

The Scam Has Been Confirmed: PCR Does Not Detect SARS-CoV-2

Posted on:

Monday, December 14th 2020 at 10:45 am

Posted by **Green Medicine** Information



Unofficial English translation of 'Frauds and falsehoods in the medical field' originally published on www.dsalud.com

The genetic sequences used in PCRs to detect suspected SARS-CoV-2 and to diagnose cases of illness and death attributed to [Covid-19](#) are present in dozens of sequences of the human genome itself and in those of about a hundred microbes. And that includes the initiators or primers, the most extensive fragments taken at random from their supposed "genome" and even the so-called "target genes" allegedly specific to the "new coronavirus". The test is worthless and all "positive" results obtained so far should be scientifically invalidated and communicated to those affected; and if they are deceased, to their relatives. Stephen Bustin, one of the world's leading experts on PCR, in fact says that under certain conditions anyone can test positive!

We have been warning you since March: you cannot have specific tests for a virus without knowing the components of the virus you are trying to detect. And the components cannot be known without having previously isolated/purified that virus. Since then we continue to accumulate evidence that no one has isolated SARS-CoV-2 and, more importantly, that it can never be isolated for the reasons we explained last month (read the report "[Can you prove that there are pathogenic viruses?](#)" on our

website -www.dsalud.com-). And in the present report we are going to offer new data that show that RT-PCR does not detect the so called SARS-CoV-2 as it is known, but fragments of human RNA and those of numerous microbes.

We have already explained the numerous problems that RT-PCR poses, recognized by organizations or governments such as the *WHO* or the *CDC* and by prestigious international experts such as **Dr. Stephen Bustin** who considers both the arbitrariness of establishing criteria for results and the choice of the number of cycles to be nonsense because they can lead to anyone testing positive.

In this report we are going to add the results of a particular research we have done from the data published on the alleged SARS-CoV-2 and on the protocols endorsed by the *WHO* for the use of RT-PCR as well as the data corresponding to the rest of the "human coronaviruses". And the conclusions are extremely serious: **none of the seven "human coronaviruses" have actually been isolated** and all the sequences of the primers of their respective PCRs as well as those of a large number of fragments of their supposed genomes are found in different areas of the human genome and in genomes of bacteria and archaea, such as these: *Shewanella marina JCM*, *Dialister succinatiphilus*, *Lactobacillus porcine*, *Lactobacillus manihotivorans*, *Leptospira sarikeiensis*, *Bizionia echini*, *Sanguibacteroides justesenil*, *Bacteroides massiliensis*, *Lacinutrix venerupis*, *Moraxella bovis*, *Leptospira saintgironisiae*, *Winogradskyella undariae*, *Acetobacterium puteale*, *Chryseobacterium hispanicum*, *Paenibacillus koleovorans*, *Tamiana fuccidanivorans*, *Fontibacillua panacisegetis*, *Ru bacter ruber*, *Skemania piniformis*, *Chryseobacterium shigense*, *Caloramator peoteoclasticus*, *Cellulosilyticum ruminicola*, *Nitrosopumilius evryensis* and a long list of others.

We are going to explain step by step the research that has led us to such an unusual conclusion.

HAVE ANY HUMAN CORONAVIRUSES BEEN ISOLATED?

During the first half of April, when the first research we conducted indicated that SARS-CoV-2 had not been isolated and since those who claimed to have done so were relying on "isolates" of previous "human coronaviruses", we began to do a thorough review of those claimed isolates. Specifically, we reviewed the alleged isolation work of suspected human coronaviruses **229E** (said to have been isolated in 1965), **OC43** (in 1967), **SARS-CoV** (in 2003), **NL63** (in 2004), **HKU1** (in 2005) and **MERSCoV** (in 2012). And these have been the results:

Coronavirus 229E.

Reference article: Dorothy Hamre and John Procknow. *A new virus isolated from the human respiratory Tract*. Proceedings of the Society for Experimental Biology and Medicine, 121: 1:190-193. January 1, 1966.

Since the authors refer to other articles to explain the method of isolation - which they call *Complement Fixation* - we consulted a reference article for that method: that of **Janet W. Hartley et al.** *Complement Fixation and tissue culture assay for mouse*

leukaemia viruses PNAS, 53(5):931-938, May 1965. This is a procedure already in disuse that uses the antigen-antibody reaction to detect either one or the other. In the case we are dealing with, the aim was to detect the antigens of the supposed new virus but, as we have already explained, specific antibodies are needed which cannot be obtained the first time a virus is detected.

Coronavirus OC43.

Reference article: Paul Lee. *Molecular epidemiology of human coronavirus OC43 in Hong Kong*. Thesis for the Department of Microbiology, University of Hong Kong, August 2007. The HKU Scholars Hub.

What was considered to be viral RNA was extracted from cultures without any proof that the RNA belongs to a virus. The tool used - a QIAamp kit - removes reagents, inhibitors and contaminants but what it cannot do is determine where the extracted RNA comes from. And there are no controls. It is then amplified by PCR and sequenced assuming (!) that it is genetic information of a virus. Finally, the author speculates about mutations, recombinations, genotypes, molecular evolution, strains and other jargon that conveys the idea -unproven- that a "virus" is being worked with.

