

The Failure to Prove the Virus Exists

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Over a year ago, I proposed (insisted on) a procedure to prove SARS-COV-2 exists.

This procedure is essential—and needless to say, it hasn't been done, and will never be done.

Why? Because the outcome could completely and utterly destroy the COVID narrative.

Here is the procedure: You line up 500 people who have been diagnosed with COVID-19, and you take tissue samples from them.

You properly process these samples, through centrifuging, etc., in order to extract and arrive at what you believe is the virus.

You put that material under an electron microscope and photograph it.

You then place the 500 photos from the 500 “pandemic patients” side by side.

You ask yourself three burning questions.

One: In each and every photo, are there many identical viruses?

Two: Are these viruses in every one of the 500 photos?

Three: Is the virus one you've never seen before?

If the answer to question one and two is yes, you appear to have found a common virus for the 500 patients. If the answer to three is yes, it's a virus never seen before.

If the answer to either question one or two is no, you've failed to find the common virus you're looking for. You've failed to prove a viral cause for what you're calling COVID-19.

If you see many identical virus particles in some, but not all, of the photos, you may or may not have found a virus. To decide that issue, you need three conditions: the researchers are honest and independent; a new team of such researchers will repeat the whole procedure, from the beginning, to see whether their findings match those of the original team; and you need truly qualified experts to determine whether the particles in the electron microscope photos are actually viruses or something else.

Note: This is why one or two photos from a study mean NOTHING.

All right. Moving on, there are other factors involved in the process of discovering a virus. These factors are ISOLATION and GENETIC SEQUENCING.

They're both covered in a Statement on Virus Isolation, authored and published by Dr. Andrew Kaufman, Dr. Tom Cowan, and Sally Fallon Morell. I reprint it here in full:

Statement On Virus Isolation (SOVI) [1]

"Isolation: The action of isolating; the fact or condition of being isolated or standing alone; separation from other things or persons; solitariness." – Oxford English Dictionary

The controversy over whether the SARS-CoV-2 virus has ever

been isolated or purified continues. However, using the above definition, common sense, the laws of logic and the dictates of science, any unbiased person must come to the conclusion that the SARS-CoV-2 virus has never been isolated or purified. As a result, no confirmation of the virus' existence can be found. The logical, common sense, and scientific consequences of this fact are:

- * the structure and composition of something not shown to exist can't be known, including the presence, structure, and function of any hypothetical spike or other proteins;
- * the genetic sequence of something that has never been found can't be known;
- * "variants" of something that hasn't been shown to exist can't be known;
- * it's impossible to demonstrate that SARS-CoV-2 causes a disease called Covid-19.

In as concise terms as possible, here's the proper way to isolate, characterize and demonstrate a new virus. First, one takes samples (blood, sputum, secretions) from many people (e.g. 500) with symptoms which are unique and specific enough to characterize an illness. Without mixing these samples with ANY tissue or products that also contain genetic material, the virologist macerates, filters and ultracentrifuges i.e. purifies the specimen. This common virology technique, done for decades to isolate bacteriophages [2a] and so-called giant viruses in every virology lab, then allows the virologist to demonstrate with electron microscopy thousands of identically sized and shaped particles. These particles are the isolated and purified virus.

These identical particles are then checked for uniformity by physical and/or microscopic techniques. Once the purity is determined, the particles may be further characterized. This would include examining the structure, morphology, and

chemical composition of the particles. Next, their genetic makeup is characterized by extracting the genetic material directly from the purified particles and using genetic-sequencing techniques, such as Sanger sequencing, that have also been around for decades. Then one does an analysis to confirm that these uniform particles are exogenous (outside) in origin as a virus is conceptualized to be, and not the normal breakdown products of dead and dying tissues. [2b] (As of May 2020, we know that virologists have no way to determine whether the particles they're seeing are viruses or just normal break-down products of dead and dying tissues.) [2c]

If we have come this far then we have fully isolated, characterized, and genetically sequenced an exogenous virus particle. However, we still have to show it is causally related to a disease. This is carried out by exposing a group of healthy subjects (animals are usually used) to this isolated, purified virus in the manner in which the disease is thought to be transmitted. If the animals get sick with the same disease, as confirmed by clinical and autopsy findings, one has now shown that the virus actually causes a disease. This demonstrates infectivity and transmission of an infectious agent.

None of these steps has even been attempted with the SARS-CoV-2 virus, nor have all these steps been successfully performed for any so-called pathogenic virus. Our research indicates that a single study showing these steps does not exist in the medical literature.

