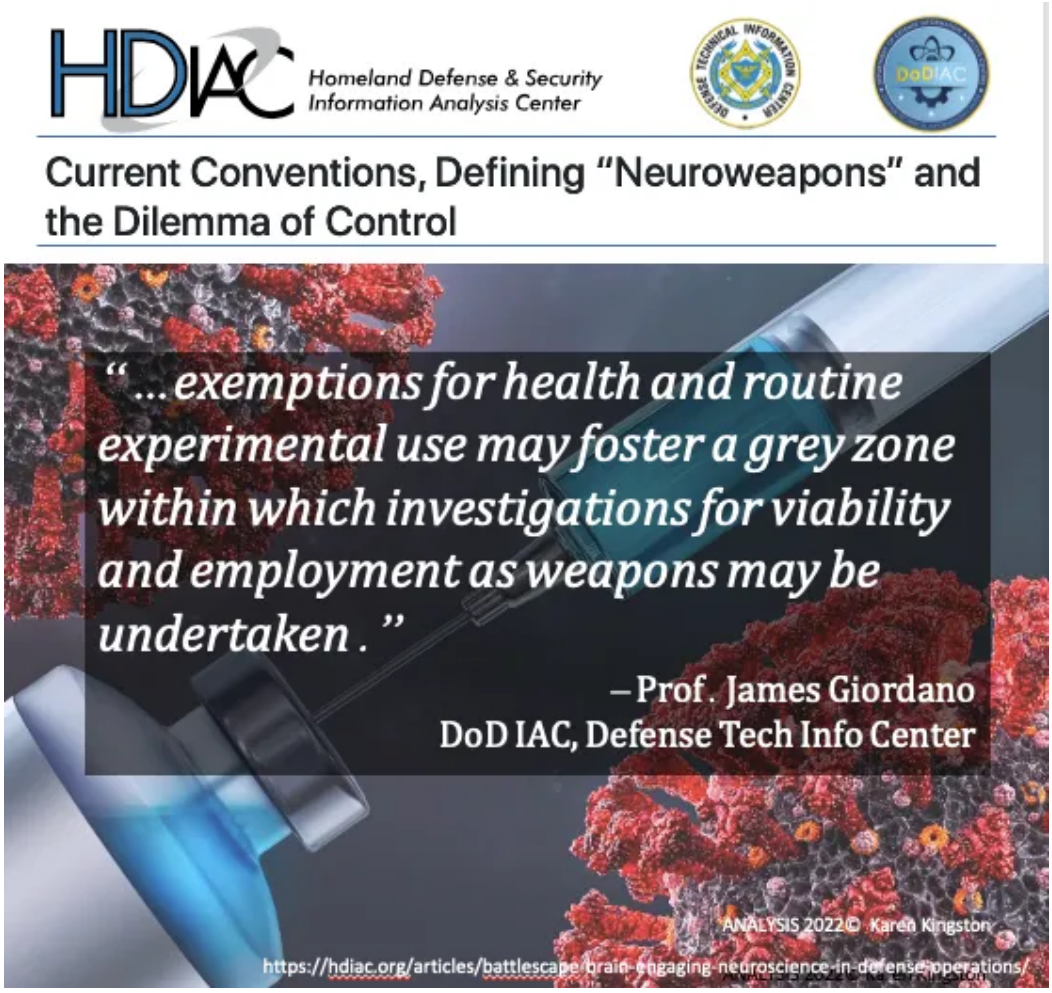


To put the magnitude of the pure evil genius of the LNP technology, mRNA was not able to infect any cells, including human cells, due to a four-billion year evolved lipid bilayer that protects cells, all cells, until this technology was invented.

The lipid nanoparticle (LNP) technology is pure evil. LNPs are also considered bio-tagging neuro-technologies and nanoweapons, that happen to look exactly like the COVID-19 spike proteins.

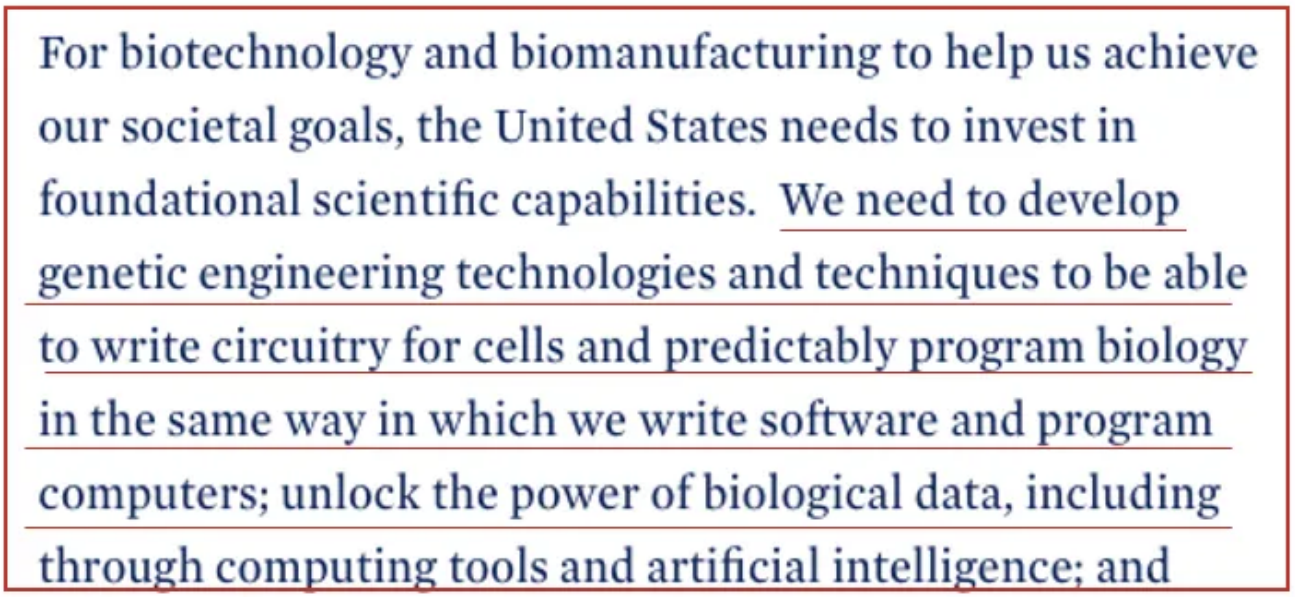


Per a 2018, West Point lecture given by Professor James Giordano at the Modern War Institute, these lipid nanoparticles (LNPs) have been used as nanoweapons to emotionally and mentally hijack influential leaders and can be used to induce a pandemic of 'worried well' or 'pandemic of strokes.'



Another deception we were led to believe is that *the mRNA sequence is an actual synthetic virus* preloaded in the 'vaccines'. The mRNA sequence is not a synthetic virus. It's a computer program sequence. The lipid nanotechnology does the gene editing and produces the spike proteins directly inside cells inside the body.

President Joe Biden signed an Executive Order on September 12, 2022, calling for an 'all of government approach' to ensure the bodies of *Americans will be installed with genetic engineering technologies* (aka, *lipid nanoparticles*) to program their cells the same way we write software and program computers because the gene-editing technology has already been installed in most Americans.


For biotechnology and biomanufacturing to help us achieve our societal goals, the United States needs to invest in foundational scientific capabilities. We need to develop genetic engineering technologies and techniques to be able to write circuitry for cells and predictably program biology in the same way in which we write software and program computers; unlock the power of biological data, including through computing tools and artificial intelligence; and

> We were informed that the mRNA SARS-CoV-2 sequence in the 'vaccines' produces spike proteins that are biological, in other words, organic in nature. This is another deception. What is produced by the LNPs and programmed sequences are genetically coded biosynthetic proteins and structures that are part biology and part technology, both organic and inorganic.

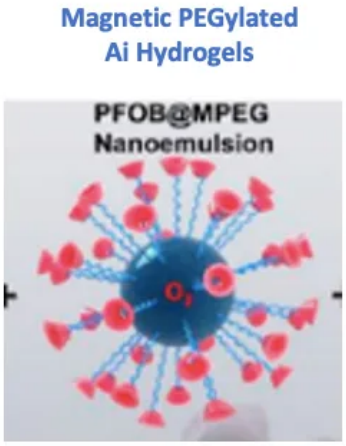
The pegylated LNPs contain magnetic hydrogels, a smart technology that is perceptive, responsive, and intelligent. Magnetic hydrogels produce can produce viruses, bacteria, toxins, and biosynthetic organisms and organelles inside the human body. Smart magnetic hydrogels can also send and receive signals through optics and electromagnetic frequencies.