SARS-CoV Coronavirus.

Reference article: J. S. M. Peiris and others. *Coronavirus as a possible cause of SARS*. Lancet 361: 1319-25, April 2003.

There is no mention of purification in the article. There is not even any mention of filtration or centrifugation. It is only stated that "*the viruses were isolated in fetal monkey liver cells from nasopharyngeal aspirates and lung biopsies of two patients*". There are no controls. The only mention is of a "cytopathic effect" that is attributed to a virus and that PCR was done for known viruses and retroviruses without obtaining results. Finally, RT-PCR was done with "random initiators" and a sequence "of unknown origin" is detected to which "*a weak homology with the coronaviridae family*" is found. Then they designed primers for that sequence and when testing 44 samples from [SARS](#) patients only 22 were positive.

Coronavirus NL63.

Reference article: Lia van der Hock and others. *Identification of a new human coronavirus*. Nature Medicine, 10, 4 April 2004.

The authors state that "*the identification of unknown pathogens using molecular biology tools is difficult because the target sequence is not known so that PCR-specific initiators cannot be designed*".

What they used is a tool they developed themselves called VIDISCA which, they claim, does not require prior knowledge of the sequence! Is that possible? Let's see how it works: first the culture is prepared and it is assumed that a virus is present due to the evidence of "cytopathic effect". The novelty introduced by this method is that "restriction enzymes" are added, enzymes that cut the nucleic acid molecules at certain locations and always by the same length. In this way, if after the action of these enzymes they

observe many fragments of DNA or RNA that are the same or very similar, they deduce that it comes from a virus, since the host genome would present random cuts, while the virus genome presents a large number of copies that are the same due to the replication of the virus. And is such a deduction correct? Of course not! This assumption (which adds to the previous assumption that there is a virus) does not take into account that there are "virus-like particles", "retrovirus-like particles", "endogenous retroviruses", "exosomes", "extracellular" particles and even mitochondrial DNA. In denial, there are a multitude of particles that possess the same reproductive characteristics in large quantities as "viruses" and therefore can falsify results by producing large numbers of identical copies when cut by enzymes as recognized in an article on the VIDISCA technique entitled *Enhanced bioinformatic proSling of VIDISCA libraries for virus detection and Discovery. It was published in volume 263 of Virus Research on April 2, 2019, and it's authors-Cormac M. Kinsella et al.-recognize that "no redundancy is expected in the VIDISCA insert from the host background nucleic acid except in the case of 'virus-like' characteristics, i.e., high copy numbers as in mitochondrial DNA.*

Coronavirus HKU1.

Reference article: Patrick C. Y. Woo and others. *Characterization and Complete Genome Sequence of a Novel Coronavirus, Coronavirus HKU1, from Patients with Pneumonia.* Journal of Virology, 79, 2, January 2005.

The article, incredibly, begins with these words: "*Despite extensive research in patients with respiratory tract infections, no microbiological cause has been identified in a significant proportion of patients. RNA is extracted from non-purified cultures.*" And a PCR with coronavirus genes is used. For the sequencing they use two protein databases organized in families, domains and functional sites -PFAM and INterProScan- combined with two computer programs that carry out "predictions" on how nucleotides should be combined. The text adds: "*The sequences were manually assembled and edited to produce a final sequence of the viral genome.*" And once again there are no controls.

MERS-CoV Coronavirus.

Reference article: Ali Moh Zaki and others. *Isolation of a Novel Coronavirus from a Man with Pneumonia in Saudi Arabia.* The New England Journal of Medicine, 367:19, November 2012.

The genetic material is extracted directly from the culture supernatant and sputum sample with a tool called *High Puré Viral Nucleic Acid Kit* and then tested with different PCRs for various known microorganisms. There is no mention of purification and there are no controls.

In short, what had been done with the first coronaviruses -and with many other supposed viruses- is to cultivate supposedly infected tissues - any "cytopathic effect" was attributed to the presence of a virus only - and then either some proteins are obtained which without any test are considered "virus antigens" and when these

"antigens" are detected in cultures it is interpreted as "isolation", or fragments of nucleic acids are extracted assuming that they belong to a virus.

We already explained in the article published in the previous issue of the magazine that according to Dr. Stefan Lanka the so-called "cytopathic effect" is actually an effect caused by the conditions of the culture itself. This is recognized for example in the article *Antibiotic-induced release of small extracellular vesicles (exosomes) with surface-associated DNA* published on August 15, 2017 on the website of *Nature* and signed by Andrea Németh and others

https://www.ncbi.nlm.nih.gov/pmc/articles/PMC5557920/pdf/41598_2017_Article_8392.pdf

It explains that certain substances -such as antibiotics- added to in vitro experiments can stress the cell cultures so that they generate new sequences that had not been previously detected. This had already been noticed by none other than Dr. Barbara McClintock in 1983 during her Nobel Prize lecture, as can be seen at <https://www.nobelprize.org/uploads/2018/06/mcclintock-lecture.pdf>

In essence, **NOT ONE OF THE SEVEN SUPPOSED HUMAN CORONAVIRUS HAS REALLY BEEN ISOLATED.** The only thing that has been different between them are the laboratory procedures and techniques that were becoming progressively more sophisticated which, in this case, has implied not a greater accuracy but a greater capacity for deception and self-deception that has culminated in the virtual manufacture of the SARS-CoV-2.