Instead, since 1954, virologists have taken unpurified samples from a relatively few people, often less than ten, with a similar disease. They then minimally process this sample and inoculate this unpurified sample onto tissue culture containing usually four to six other types of material – all of which contain identical genetic material as to what is called a “virus.” The tissue culture is starved and poisoned and naturally disintegrates into many types of particles, some

of which contain genetic material. Against all common sense, logic, use of the English language and scientific integrity, this process is called "virus isolation." This brew containing fragments of genetic material from many sources is then subjected to genetic analysis, which then creates in a computer-simulation process the alleged sequence of the alleged virus, a so called in silico genome. At no time is an actual virus confirmed by electron microscopy. At no time is a genome extracted and sequenced from an actual virus. This is scientific fraud.

The observation that the unpurified specimen – inoculated onto tissue culture along with toxic antibiotics, bovine fetal tissue, amniotic fluid and other tissues – destroys the kidney tissue onto which it is inoculated is given as evidence of the virus' existence and pathogenicity. This is scientific fraud.

From now on, when anyone gives you a paper that suggests the SARS-CoV-2 virus has been isolated, please check the methods sections. If the researchers used Vero cells or any other culture method, you know that their process was not isolation. You will hear the following excuses for why actual isolation isn't done:

1. There were not enough virus particles found in samples from patients to analyze.
2. Viruses are intracellular parasites; they can't be found outside the cell in this manner.

If No. 1 is correct, and we can't find the virus in the sputum of sick people, then on what evidence do we think the virus is dangerous or even lethal? If No. 2 is correct, then how is the virus spread from person to person? We are told it emerges from the cell to infect others. Then why isn't it possible to find it?

Finally, questioning these virology techniques and conclusions is not some distraction or divisive issue. Shining the light

on this truth is essential to stop this terrible fraud that humanity is confronting. For, as we now know, if the virus has never been isolated, sequenced or shown to cause illness, if the virus is imaginary, then why are we wearing masks, social distancing and putting the whole world into prison?

Finally, if pathogenic viruses don't exist, then what is going into those injectable devices erroneously called "vaccines," and what is their purpose? This scientific question is the most urgent and relevant one of our time.

We are correct. The SARS-CoV2 virus does not exist.

—end of Kaufman, Cowan, Morell Statement—

Finally, here is a repost of my article about a claim of virus isolation. Dr. Kaufman does a step-by-step analysis of a quote from a typical study that purports to describe how SARS-CoV-2 was isolated:

—Dr. Andrew Kaufman refutes "isolation" of SARS-Cov-2; he does step-by-step analysis of a typical claim of isolation; there is no proof that the virus exists—

The global medical community has been asserting that "a pandemic is being caused by a virus, SARS-Cov-2."

But what if the virus doesn't exist?

People have been asking me for a step-by-step analysis of a mainstream claim of virus-isolation. Well, here it is.

"Isolation" should mean the virus has been separated out from all surrounding material, so researchers can say, "Look, we have it. It exists."

I took a typical passage from a published study, a "methods" section, in which researchers describe how they "isolated the

virus.” I sent it to [Dr. Andrew Kaufman](#) [3], and he provided his analysis in detail.

I found several studies that used very similar language in explaining how “SARS-CoV-2 was isolated.” For example, “*Severe Acute Respiratory Syndrome Coronavirus 2 from Patient with Coronavirus Disease, United States, (Emerging Infectious Diseases, Vol. 26, No. 6 – June 2020)*” [4].

First, I want to provide a bit of background that will help the reader understand what is going on in the study.

The researchers are creating a soup in the lab. This soup contains a number of compounds. The researchers assume, without evidence, that “the virus” is in this soup. At no time do they separate the purported virus from the surrounding material in the soup. Isolation of the virus is not occurring.

They set about showing that the monkey (and/or human cells) they put in the soup are dying. THAT’S THEIR KEY “EVIDENCE.” This cell-death, they claim, is being caused by “the virus.” However, as you’ll see, Dr. Kaufman dismantles this claim.

There is no reason to infer that SARS-CoV-2 is in the soup at all, or that it is killing cells.

Finally, the researchers assert, with no proof or rational explanation, that they were able to discover the genetic sequence of “the virus” they never isolated. “We didn’t find it, we don’t know anything about it, but we sequenced it.”

Here are the study’s statements claiming isolation, alternated with Dr. Kaufman’s analysis:

STUDY: “We used Vero CCL-81 cells for isolation and initial passage [in the soup in the lab]...”