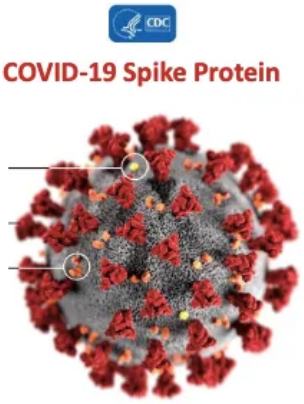
Multi-functional magnetic hydrogel: Design strategies and applications

2.5 | Intelligent response

Smart hydrogel is a kind of material that can perceive small physical/chemical stimuli (such as temperature, light, magnetism, pH) and make significant response behaviors.[65] Because of this intelligence, hydrogel has a fascinating application prospect in tissue engineering, drug-controlled release and soft actuators. Especially, as an external stimulus of stimulus-responsive materials, magnetic field has the advantages of instant action, contactless control and easy integration into electronic devices. Therefore, the research and development of smart MHs has been very active in recent years.[66]

Over the past few decades, tissue engineering has been successfully applied to the repair of various tissues (retinas, ligaments, fats, blood vessels, etc.). With the potential of hydrogel to construct microenvironment, the scaffolds based on multi-functional MHs have attracted much attention due to their intelligence. On the one hand, under the guidance of magnetic field, MHs can move directionally or be induced into specific tissue-like microstructure,[67] providing a suitable growth environment for tissue reconstruction. Schmidt proposed a novel magnetic templating technology which can induce highly aligned 3D tubular microstructures in naturally derived hydrogel scaffolds.[68]





https://dnascience.plos.org/2020/02/20/covid-19-vaccine-will-close-in-on-the-spikes

https://onlinelibrary.wiley.com/doi/epdf/10.1002/nano.202100139/

More and more people are becoming magnetic because of this Ai nanoweapon technology. Magnetic humans are not a consequence of an mRNA gain of function virus or vaccine. It's a consequence of magnetic-hydrogel Ai nanoweapons infiltrating the human body.

The SARS-CoV-2 mRNA virus sequence that was released on the globe and is found in the 'vaccines' would be completely useless, unable to infect cells and have no ability to produce antibodies without the lipid nanoparticles (LNPs) and magnetic Ai hydrogels. This also means that the COVID-19 pandemic was impossible without the LNPs. ***COVID-19 was not caused by a mRNA gain-of-function SARS-CoV-2 virus. COVID-19 was caused by an Ai nanoweapon.***

Another leading mRNA expert is Ralph Baric from the University of North Carolina. He is the godfather of coronavirus mRNA gain-of-function research and chimeric spike proteins. He has received over $100 million for coronavirus research from Fauci's NIAID/NIH and is a named inventor on dozens of patents for coronaviruses, chimeric spike proteins, and other gain-of-function bioweapon-based technologies.

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Compositions of Chimeric Spike Proteins

Methods and compositions for chimeric coronavirus spike proteins
Patent number: 9884895
Abstract: The present invention provides compositions and methods comprising a chimeric coronavirus spike protein.
Type: Grant
Filed: March 20, 2015
Date of Patent: February 6, 2018
Assignee: The University of North Carolina at Chapel Hill
Inventors: Ralph Baric, Sudhakar Agnihothram, Boyd Yount

STATEMENT OF FEDERAL SUPPORT

This invention was made with government support under Grant No. U54AI057157 awarded by the National Institutes of Health. The government has certain rights in the invention.

FIELD OF THE INVENTION

The present invention relates to methods and compositions comprising a chimeric coronavirus spike protein for treating and/or preventing a disease or disorder caused by a coronavirus infection.

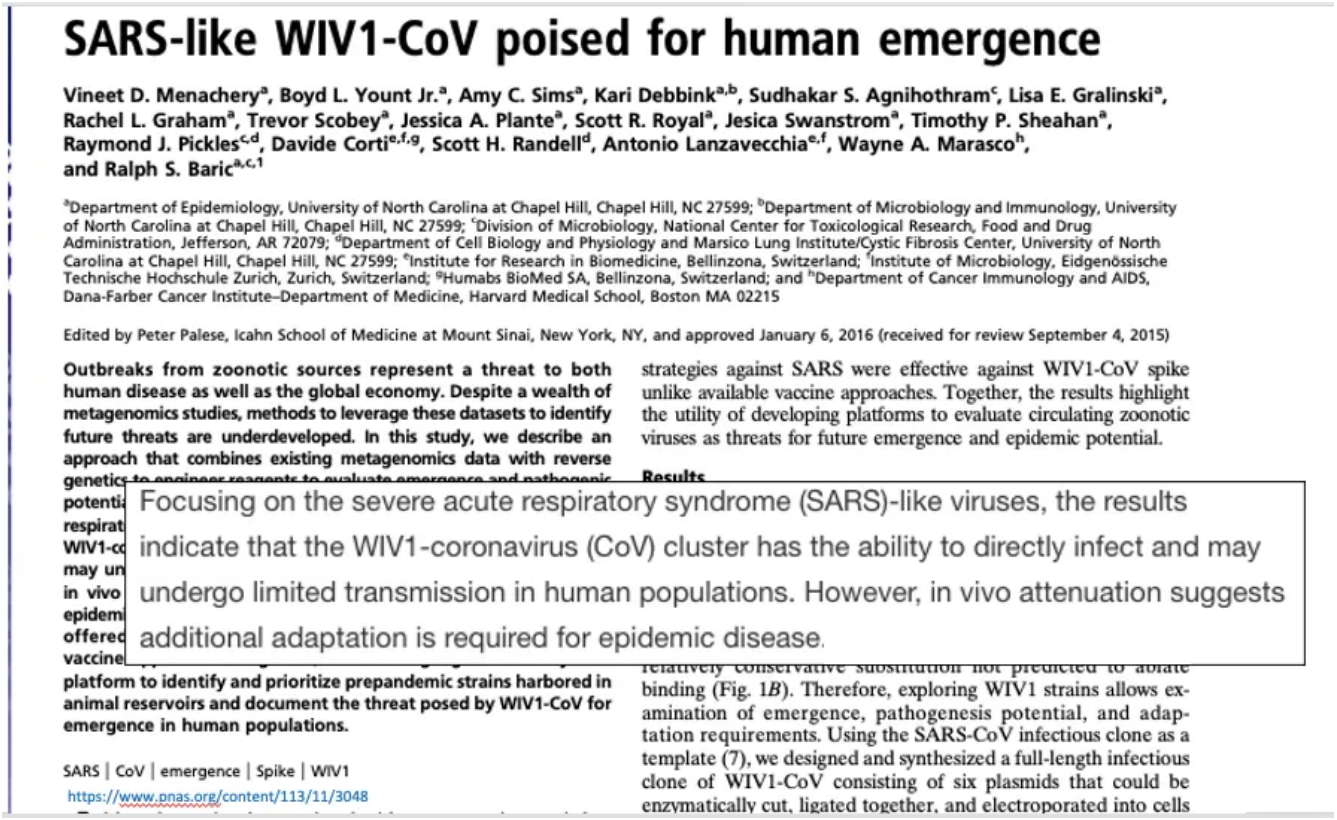
SUMMARY OF THE INVENTION

In one aspect, the present invention provides a chimeric coronavirus spike protein comprising, in orientation from amino to carboxy terminus: a) a first region comprising a portion of a coronavirus spike protein ectodomain that precedes the receptor binding domain (RBD) as located in a nonchimeric coronavirus spike protein, of a first coronavirus; b) a second region comprising a coronavirus spike protein receptor binding domain (RBD) of a second coronavirus that is different from said first coronavirus; c) a third region comprising a portion of a coronavirus spike protein S1 domain as located in a nonchimeric coronavirus spike protein immediately downstream of the RBD, contiguous with a portion comprising a coronavirus spike protein S2 domain as located immediately upstream of a fusion protein domain in a nonchimeric coronavirus spike protein, wherein said third region is of said first coronavirus; and d) a fourth region comprising a portion of a coronavirus spike protein from the start of the fusion protein domain through the carboxy terminal end as located in a nonchimeric coronavirus spike protein of a third coronavirus that is different from said first coronavirus and said second coronavirus.