And the obvious consequence of the lack of evidence of its isolation is that such "coronaviruses" cannot be held responsible for any disease. Moreover, all tests - of whatever kind - based on the presumed components of these "viruses" (nucleic acids or proteins) are completely disqualified as "infection tests" and even more as "diagnostics" of diseases.

MORE UNANSWERED REQUESTS

In the previous issue we already collected the answers given by the authors of several articles that supposedly described the isolation of SARS-CoV-2 in which they acknowledged that they had not "purified" which implicitly means acknowledging that the virus was not isolated. And now we are going to add one more piece of evidence: the responses given by different authorities - political and health - from various countries about the purification and isolation of SARS-CoV-2.

James McCumiskey -author of the book *'The Latest Conspiracy: The Biomedical Paradigm'*- tells us that the *National Virus Reference Laboratory of Ireland* requested information about it from the *University of Dublin* and the latter responded that "*it has no records that could provide an answer to their request*". The director of legal services of the laboratory insisted on his request to the university and the university responded as follows: "*The position of the university is that material of academic debate cannot be subject to the Freedom of Information Act*". It follows from the NVR's request that they

have not cultivated SARS-CoV-2 or purified it. They only acknowledge having "*detected SARS-CoV-2 RNA in diagnostic samples.*"

On June 22, a group of experts sent a consultation in similar terms to British Prime Minister Boris Johnson. The letter was signed by Dr. Kevin Corbett, Piers Corbyn - professor at *Imperial College* London -, the engineer and independent researcher - who we interviewed in the journal at the time - David Crowe, Dr. Andrew Kaufman, the Edinburgh professor of biology Roger Watson and the biologist and chemist David Rasnick - and to this day they still have not received a reply!

Another similar request - in this case to the *National Research Council of Canada* - received the following response: "*We have not been able to carry out a complete search of the NRC's records so we regret to inform you that no records have been identified that respond to your request.*"

We will add that two journalists have been sending similar requests - under the Freedom of Information Act - to various institutions in Canada, New Zealand, Australia, Germany, the United Kingdom and the United States, and as of September 5, twelve institutions have responded, all indicating the same thing: that they have no record of work describing the isolation of the virus that is supposed to cause Covid-19. The details and the answers can be seen at

<https://www.fluoridefreepeel.ca/u-k-dept-of-health-and-social-care-has-no-record-of-covid-19-virus-isolation/>

LOOKING FOR THE ORIGIN OF THE FALSE GENOME

The question we asked ourselves then was: if the sequences that have been published do not belong - as claimed- to new viruses, where do they come from? And to try to answer that question we decided to carry out a search with a computer program called *Basic Local Alignment Search Tool (BLAST)*, a sequence alignment search tool that allows us to compare a given sequence with all the sequences stored in the *National Institutes of Health of the United States* (it is public and can be consulted at <https://blast.ncbi.nlm.nih.gov/Blast.cgi>). We explain step by step what we did so that our readers can repeat the search for themselves and check the results.

First we collected all the initiators of the PCRs described in the protocols hosted on the *WHO* website at the time which were these:

China *CDC* protocol: uses ORF1ab and N genes as target.

Protocol of the *Pasteur Institute* (France): uses two fragments of the RdRP (which is supposed to be SARS.CoV-2 specific).

United States *CDC* protocol: uses three fragments of the N gene.

Protocol of the *National Institute of Infectious Diseases* of Japan: it is the only one that has as target the S gene together with other genes supposedly shared with other coronaviruses.

Charite Protocol (*Germany*): uses the E, N and RdRP genes.

Hong Kong University Protocol: uses ORF1b-nsp14 and N gene.

National Institute of Health Thailand protocol: uses the N gene.

We then introduced the sequence of the primers - the one that indicates the beginning of the sequence to be detected (forward) and the one that indicates the final (reverse) - into the *BLAST* so that it could search for them in two databases: a collection of microbe genomes and the one corresponding to the human genome.

THE SEQUENCES OF THE SO-CALLED SARS-COV-2 ARE FOUND BOTH IN HUMANS AND IN NUMEROUS MICROBES!

Let's see in detail the procedure taking as an example the initiators of the French protocol. Once on the *BLAST* website, we chose *Microbes* to search the microbial genome databases and moved to the next page. Then a form appeared in which we entered the sequence of the forward initiator of the French protocol -that is ATGAGCTTAGTCCTGTG-, we selected the option *Highly similar sequences* and pressed the *BLAST* key. Just a few seconds later the results appeared -we took a screenshot (*image 1*)- and we were shown 100 sequences of microbes -particularly bacteria and archaea- with a coincidence of between 77% and 100% with an identity percentage of 100%.

We then returned to the home page and that second time we chose *Human* to search the human genome, we repeated the same operation and after a few seconds the result appeared which we screen captured again (*image 2*). And it turns out that the sequence entered coincides with 74 sequences of the human genome, with a coincidence of between 66% and 100% and a percentage of identity of 100%.

And that indicates that the sequence of that initial PCR primer that is supposed to be specific to SARS-CoV-2 actually corresponds to 74 fragments of the human genome and a hundred microbial fragments as well!

We then decided to repeat the operation but with the final or reverse primer - which is CTCCCTTTGTGTGTGT - and the results were similar.

Since these were very short sequences -about twenty genetic letters or nucleotides- we decided to try again but with the target sequence defined by these two primers, i.e. the sequence of the supposed SARS-CoV-2 genome that is between the initial primer and the final primer. Obviously, for this we needed the sequence that is officially claimed to be the "SARS-CoV-2 genome" and although thousands of laboratories claim to have isolated and sequenced it -a false claim as we have explained in previous reports- we decided to go to the *National Centre for Biotechnology Information* website: https://www.ncbi.nlm.nih.gov/nuccore/NC_045512.2?report=genbank&to=29903.

Once there, we located the "target sequence", a fragment of 108 nucleotides located between positions 12,690 and 12,797 of the "genome", which is this one:

ATGAGCTTAGTCCTGTTGCACTACGACAGATGTTGTGCCGGTACACAAACTGCTTG
CACTGAT GACAATGCGTTAGCTTACAACAACAAAGGGAG.

With this we repeated the steps previously described and the results were again surprising since there appeared again a hundred microbe sequences with a percentage of a match of 100% and four sequences of the human genome with an identity percentage between 83% and 95%. The matches were therefore lower but the important thing is that we continue to find fragments of the supposed "target sequence" of SARS-CoV-2 both in microbes and in our own genome.

Truly astonished we took a further step and tested with the gene considered at that time as the most specific of SARS-CoV-2, the E gene that is supposed to generate the envelope proteins and is located between positions 26,245 and 26,472:

ATGTA CTCA TTCGTTTCGGAAGAGACAGGTACTACGTTAATAGTTAATAGCGTACTT
CTCTTGCTTTCGTGGTATTCTTGCTAGTTACTAGCCATCCTGCTTCGATTGTGCG
TACTGCTGCAATATTGTTAACGTGAGTCTTGTA AACCTTTACGTTTACTCGTGTTA
AAATCTGAATTCTTCTAGAGTTTCGATTCTGGTCTAA.

We repeated with it the steps already described and the result was even more surprising because despite its length another hundred microbe sequences appeared with a percentage of identity of 100% and 10 sequences of the human genome with a percentage of identity between 80% and 100%. And similar results were obtained with a fragment chosen at random and with the N gene which they say corresponds to the proteins of the SARS-CoV-2 nucleocapsid.

We finally decided to test with the S gene which is said to generate the structural "spike" proteins that are key to entry into the cell and was subsequently considered to be the most specific SARS-CoV-2 gene. Since it is a gene whose sequence is much longer - 3821 nucleotides between positions 21,563 and 25,384 - we tested with two fragments chosen at random within that gene and the first - TTGGCAAATTCAAGACTCACTTTC - resulted in another hundred microbe sequences and 93 sequences of the human genome and the second - CTTGCTGCTACTAAATGCAGAGTGT - a hundred microbial sequences and 90 of the human genome.

Finally we decided to test with the initiators of the Japan Protocol, the only one that includes target sequences of the S gene and, the reader will have already guessed, the results were once again similar: a hundred microbe sequences and 93 sequences of the human genome with an identity percentage between 94.12% and 100%!

CONCLUSIONS

The consequence of all that we have just explained is clear and immediate: **THERE IS NO VALID TEST TO DETECT SARS-COV-2**, neither antibody or antigen tests nor RT-PCR. And we included those based on the supposed gene that codes for the S1 or spike protein. And that means that

ALL THE NUMBERS OF "CASES", "INFECTED", "SICK", "Asymptomatic" OR "DEAD DUE TO COVID-19" LACK A SCIENTIFIC BASE AND ALL "POSITIVES" ARE FALSE POSITIVES, something that should be communicated immediately to those affected and those responsible should be held accountable.

We end by adding that even the *WHO* itself does not really believe in these tests. Just read the document published last September 11 as a laboratory guide for SARS-CoV-2 entitled *Diagnostic tests for SARS-CoV-2* - it is available at <https://apps.who.int/iris/rest/bitstreams/1302661/retrieve> - and it literally says on page 5: "*Whenever possible, suspected active infection should be tested with a nucleic acid amplification test (NAAT) such as RT-PCR. NAAT tests should target the SARS-CoV-2 genome but since there is no known global circulation of SARS-CoV-1 a Sarbecovirus sequence (presumed to include at least five human and animal coronaviruses including SARS-CoV-1 and SARS-Cov-2) is also a reasonable target*". That is, *WHO* agrees to use non-specific sequences to detect SARS-CoV-2.