1. https://patents.justia.com/patent/9884895

In his paper submitted to *PNAS* on September 4, 2015 entitled, "*SAR-like WIV1-CoV Poised for Human Emergence*," Ralph Baric recognizes that his coronavirus mRNA viruses *(**including SARS-CoV-2) are not capable of significant human-to-human transmission, if any transmission at all.***

- The paper states, "***WIV1-coronavirus** (CoV) cluster…may undergo limited transmission in human populations…**in vivo attenuation** (attenuation means that **SARS-CoV-2 becomes extremely weak once inside the human body) suggests additional adaptation is required for epidemic disease.***"



SARS-like WIV1-CoV poised for human emergence

Vineet D. Menachery[a], Boyd L. Yount Jr.[a], Amy C. Sims[a], Kari Debbink[a,b], Sudhakar S. Agnihothram[c], Lisa E. Gralinski[a], Rachel L. Graham[a], Trevor Scobey[a], Jessica A. Plante[a], Scott R. Royal[a], Jesica Swanstrom[a], Timothy P. Sheahan[a], Raymond J. Pickles[c,d], Davide Corti[e,f,g], Scott H. Randell[d], Antonio Lanzavecchia[e,f], Wayne A. Marasco[h], and Ralph S. Baric[a,c,1]

[a]Department of Epidemiology, University of North Carolina at Chapel Hill, Chapel Hill, NC 27599; [b]Department of Microbiology and Immunology, University of North Carolina at Chapel Hill, Chapel Hill, NC 27599; [c]Division of Microbiology, National Center for Toxicological Research, Food and Drug Administration, Jefferson, AR 72079; [d]Department of Cell Biology and Physiology and Marsico Lung Institute/Cystic Fibrosis Center, University of North Carolina at Chapel Hill, Chapel Hill, NC 27599; [e]Institute for Research in Biomedicine, Bellinzona, Switzerland; [f]Institute of Microbiology, Eidgenössische Technische Hochschule Zurich, Zurich, Switzerland; [g]Humabs BioMed SA, Bellinzona, Switzerland; and [h]Department of Cancer Immunology and AIDS, Dana-Farber Cancer Institute–Department of Medicine, Harvard Medical School, Boston MA 02215

Edited by Peter Palese, Icahn School of Medicine at Mount Sinai, New York, NY, and approved January 6, 2016 (received for review September 4, 2015)

Outbreaks from zoonotic sources represent a threat to both human disease as well as the global economy. Despite a wealth of metagenomics studies, methods to leverage these datasets to identify future threats are underdeveloped. In this study, we describe an approach that combines existing metagenomics data with reverse genetics to engineer reagents to evaluate emergence and pathogenic potenti respirat WIV1-c may un in vivo epidem offered vaccine platform to identify and prioritize prepandemic strains harbored in animal reservoirs and document the threat posed by WIV1-CoV for emergence in human populations.

SARS | CoV | emergence | Spike | WIV1
https://www.pnas.org/content/113/11/3048

strategies against SARS were effective against WIV1-CoV spike unlike available vaccine approaches. Together, the results highlight the utility of developing platforms to evaluate circulating zoonotic viruses as threats for future emergence and epidemic potential.

Results

Focusing on the severe acute respiratory syndrome (SARS)-like viruses, the results indicate that the WIV1-coronavirus (CoV) cluster has the ability to directly infect and may undergo limited transmission in human populations. However, in vivo attenuation suggests additional adaptation is required for epidemic disease.

relatively conservative substitution not predicted to ablate binding (Fig. 1*B*). Therefore, exploring WIV1 strains allows examination of emergence, pathogenesis potential, and adaptation requirements. Using the SARS-CoV infectious clone as a template (7), we designed and synthesized a full-length infectious clone of WIV1-CoV consisting of six plasmids that could be enzymatically cut, ligated together, and electroporated into cells

In 2015, Baric recognized that mRNA coronaviruses were useless on their own. He clearly states that mRNA coronaviruses (including SARS-CoV-2) are not of 'epidemic potential.'

In order to capitalize on his decades of research, patents, and royalty agreements, Baric needed find a way to use his mRNA gain-of-function coronaviruses to create a bioweapon that would be of 'epidemic potential'.

Baric has a close-knit team for his gain-of-function mRNA coronavirus research. He's been decades-long partners with Peter Daszak of EcoHealth Alliance, and Dr. Zhenghli Shi of the Wuhan Institute of Virology (WIV). This evil trio has created countless gain-of-function weaponized mRNA viruses, and not just coronaviruses; zika, ebola, influenza, HIV, and many others.

Wuhan coronavirus hunter Shi Zhengli speaks out

China's "Bat Woman" denies responsibility for the pandemic, demands apology from Trump.

JON COHEN

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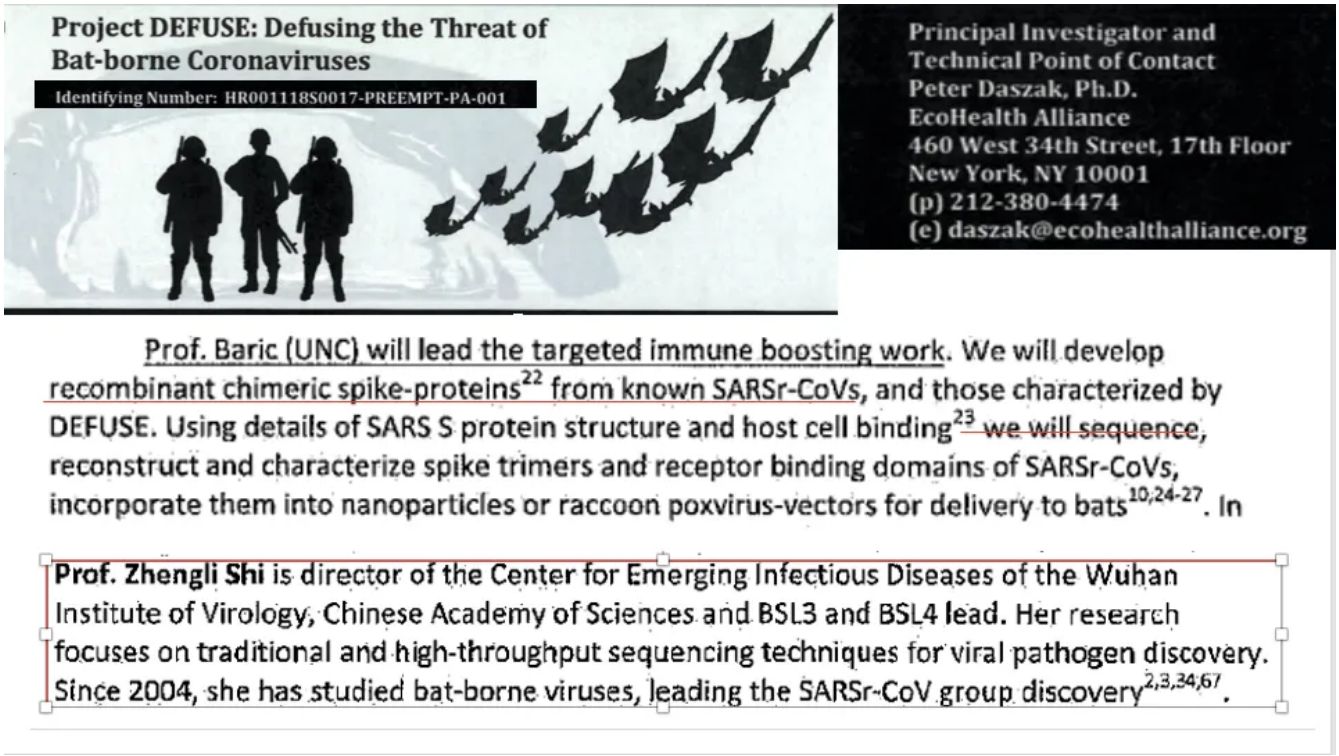
1,998 7

Science's COVID-19 reporting is supported by the Pulitzer Center and the Heising-Simons Foundation

In January of 2018, he and his buddies reached out to our military's Defense Advanced Research Projects Agency (DARPA) for funding via a proposal. In EcoHealth Alliance's proposal submitted to DARPA, Baric and Shi were listed as EcoHealth Alliance's team members to develop the ***chimeric gain-of-function spike protein bioweapons*** from the mRNA coronavirus library the team had already created.

Per the DARPA proposal, the next phase of their bioweapons research was focused on how to take Baric's virtually useless mRNA gain-of-function coronaviruses (by his own admission) to create something from his mRNA viruses that was infectious, deadly and of 'pandemic potential'.