That is not all because the manual later states, "*An optimal diagnosis consists of a NAAT test with at least two genome-independent targets of the SARS-CoV-2; however, in areas where transmission is widespread, a simple single-target algorithm can be used.*"

The *WHO* manual states, "*One or more negative results do not necessarily rule out SARS-CoV-2 infection. There are a number of factors that can produce a negative result in an infected individual including poor quality of the sample, late collection of the sample, inadequate handling, or technical reasons inherent in the test, such as mutation of the virus or inhibition of PCR.*"

What are the judges waiting for to act on their own initiative?
Jesus Garcia Blanca

Note: the author publicly thanks Juan Pedro Aparicio Alcaraz for his patient and meticulous collaborative work in the search for scientific articles and for his tedious work with the BLAST.

THIS REPORT APPEARS IN [\(https://www.dsalud.com/revistas/numero-242-noviembre-2020/\)](https://www.dsalud.com/revistas/numero-242-noviembre-2020/)

The media hyping of a vaccine is getting pushback all around the world. Mike Lord writing for 'The Liberty Beacon' shares how this Government's COVID Hoax hangs by a thread as credibility evaporates:

"The British government's offensive against its own people ran into deeper trouble today as its credibility continued to evaporate.

Trust in the government as a source of information is a key and indispensable component of any propaganda war. If the source putting out the propaganda becomes mistrusted, disbelieved or discredited, the propaganda war is lost.

The Johnson regime's problems center around its continued reliance upon the PCR test to produce stats that purport to show a "spread" of infection.

It was a bitter blow for the government's psychological warfare operation when the PCR test became exposed a fraudulent. The setback arose from revelations of gross inaccuracy, its tendency to produce false positives by the thousand and – this was the utterly damning clincher – the government admitting on one of its own websites that the PCR test actually **CANNOT** detect a live infectious virus.

This renders all "positives" that relied on the PCR test invalid.

That in turn renders all diagnoses of a COVID19 "case" based on a PCR positive result fraudulent.

It is believed now that the number of "false cases" may number tens of thousands and may include the overwhelming majority of all alleged cases.

One observer has pointed out that, "if on a particular day the government reports, say, 500 new cases based on the results of PCR tests, the actual number of real cases is far less and could, theoretically, even be zero."

This in turn means that the "infections" stat used as the justification for the damaging lockdowns such as the one that London has just been hit with are fabrications.

The PCR test is considered the most inaccurate of all the tests in use yet none of the others are particularly reliable: the PM and ministers have for instance all admitted that the test used at airports is only 7% accurate, which is still superior to the PCR test, although tossing a coin has been described as about seven times more likely to return a correct diagnosis than the PCR test.

The writer and keen observer of government, Kieron McFadden said that,

"It is entirely possible, considering the results of a test so wildly misleading have been depended on as the primary tactic for convincing the nation that an epidemic is progress, there may be no epidemic at all. A bug, yes, that causes severe problems for a small number of already frail people – as do many other bugs – but an epidemic? No. What is coming to light is a hoax of almost unimaginable duplicity and inflicted damage." Yet the government continues to use it and its false results to perpetuate one of the most transparent and destructive frauds in British history.

It is unclear at this time why the government does so and the government does not seem about to own up and come clean any time soon.

One theory is that it is reluctant to change because the PCR test has enabled it to manipulate the stats for many months – or at least it did until the scam was exposed.

Meanwhile, the criminals operating under the cover of government continue to declare stringent and damaging lockdowns on the basis of the false tests, a further fraud involving massive costs to the nation and its citizens.

Again, the reasons for doing this are unclear except that the continued desperate portrayal of an epidemic is considered vital to enforcing dangerous vaccines on the populace.

That drive has itself run into trouble as many scientists and medical professionals have sounded the alarm over the dangers of injecting a large number of people with vaccines that have not been fully studied and tested both for safety and efficacy.

Should serious adverse reactions become apparent over the next months and years, many people involved in or collaborating with the drive to inject these substances into people may become liable to criminal charges involving humanitarian crimes.

In an effort to somehow keep its flawed show on the road, the government today invented the “discovery” of a new strain of the virus to “explain” the rise in its false infections stats.

One of the many problems the government is facing is that it needs “rising” infections to justify hitting areas and cities with more lockdowns and restrictions but those alleged rising infections – if believed – declare that previous measures involving lockdowns, distancing, masks and so forth were of no use.

Therefore declaring that the virus has mutated gets the government off that particular hook for a short while – they hope . Yet this in turn has laid the Health Secretary, whose credibility is already near zero open to further ridicule in the wake of his excruciating, embarrassingly phony shedding of crocodile tears on live TV a few days earlier.

One source told us that,

“The whole government caper has descended into Pythonesque farce that would be hilarious were it not for the damage and misery this corrupt government is causing the nation.”

Other reports attest that grass roots anger at the government's deceit and trickery is reaching boiling point, as we witness the spectacle of a government that appears to combine being incredibly inept with shamelessly subversive.



The way in which this fake virus has been fear-mongered upon the public, suppression of any rebuttal, from experts in their fields of science, censorship of anything that raises question to the alleged Covid-19 virus, we must take serious the issue of vaccines. In a related report, I learned that Autism is now 1 in 18 children. This was noted in a report by a family doctor who works in the area of mitigating vaccine damaged children.

Dr. Don Boys is a Christian and ran a Christian school in Indianapolis, Indiana and understands Bible Prophecy well enough to raise questions about the insanity of the rush to produce an unproven nMRNA vaccine and disregard protocol required to produce a vaccine in virtually weeks instead of the traditional 5-7 years to bring a vaccine to market. This is all so unorthodox and disregards all safety procedures. Such suggests that there is more to the vaccine than the public has been led to believe.

The Koch Postulates have been avoided by the pharmaceutical industry. As this article has shown, the entire process of testing has been faulty from the get go. 100,000 doctors worldwide are protesting the careless method being used to roll out the vaccine. For months now, I have been unable to download and save articles from doctors, researchers, virologists, and people in fields of immunity, biochemistry, molecular biology. I have purchased nearly a dozen books on the topics and clearly, this is being driven by globalism and the World Economic Forum and the Vatican pressuring people into receiving their unproven “injections”. We have no transparency or honesty by those driven by profit motives and those with a history of eugenics in their background.

KOCH'S POSTULATES

A microorganism can be accepted as the causative agent of an infectious disease only if the following conditions are satisfied,

- ▶ **The bacterium should be consistently associated with the lesions of the disease**
- ▶ **It should be possible to isolate the bacterium in pure culture from the lesions**
- ▶ **Inoculation of such pure culture into suitable laboratory animals should reproduce the lesions of the disease.**
- ▶ **It should be possible to re isolate the bacterium in pure culture from the lesions produced in the experimental animals.**

Dr. Don Boys is the host of the web page “All News Pipeline” just posted thought provoking about the Covid-19 virus being the “Mark” of the Beast. I wrote numerous articles on Bill Gates and his emphasis on creating the Quantum Dot/Hydrogel ID2020;

and, I believe Dr. Don Boys has been one of the major voices of Truth seeking to expose what certainly resembles the Biblical “Mark” of the Beast.



MARK OF THE BEAST Makes Its Debut in 2020.....as the Untested, Toxic and Unsafe COVID-19 Vaccine Bioengineered by the *Big Pharma*, the Premier Huckster of the “Super-Vaccination Agenda”.

[State of the Nation](#)

There’s a very good reason (actually very BAD reason) why THE GREAT SCAMDEMIC, THE GREAT RESET and THE GREAT STEALECTION were all carried out by the *New World Order* globalist cabal in 2020. It’s the very same reason why 2019 saw the military deployment of 5G in key power centers around the globe.

The amount and degree of sheer chaos and confusion, distraction and diversion, which have been generated by these three “GREATs”, has permitted the power elite to both execute the military deployment of 5G worldwide and roll out a dangerously untested COVID-19 vaccine with very little scrutiny.

The Powers That Be have long planned to roll out both 5G and the COVID-19 vaccine during the same time frame because they function as a highly coordinated weapon system. See: [QUATERNARY WEAPON SYSTEM Activated Before Each Coronavirus Cluster Explosion](#)

MARK OF THE BEAST

- **Revelation 13:17 (KJV)**
17 And that no man might buy or sell, save he that had the mark, or the name of the beast, or the number of his name.
- **Revelation 14:11 (KJV)**
11 And the smoke of their torment ascendeth up for ever and ever: and they have no rest day nor night, who worship the beast and his image, and whosoever receiveth the mark of his name.

What the world community of nations has been undergoing since January of 2019 is an immense beta test involving a highly sophisticated bioweapon — COVID-19 — which has been quite stealthily launch at primarily urban targets where 5G roll-outs occurred (e.g. Wuhan, Milan, Tehran, New York City, Seattle and Guayaquil, Ecuador).

Then, against all odds and contrary to what science says is even possible, the COVID-19 vaccine has been rolled out first in Russia and then in other guinea pig nations such as the United Kingdom and United States.

How could such a highly suspect and unproven vaccine even be safely administered in the same year that the coronavirus pandemic first began?!

That's right, it can't be! Not with any degree of safety or effectiveness.

And, the only way *Big Pharma* was able to introduce any type of Covid vaccine in 2020 is because it had been undergoing years of painstaking development and advanced bioengineering in secret labs across the planet.

MARK OF THE BEAST

With this crucial understanding, every inhabitant on Earth needs to be aware that the COVID-19 vaccine is not what they say it is. And, that it is dangerous to the extreme with a nefarious agenda behind it that should scare the bejesus out of everyone.