The 2018 EcoHealth Alliance DARPA proposal clearly states,

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*"**WE WILL DEVELOP** recombinant **CHIMERIC SPIKE-PROTEINS** from known SARSr-CoV ...*
***AND INCORPORATE THEM INTO NANOPARTICLES...**"*

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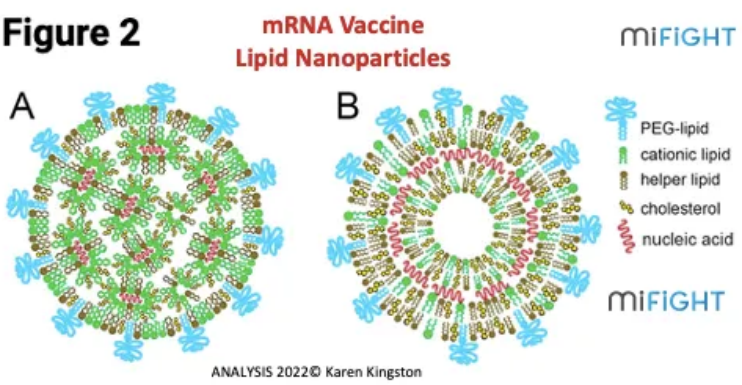
Lipid Nanoparticles—From Liposomes to mRNA Vaccine Delivery, a Landscape of Research Diversity and Advancement

Rumiana Tenchov, Robert Bird, Allison E. Curtze, and Qiongqiong Zhou*

Figure 2



SPIKE Protein Image

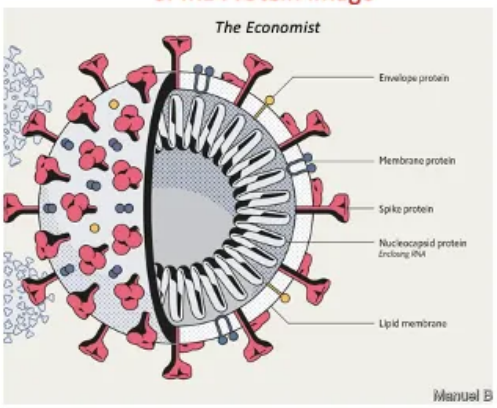


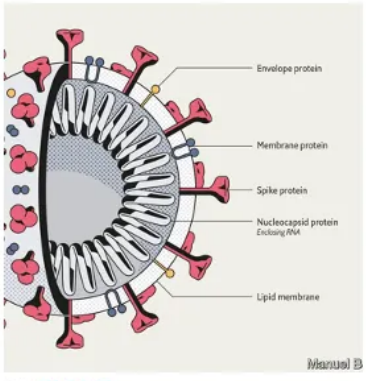
Figure 2. Suggested structures of lipid nanoparticle nucleic acid carriers: nucleic acids organized in inverse lipid micelles inside the nanoparticle (A); nucleic acids intercalated between the lipid bilayers (B). (26−29)

https://pubs.acs.org/doi/10.1021/acsnano.1c04996

Cite this: *ACS Nano* 2021, 15, 11, 16982−17015
Publication Date: June 28, 2021
https://doi.org/10.1021/acsnano.1c04996

The lipid nanoparticles (LNPs) are the Ai bioweapons as are the spike proteins.

SARS-CoV-2 Spike Protein



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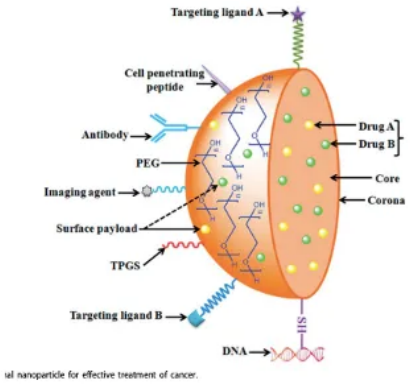
'Vaccine Lipid' Technology



Military NANOWEAPON

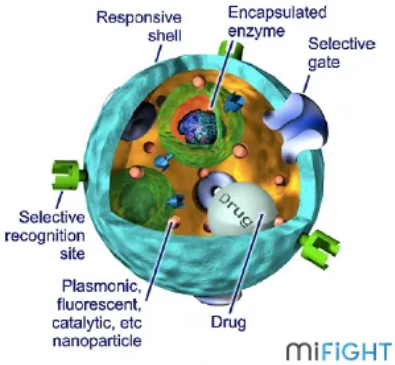


Fig. 1. Futuristic picture of the intelligent multifunctional particle.

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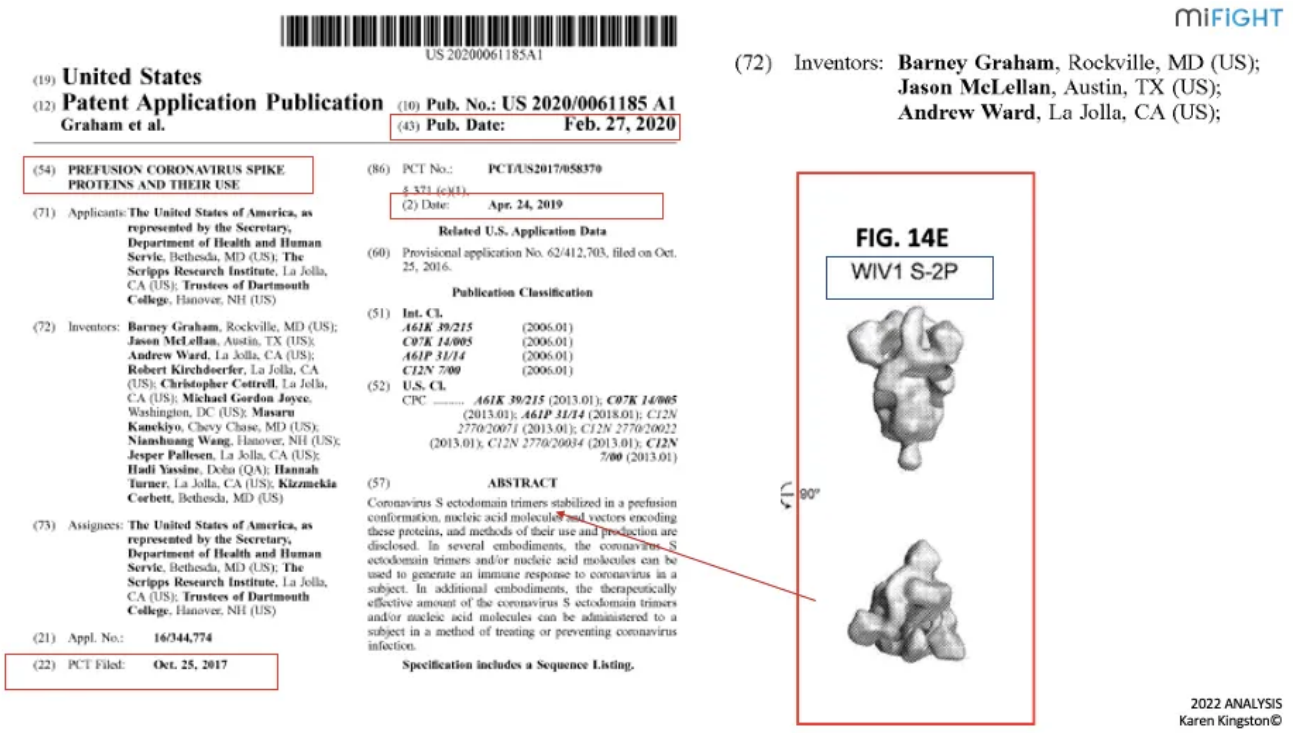
The DARPA proposal is further evidence that **the SARS-CoV-2 mRNA virus was never infectious or deadly.** It's the WIV1 2-SP spike protein (literally named after the Wuhan Institute of Virology) that's highly-inflammatory and deadly, not the mRNA SARS-CoV-2 virus. The WIV1 2-SP 'spike protein' is a product of the parasitic Ai bioweapon, as are all other COVID-19 spike proteins.

IMPORTANT: **The highly-deadly WIV-1 S-2P spike protein is also a separate invention from SARS-CoV-2. The WIV-S-2P spike and other S-2P spike proteins are made in labs and have their own patent filed by Barney Graham from NIAID/NIH and Jason McLellan from University of Texas.**



US 20200061185A1

(19) **United States**

(12) **Patent Application Publication**

Graham et al.