In fact, everything now points to a Covid vaccine that not only functions as the long anticipated MARK OF THE BEAST, it's also a means by which TPTB will implement their high-priority agenda of transhumanism. See: [COVID-19 Vaccine Utilizes Nano Tech to Greatly Advance Transhuman Agenda](#)

Perhaps the central pillar of that agenda is to remove any tendency on the part of a normal human being to pursue spiritual advancement or religious endeavors. This is where the COVID-19 vaccine comes in because its primary function is to permanently turn off the famous GOD gene. This will be accomplished through the systematic vaccination of humanity using the FunVax jab, also known as the COVID-19 vaccine. As follows:

The FunVax bioweapon system is using at least 4 different modalities. The perpetrators are (i) using chemtrails to spray flu viruses and aerosol vaccines, (ii) administering adult flu shot programs and childhood vaccination schedules, (iii) releasing several variants of the COVID-19 bioweapon in hospitals, nursing homes, assisted-living facilities (and on cruise ships when they were operating), and (iv) utilizing the intensive 5G roll-outs to weaken the immune system as well as amplify the electromagnetic frequencies and microwave transmissions that facilitate the clandestine FunVax vaccination process. Also, the perps have planned a fifth component whereby they are (v) formulating a mandatory COVID-19 vaccine which will ensure that every individual receives the VMAT2 'immunization' that will be enforced via an immunity certificate. (Source: [Here's the main reason why the NWO globalist cabal is dead set on mandating a universal COVID-19 vaccination program.](#))

One of the unique features of the MARK OF THE BEAST is Luciferase.

Bill Gates is building something that we call the Human Implantable Quantum Dot Microneedle Vaccination Delivery System, and it is composed of multiple things. I want to draw your attention to one component, the quantum dot microneedles that will deliver the vaccines, and a very, very unique biochemical that makes it all work. If you're standing up while reading this, you might want to sit down. Today we bring you 'near infrared bioluminescence enzyme luciferase' which is the chemical that will make the quantum dot vaccination readable through a special mobile device app. That's right, the enzyme that will light up Bill Gates Human Implantable Quantum Dot Microneedle Vaccination Delivery System is called Luciferase, that's what makes the vaccination readable long after the victim has been injected. Luciferase.(Source: [SHOCK AS IT'S REVEALED AN ENZYME CALLED LUCIFERASE IS WHAT MAKES BILL GATES IMPLANTABLE QUANTUM DOT MICRONEEDLE VACCINE DELIVERY SYSTEM WORK](#))

COVID-19 Vaccine Compliance

The highly organized global criminal enterprise to vaccinate every human being has been in the works for decades. It's why vax super-pitchman Bill Gates was made king of Microsoft and then the world's richest billionaire. See: [Yes, Bill Gates' invention really is an injectable, permanent, under-the-skin, biochip sensor?](#)

In point of fact, Gates' carefully guided transition from covertly constructing back doors for computer viruses to deceitfully creating back doors for corona viruses took place with highly purposeful design.

[MARK OF THE BEAST: Brought to you by everyone's Big Brother—BILL GATES](#)

However, it's one thing to fabricate a coronavirus pandemic out of nowhere as a means to justify a new mandatory COVID-19 vaccine ... and ... another thing to enforce the administration of that untested jab on every inhabitant of planet Earth.

This is where things are about to get very challenging in view of the fact that [two-thirds of voters say they won't get coronavirus vaccine right away or at all.](#)

The psychopathic perpetrators of [OPERATION COVID-19](#) know full well that many folks are extremely leery of a Covid vaccination. So many Americans are suspicious of the COVID-19 vaccine because of the association it has with the MARK OF THE BEAST (MOTB). This poses a great obstacle which has already been thought through by the likes of Bill Gates, Dr. Fauci, Deborah Birx, Robert Redfield, Tedros Adhanom & Company.

That "Company" is, of course, *Big Pharma*. And *Big Pharma* is the key corporate player in the all-powerful [Government-Corporate Complex That Took Complete Control Of The USA.](#)

What's the critical point?

Every U.S. citizen will soon need a COVID-19 vaccine to live a normal life in America...
... unless *We the People* terminate this steamrolling juggernaut post-haste.

Slowly but surely both *Corporate America* and the U.S. Federal Government are floating different proposals about how an immunity certificate will soon be required.

For example, if you want to work for any Fortune 1000 company, proof of vaccination will be requested if the perps get their way. Eventually every publicly traded corporation will be coerced into this pen of vaccine compliance. So will small companies that need a public health inspection and/or certification to 'legally' operate in a city, county or state.

Retail stores may begin to require that all customers produce proof of Covid immunity to buy their goods. So might every service business such as medical, dental, mental health and other healthcare delivery offices both allopathic and alternative.

Then there are workout gyms and sport clubs, yoga and Pilates studios, massage parlors and hair salons. Getting in any of these is already quite difficult without a mask. Then there are the calls for all school children to be vaccinated in order to enter kindergarten, elementary school and junior high. Teenagers will be vaxxed first to enroll

in high schools and young adults will be required to present an immunity certificate to even apply to any American college or university.

The list of organizations throughout civil society as well as governmental institutions is even longer, as are all the globalist NGOs that will eventually demand a Covid vaccine to be employed by them.

SPECIAL NOTE: People, the gravity of this swiftly developing predicament whereby proof of a COVID-19 vaccination will soon be required to “buy and sell” anything cannot be overstated. Just consider where the Coviteers are ultimately taking this: A Covid immunity certificate will soon be required to enter any establishment; attend any event; matriculate in any program; fly, train or bus to any destination; be treated by any hospital, clinic, physician or other healthcare practitioner, etc. They named it *Operation Warp Speed* for a reason; they want all 50 states operating in strict COVID-19 vaccine compliance as a precursor to legislating mandatory vaccines for every citizen and non-citizen alike. The sheer speed and intensity with which the perpetrators of OPERATION COVID-19 are rolling over the American people with their Covid juggernaut is as incomprehensible as it is staggering. Hence, every individual needs to be on guard like never before. Blindly complying with all of the illicit orders and other Covid dictates ought to be vigilantly avoided. The more folks bend to the will of the *Plandemic* tyrants, the more easily it will be for them to implement their pre-planned compulsory COVID-19 vaccination programs that will inject the dreaded MOTB. Most importantly, this insidious vaccination scheme will be ever-so-gradually carried out so that very few even understand that it will eventually be necessary to live your life. That’s how they always introduce destructive, dangerous and deadly programs into society.

This is exactly why the authorities are demanding masks and social distancing with such a vengeance: so they can eventually offer relief from these terribly oppressive measures after everyone has been ‘properly’ vaccinated.

This is their PLAN!

And it ain’t good. Not good at all.

These perfidious perps are quite cunningly choreographing and staging every phase of this transparently manufactured manmade *Plandemic* in the interest of implementing various mandates and measures which, they hope, will eventually push every person in the pen of vaccine compliance. As always, the *Mainstream Media* is used to hype every phase of naked fear-mongering in order to stampede the unaware and uneducated over the cliff of inevitable vaccine injury and permanent transhumanistic alteration.

The government knows it cannot force vaccines on everyone; but they have confidence that a critical mass of the citizenry will demand them; and to the extent that the overwhelming peer pressure exerted by the insane pro-vax crowd will compel even the most ardent anti-vaxxers into a receiving their annual Covid vaccination.

KEY POINT: *Big Pharma* is being used by the Central Banking Cartel and International Banking Crime Syndicate to roll out the COVID-19 vaccine under the governmental rubric of *Operation Warp Speed*. At “warp speed” because the power elite are so desperate to vaccinate the global population in order to effectively neutralize them because of the rapidly emerging context defined by the ongoing and raging war between the 1% and 99%. See: [So this is what they mean by “WARP SPEED”!](#)

Conclusion

The NWO globalists are on a serious mission to vaccinate every American with all deliberate speed. The US government will employ the 4 major branches of the Armed Forces to accomplish this mission.

TPTB know that when a sufficient number of Americans are regularly vaccinated, they will no longer be a threat to their kleptocratic oligarchy that intends to rule over the entire planetary civilization via a **One World Government** and Global Security Superstate.

This *New World Order* is fundamentally defined by a world community of nations, each of which is run by totalitarian communist government. That’s why the despotic Democrats are so determined to steal the 2020 election and that this **RED ALERT** has been issued:

[THE 2020 COVID-COMMUNIST PLOT TO OVERTHROW THE AMERICAN REPUBLIC ...](#)
[... AS A PRELUDE TOWARD ESTABLISHING A ONE WORLD GOVERNMENT AND GLOBAL TOTALITARIAN SUPERSTATE](#)

The globalists also know that all resistance to such a totalitarian tyranny will be removed once the MARK OF THE BEAST is put into full operation, **where no one can buy or sell without it.**

[COVID-19 Vaccine is the “Mark of the Beast”](#)

Warning: There is now a HUGE vaccination campaign going on across the USA within the Christian churches. It’s being craftily spearheaded by none other than Dr. Fauci’s boss at the NIH. This particular government-religious initiative is being deceptively conducted to corral the entire Christian community into the pen of vaccine compliance because they pose the greatest impediment to the communist scheme to collapse the American Republic. Those Bible-reading folks also know more about the MARK OF THE BEAST than any other group in the country See: [HUGE COVID SCANDAL AT THE NIH: Why has no one investigated Fauci’s boss?](#)

[State of the Nation](#)

December 13, 2020

Reference

[Is the COVID-19 vaccine really the “Mark Of The Beast”?](#)

Recommended Reading

FUNVAXgate: Bill Gates' Implantable Quantum Dot, Microneedle Vaccine Delivery System & Luciferase

TinyURL: <https://bit.ly/2WeaQWH>

<http://stateofthenation.co/?p=41277>

Blessings,

Pastor Bob, EvanTeachr@aol.com

www.pastorbobreid.com

<http://jesusisthewaythetruththelife.com/node/22>