(10) **Pub. No.: US 2020/0061185 A1**

(43) **Pub. Date: Feb. 27, 2020**

(54) **PREFUSION CORONAVIRUS SPIKE PROTEINS AND THEIR USE**

(71) Applicants: **The United States of America, as represented by the Secretary, Department of Health and Human Servic**, Bethesda, MD (US); **The Scripps Research Institute**, La Jolla, CA (US); **Trustees of Dartmouth College**, Hanover, NH (US)

(72) Inventors: **Barney Graham**, Rockville, MD (US); **Jason McLellan**, Austin, TX (US); **Andrew Ward**, La Jolla, CA (US); **Robert Kirchdoerfer**, La Jolla, CA (US); **Christopher Cottrell**, La Jolla, CA (US); **Michael Gordon Joyce**, Washington, DC (US); **Masaru Kanekiyo**, Chevy Chase, MD (US); **Nianshuang Wang**, Hanover, NH (US); **Jesper Pallesen**, La Jolla, CA (US); **Hadi Yassine**, Doha (QA); **Hannah Turner**, La Jolla, CA (US); **Kizzmekia Corbett**, Bethesda, MD (US)

(73) Assignees: **The United States of America, as represented by the Secretary, Department of Health and Human Servic**, Bethesda, MD (US); **The Scripps Research Institute**, La Jolla, CA (US); **Trustees of Dartmouth College**, Hanover, NH (US)

(21) Appl. No.: **16/344,774**

(22) PCT Filed: **Oct. 25, 2017**

(86) PCT No.: **PCT/US2017/058370**

§ 371 (c)(1),

(2) Date: **Apr. 24, 2019**

Related U.S. Application Data

(60) Provisional application No. 62/412,703, filed on Oct. 25, 2016.

Publication Classification

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C07K 14/005 (2006.01)
A61P 31/14 (2006.01)
C12N 7/00 (2006.01)

(52) **U.S. Cl.**

CPC ***A61K 39/215*** (2013.01); ***C07K 14/005*** (2013.01); ***A61P 31/14*** (2018.01); *C12N 2770/20071* (2013.01); *C12N 2770/20022* (2013.01); *C12N 2770/20034* (2013.01); ***C12N 7/00*** (2013.01)

(57) **ABSTRACT**

Coronavirus S ectodomain trimers stabilized in a prefusion conformation, nucleic acid molecules and vectors encoding these proteins, and methods of their use and production are disclosed. In several embodiments, the coronavirus S ectodomain trimers and/or nucleic acid molecules can be used to generate an immune response to coronavirus in a subject. In additional embodiments, the therapeutically effective amount of the coronavirus S ectodomain trimers and/or nucleic acid molecules can be administered to a subject in a method of treating or preventing coronavirus infection.

Specification includes a Sequence Listing.

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(72) Inventors: **Barney Graham**, Rockville, MD (US); **Jason McLellan**, Austin, TX (US); **Andrew Ward**, La Jolla, CA (US);

FIG. 14E

WIV1 S-2P

90°

2022 ANALYSIS
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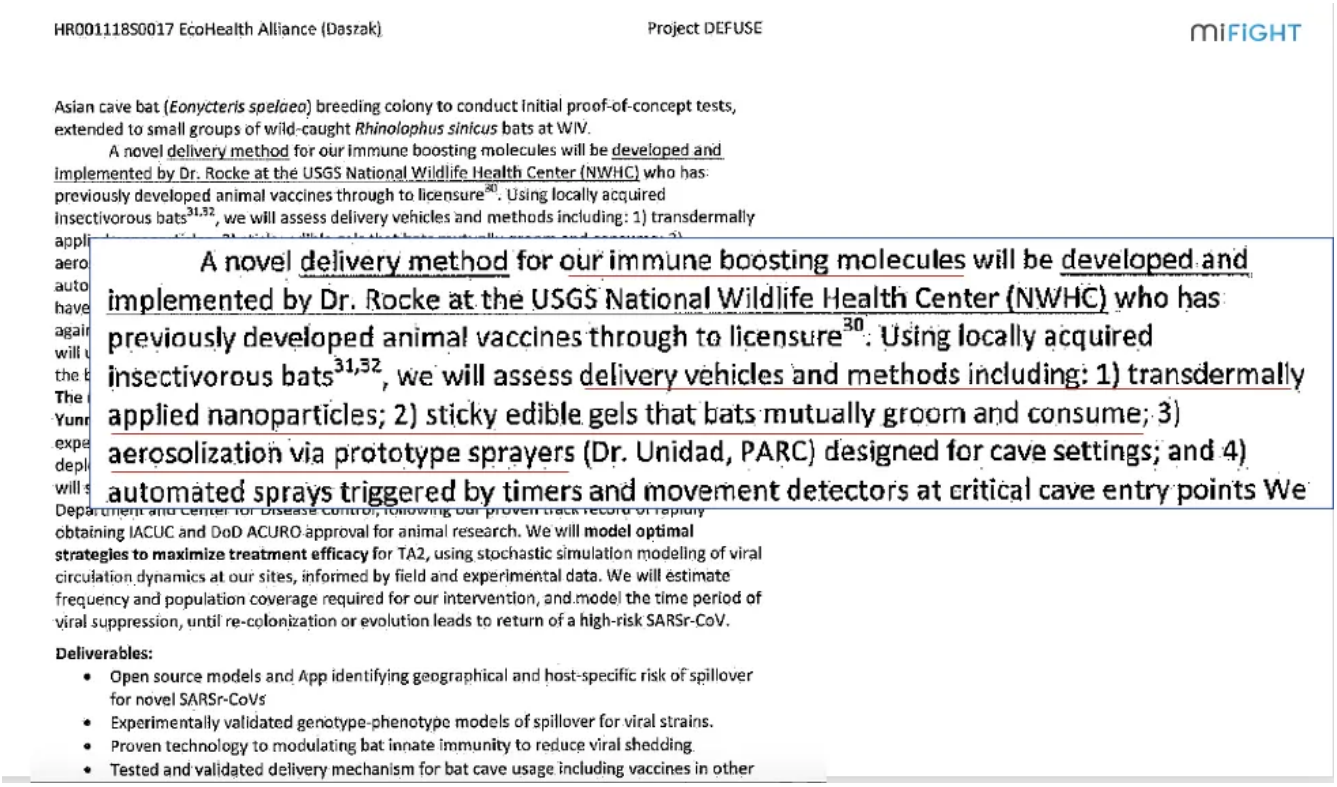
In the DARPA proposal, Daszak also states they are going to infect the bats by delivering the ***lipid nanoparticle (LNP) technology encapsulated spike proteins*** in the form of a transdermal (skin) patch, a gel for the bats to eat, a gel for their skin, or inhaled aerosol sprays.



HR001118S0017 EcoHealth Alliance (Daszak) Project DEFUSE MIFIGHT

Asian cave bat (*Eonycteris spelaea*) breeding colony to conduct initial proof-of-concept tests, extended to small groups of wild-caught *Rhinolophus sinicus* bats at WIV.

A novel delivery method for our immune boosting molecules will be developed and implemented by Dr. Rocke at the USGS National Wildlife Health Center (NWHC) who has previously developed animal vaccines through to licensure[30]. Using locally acquired insectivorous bats[31,32], we will assess delivery vehicles and methods including: 1) transdermally applied nanoparticles; 2) sticky edible gels that bats mutually groom and consume; 3) aerosolization via prototype sprayers (Dr. Unidad, PARC) designed for cave settings; and 4) automated sprays triggered by timers and movement detectors at critical cave entry points We

obtaining IACUC and DoD ACURO approval for animal research. We will **model optimal strategies to maximize treatment efficacy** for TA2, using stochastic simulation modeling of viral circulation dynamics at our sites, informed by field and experimental data. We will estimate frequency and population coverage required for our intervention, and model the time period of viral suppression, until re-colonization or evolution leads to return of a high-risk SARSr-CoV.

Deliverables:

- Open source models and App identifying geographical and host-specific risk of spillover for novel SARSr-CoVs
- Experimentally validated genotype-phenotype models of spillover for viral strains.
- Proven technology to modulating bat innate immunity to reduce viral shedding.
- Tested and validated delivery mechanism for bat cave usage including vaccines in other

The hypothesis that the original 2019 and 2020 SARS-CoV-2 outbreaks involved aerosolized bioweapons attacks is more than highly probable.

> The DARPA proposal states that they are going to infect the bats with, "*aerosolization sprayers designed for cave settings; and automated sprays triggered by timers and movement detectors at critical cave entry points.*"

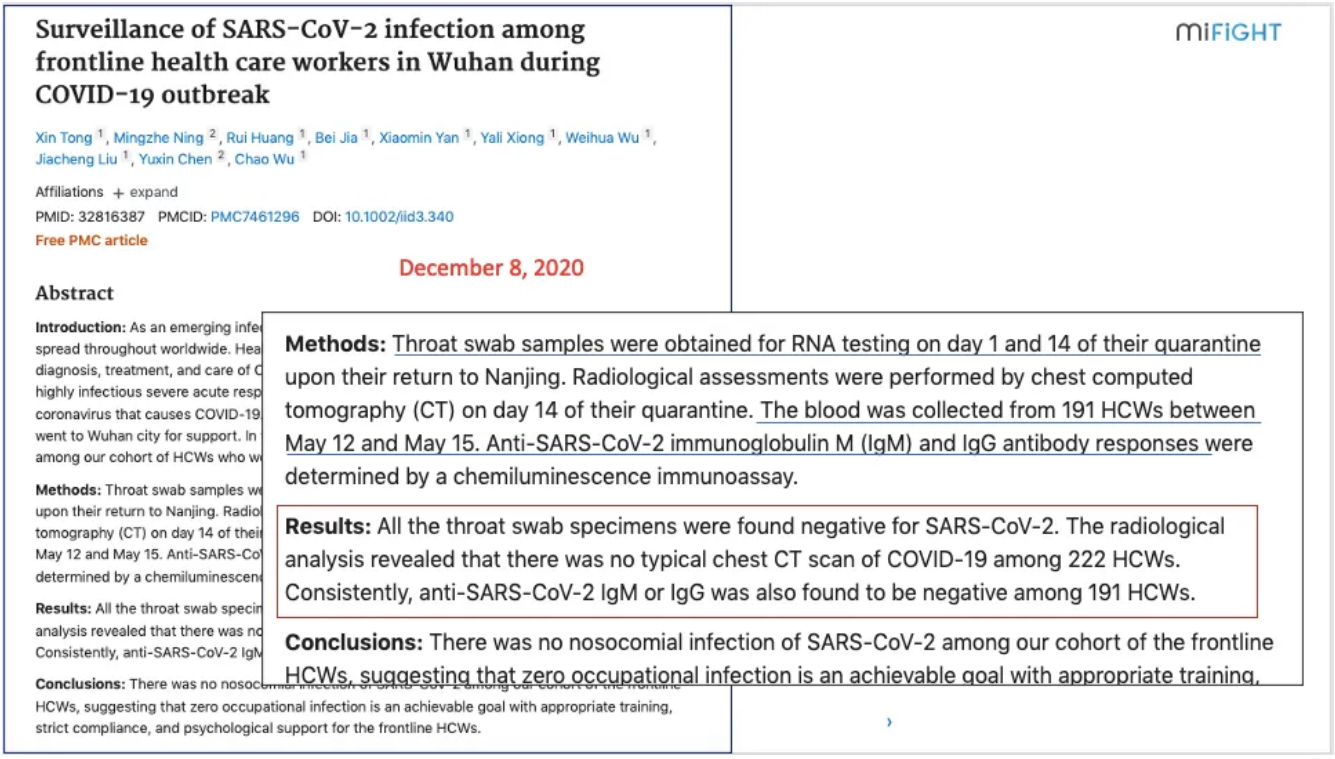
The SARS-CoV-2 mRNA virus was never deadly, nor was it contagious. I can't emphasize this enough, it wasn't SARS-CoV-2 that caused COVID-19. COVID-19 disease, disabilities and death are the product of Ai bioweapons.

The SARS-CoV-2 mRNA virus inventor's own research, Ralph Baric, stated that even if humans are infected with SARS-CoV-2 through a direct man-made delivery mechanism, human-to-human transmission was proven to be impossible. This is why the theory that SARS-CoV-2 was a lab leak is completely debunked.

> COVID-19 had to be an intentional Ai bioweapon attack to cause the type of injury and death first reported on around the globe in early 2020. The death and injury has exponentially grown due to the Ai bioweapon being in the COVID-19 shots and circulating in our environments.

The lack of human-to-human transmission is further validated by real world evidence data of hospital workers' infection rates at the hospital next to the Wuhan Market. The hospital worker infection rate at ground zero was 0.000%.

Published on December 8, 2020, in *Immunity, Inflammation, and Disease*, there were 191 Wuhan hospital workers who were tested for SARS-CoV-2 antibodies via throat swabs and blood work . **All tests came back negative**. Two-hundred and twenty-two (222) hospital workers also had chest X-rays. **All chest x-rays came back clean.**



Surveillance of SARS-CoV-2 infection among frontline health care workers in Wuhan during COVID-19 outbreak

Xin Tong [1], Mingzhe Ning [2], Rui Huang [1], Bei Jia [1], Xiaomin Yan [1], Yali Xiong [1], Weihua Wu [1], Jiacheng Liu [1], Yuxin Chen [2], Chao Wu [1]

Affiliations + expand

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Free PMC article

December 8, 2020

MiFIGHT

Abstract

Introduction: As an emerging infe
spread throughout worldwide. Hea
diagnosis, treatment, and care of C
highly infectious severe acute resp
coronavirus that causes COVID-19
went to Wuhan city for support. In
among our cohort of HCWs who w

Methods: Throat swab samples w
upon their return to Nanjing. Radio
tomography (CT) on day 14 of thei
May 12 and May 15. Anti-SARS-Co
determined by a chemiluminescen

Results: All the throat swab specim
analysis revealed that there was no
Consistently, anti-SARS-CoV-2 IgM

Conclusions: There was no nosoc
HCWs, suggesting that zero occupational infection is an achievable goal with appropriate training, strict compliance, and psychological support for the frontline HCWs.

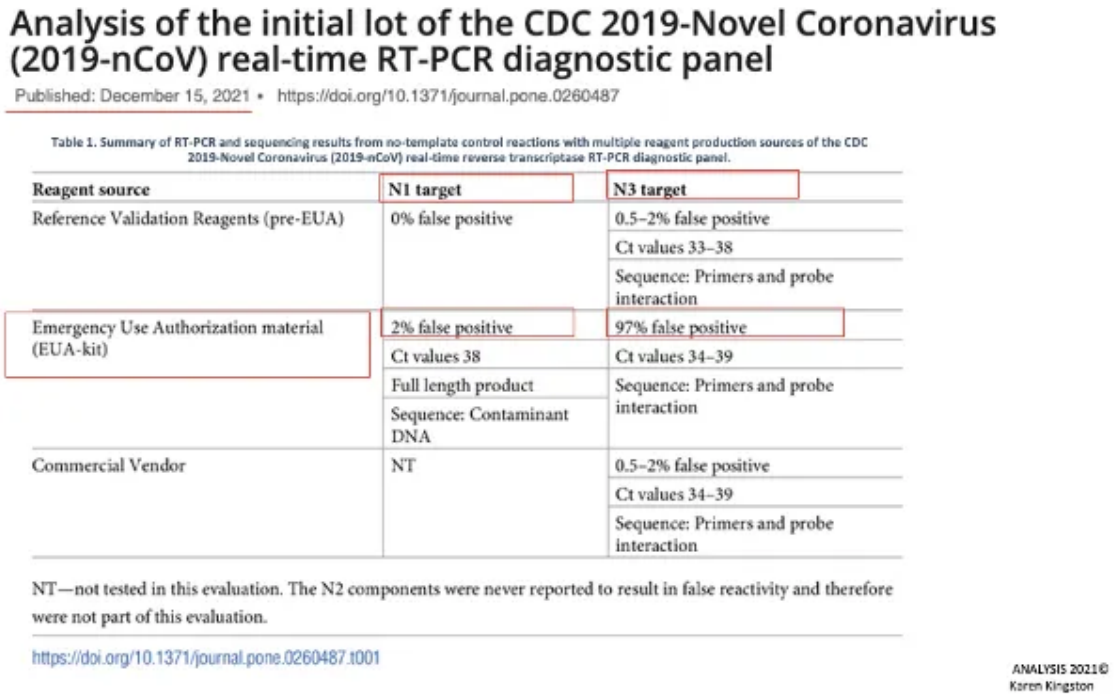
Methods: Throat swab samples were obtained for RNA testing on day 1 and 14 of their quarantine upon their return to Nanjing. Radiological assessments were performed by chest computed tomography (CT) on day 14 of their quarantine. The blood was collected from 191 HCWs between May 12 and May 15. Anti-SARS-CoV-2 immunoglobulin M (IgM) and IgG antibody responses were determined by a chemiluminescence immunoassay.

Results: All the throat swab specimens were found negative for SARS-CoV-2. The radiological analysis revealed that there was no typical chest CT scan of COVID-19 among 222 HCWs. Consistently, anti-SARS-CoV-2 IgM or IgG was also found to be negative among 191 HCWs.

Conclusions: There was no nosocomial infection of SARS-CoV-2 among our cohort of the frontline HCWs, suggesting that zero occupational infection is an achievable goal with appropriate training,

Hospital workers from Wuhan never got infected. Experts who proclaimed that COVID-19 was or is a highly-infectious virus were making false statements not based in real world evidence.

The CDC was intentionally misleading the US public into believing that the SARS-CoV-2 was capable of human-to-human transmission. Per the CDCs own data published on December 15, 2021, the emergency use authorized (EUA) PCR-tests used throughout 2020 and 2021 gave false positive tests of up to 97%.



Analysis of the initial lot of the CDC 2019-Novel Coronavirus (2019-nCoV) real-time RT-PCR diagnostic panel

Published: December 15, 2021 • https://doi.org/10.1371/journal.pone.0260487

Table 1. Summary of RT-PCR and sequencing results from no-template control reactions with multiple reagent production sources of the CDC 2019-Novel Coronavirus (2019-nCoV) real-time reverse transcriptase RT-PCR diagnostic panel.

Reagent source	N1 target	N3 target
Reference Validation Reagents (pre-EUA)	0% false positive	0.5–2% false positive
		Ct values 33–38
		Sequence: Primers and probe interaction
Emergency Use Authorization material (EUA-kit)	2% false positive	97% false positive
	Ct values 38	Ct values 34–39
	Full length product	Sequence: Primers and probe interaction
	Sequence: Contaminant DNA	
Commercial Vendor	NT	0.5–2% false positive
		Ct values 34–39
		Sequence: Primers and probe interaction

NT—not tested in this evaluation. The N2 components were never reported to result in false reactivity and therefore were not part of this evaluation.

https://doi.org/10.1371/journal.pone.0260487.t001

ANALYSIS 2021©
Karen Kingston

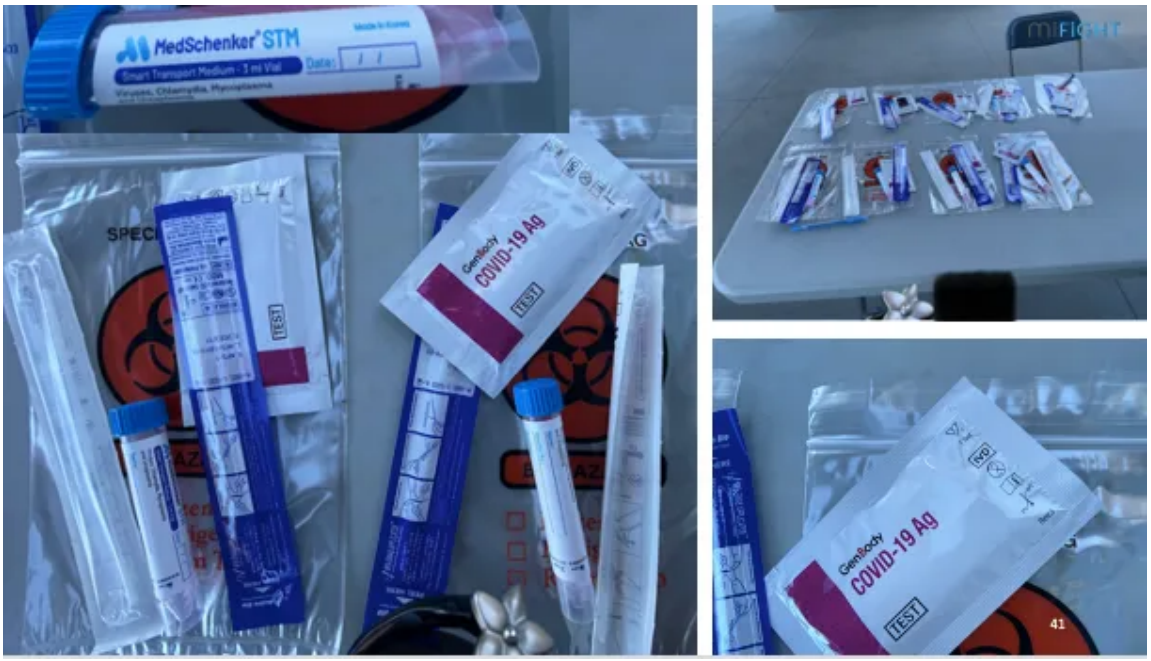
The EUA COVID-19 PCR-tests are as effective at measuring SARS-CoV-2 infections as Monopoly money is at buying a new car or home.

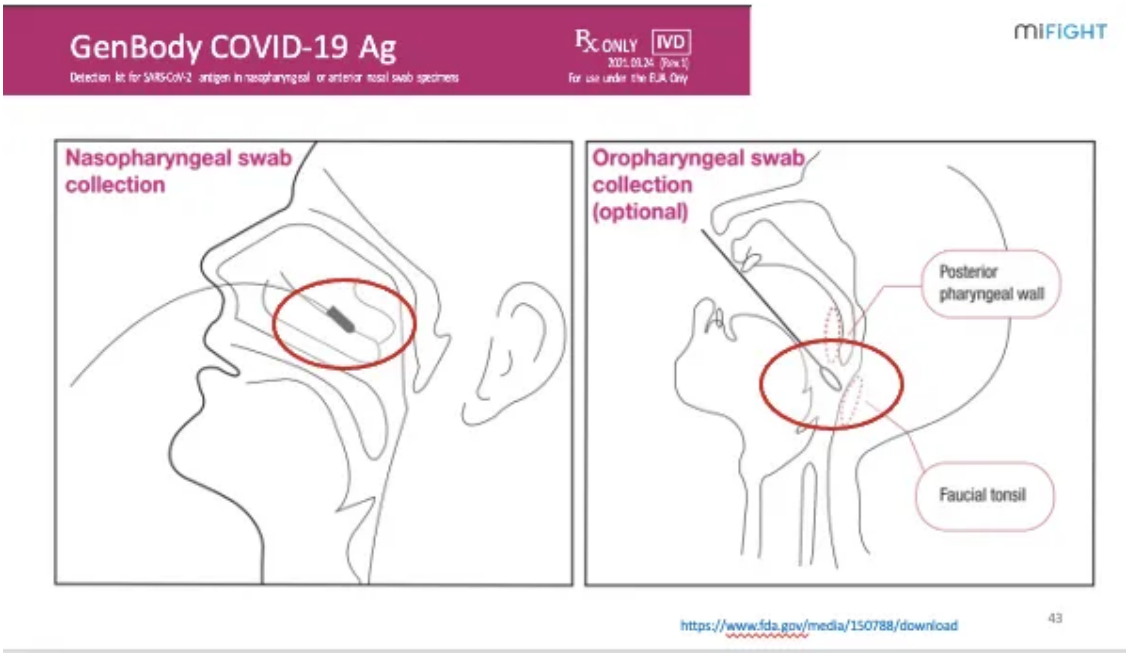
The pretend SARS-CoV-2 EUA PCR-tests were crucial 'theatre props' in convincing Americans that the *COVID-19 Story* was a real pandemic.

The demand for obsessive testing by schools was and is a very dark form of emotional and psychological abuse.

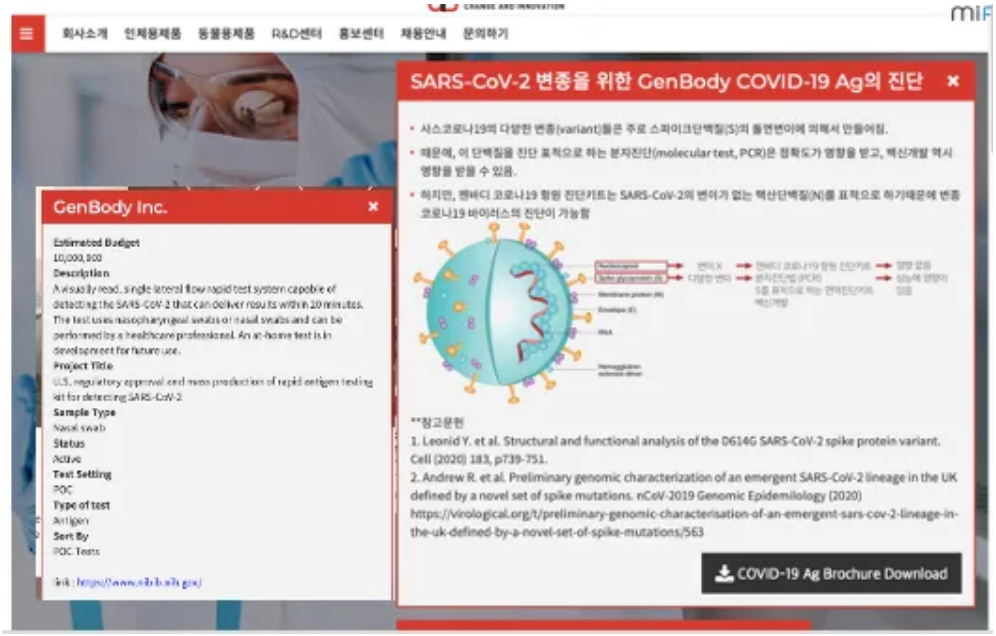
Across our nation, perfectly healthy children were 'sick until proven healthy,' being subjected to regular, invasive genetic sampling procedure in order to attend schools that we now know are 'pretend virus tests'.

These pictures are from the COVID-19 testing that was being conducted at my son's school, often *without parents' permission.*





My son's school was part of a $10,295,000 NIH funded genetic-research study being conducted by GenBody, a company based in South Korea. The student's DNA, and my son's DNA, was being sent to South Korea without parents' permission and paid for by US tax dollars.



> ***I hope this makes it clear that the PCR-testing at schools was never about the health and safety of the students and faculty. It was to collect the genetic data of America's most valuable treasure, our children, to send to foreign nations to further develop genetic weapons against America.***

The PCR-test was the perfect theatre prop to convince American's that there was an invisible killer who at first....discriminated against and targeted 'unvaxxed humans.'

Later on in the COIVD-19 story, the invisible SARS2 killer became in-discriminatory of humans' vaccine status and could strike any one, of any age, without testing or warning.

The obsessive PCR-testing increased the worry about COVID-19 to a state of paranoia.

More than two years later, after *The COVID-19 Story* debut in March of 2020 in America, we are now beginning to watch the documentary, "*The Making of the COVID-19 Story*." We see that the mRNA SARS-CoV-2 virus was not only ***not deadly and not infectious***, ***but was a fictitious story to distract us from the parasitic Ai bioweapons unleashed on humanity***.

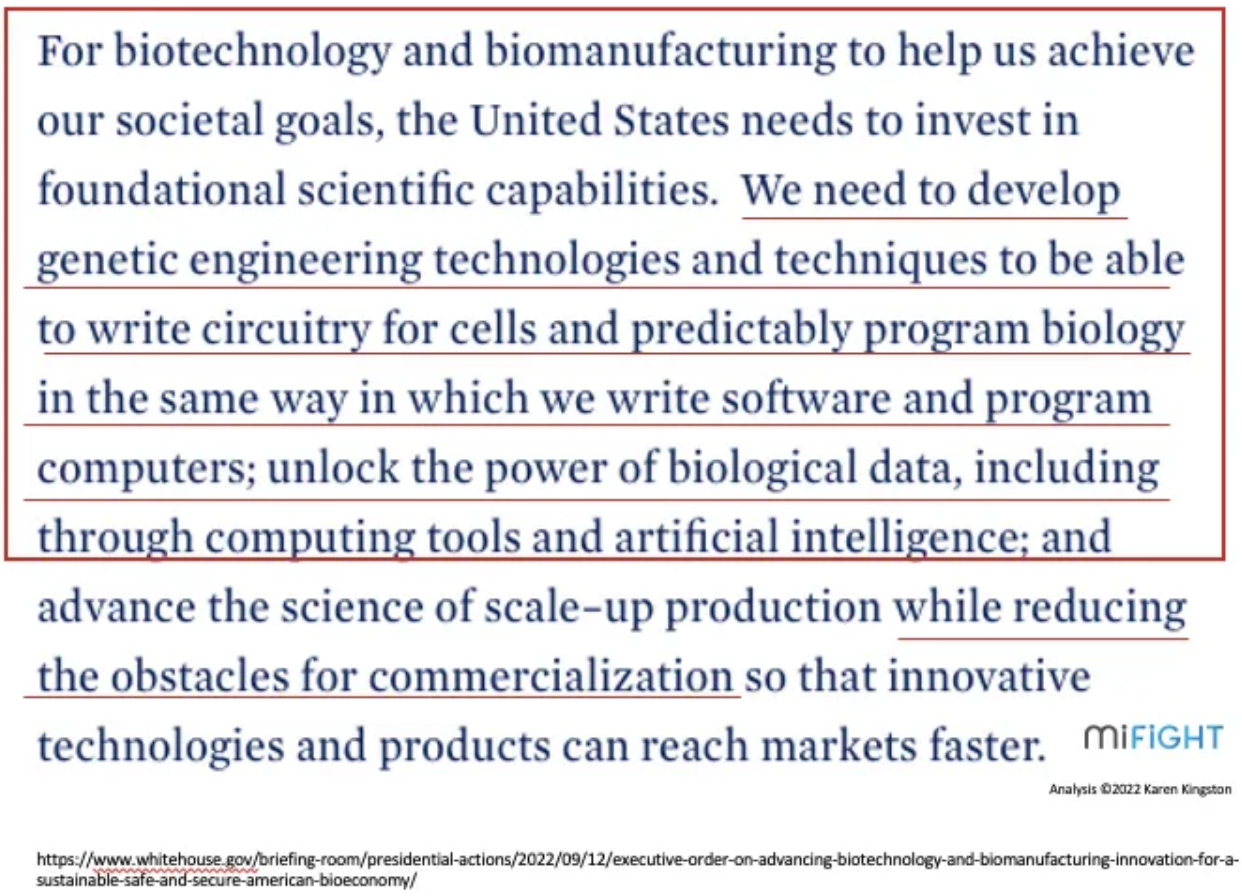
> Why are leaders and influencers focusing on a narrative about a respiratory virus, variants and the risks of biological proteins being produced by mRNA 'vaccines', which we know

> are bioweapons? Do the experts not see the evidence that people are magnetic? Do they not see that people are suffering bizarre central nervous system attacks and hallucinations?

When scientists and engineers are documenting self-assembling nanotechnology from vaccine vial samples and blood samples of those vaccinated, are we supposed to blindly follow credentialed experts instead believing of real world evidence?

We are being gaslit into believing The COVID-19 Story, featuring; the SARS-CoV-2 virus, Variants, and Spike Proteins, instead of being informed about the real threat of the parasitic Ai bioweapons that have been unleashed on Americans and all of humanity.

Perhaps Joe Biden signed the Biotechnology and Biomanufacturing Executive Order on September 12, 2022, declaring that, "*We need to develop genetic engineering technologies and techniques to be able to write circuitry for cells and predictably program biology in the same way in which we write software and program computers…*" because Americans have already been installed with the '*genetic engineering technology that can write circuitry for the cells and program biology*.'



The distraction around the SARS-CoV-2 virus, variants, and spike proteins, combined with the denial of the existence of the parasitic Ai bioweapons is one of the most evil deceptions in the history of humanity.

I pray that we repent, reunite under God, and boldly proclaim the TRUTH against the lies, deception, and evil of the COVID-19 Ai bioweapons that have been unleashed on America and humanity.

It's Time we Reunite as Humanity and Take Down COVID-19.

- STOP all COVID-19 emergency powers and financial funding
- Ban and recall all COVID-19 products and technologies
- Take Down 5G and StarLink

If you would like to support the Take Down of COVID-19, please join miFight.

A Message from the Apostle Paul

Have nothing to do with the fruitless deeds of darkness, but rather expose them. Everything exposed by the light becomes visible—and everything that is illuminated becomes a light.

This is why it is said: "Wake up, sleeper. Rise from the dead and Christ will shine on you."

The Kingston Report. TRUTH WINS.

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27 Comments

Write a comment...

Cheryll Reina Oct 17

I was born in Anniston, Alabama. We had the glorious Monsanto, Anniston Army Depot and Fort McClellan with the chemical school. I am legally disabled. I started researching in 2010. Now I know why me and so many others that spent time in Anniston are so sick. I have lost many I knew to cancer and many I know are currently battling cancer. I made a documentary on Anniston and released it in April. Through my research I realized Pfizer had purchased Monsanto