KAUFMAN: “Vero cells are foreign cells from the kidneys of monkeys and a source of contamination. Virus particles should be purified directly from clinical samples in order to prove

the virus actually exists. Isolation means separation from everything else. So how can you separate/isolate a virus when you add it to something else?"

STUDY: "...We cultured Vero E6, Vero CCL-81, HUH 7.0, 293T, A549, and EFKB3 cells in Dulbecco minimal essential medium (DMEM) supplemented with heat-inactivated fetal bovine serum (5% or 10%)..."

KAUFMAN: "Why use minimal essential media, which provides incomplete nutrition [to the cells]? Fetal bovine serum is a source of foreign genetic material and extracellular vesicles, which are indistinguishable from viruses."

STUDY: "...We used both NP and OP swab specimens for virus isolation. For isolation, limiting dilution, and passage 1 of the virus, we pipetted 50 μ L of serum-free DMEM into columns 2–12 of a 96-well tissue culture plate, then pipetted 100 μ L of clinical specimens into column 1 and serially diluted 2-fold across the plate..."

KAUFMAN: "Once again, misuse of the word isolation."

STUDY: "...We then trypsinized and resuspended Vero cells in DMEM containing 10% fetal bovine serum, 2 \times penicillin/streptomycin, 2 \times antibiotics/antimycotics, and 2 \times amphotericin B at a concentration of 2.5×10^5 cells/mL..."

KAUFMAN: "Trypsin is a pancreatic enzyme that digests proteins. Wouldn't that cause damage to the cells and particles in the culture which have proteins on their surfaces, including the so called spike protein?"

KAUFMAN: "Why are antibiotics added? Sterile technique is used for the culture. Bacteria may be easily filtered out of the clinical sample by commercially available filters (GIBCO) [5]. Finally, bacteria may be easily seen under the microscope and would be readily identified if they were contaminating the sample. The specific antibiotics used, streptomycin and

amphotericin (aka 'ampho-terrible'), are toxic to the kidneys and we are using kidney cells in this experiment! Also note they are used at '2X' concentration, which appears to be twice the normal amount. These will certainly cause damage to the Vero cells."

STUDY: "...We added [*not isolated*] 100 µL of cell suspension directly to the clinical specimen dilutions and mixed gently by pipetting. We then grew the inoculated cultures in a humidified 37°C incubator in an atmosphere of 5% CO₂ and observed for cytopathic effects (CPEs) daily. We used standard plaque assays for SARS-CoV-2, which were based on SARS-CoV and Middle East respiratory syndrome coronavirus (MERS-CoV) protocols..."

STUDY: "When CPEs were observed, we scraped cell monolayers with the back of a pipette tip..."

KAUFMAN: "There was no negative control experiment described. Control experiments are required for a valid interpretation of the results. Without that, how can we know if it was the toxic soup of antibiotics, minimal nutrition, and dying tissue from a sick person which caused the cellular damage or a phantom virus? A proper control would consist of the same exact experiment except that the clinical specimen should come from a person with illness unrelated to covid, such as cancer, since that would not contain a virus."

STUDY: "...We used 50 µL of viral lysate for total nucleic acid extraction for confirmatory testing and sequencing. We also used 50 µL of virus lysate to inoculate a well of a 90% confluent 24-well plate."

KAUFMAN: "How do you confirm something that was never previously shown to exist? What did you compare the genetic sequences to? How do you know the origin of the genetic material since it came from a cell culture containing material from humans and all their microflora, fetal cows, and

monkeys?”

—end of study quotes and Kaufman analysis—

My comments: Dr. Kaufman does several things here. He shows that isolation, in any meaningful sense of the word “isolation,” is not occurring.

Dr. Kaufman also shows that the researchers want to use damage to the cells and cell-death as proof that “the virus” is in the soup they are creating. In other words, the researchers are assuming that if the cells are dying, it must be the virus that is doing the killing. But Dr. Kaufman shows there are obvious other reasons for cell damage and death that have nothing to do with a virus. Therefore, no proof exists that “the virus” is in the soup or exists at all.

And finally, Dr. Kaufman explains that the claim of genetic sequencing of “the virus” is absurd, because there is no proof that the virus is present. How do you sequence something when you haven’t shown it exists?

Readers who are unfamiliar with my work (over 300 articles on the subject of the “pandemic” during the past year [6]) will ask: Then why are people dying? What about the huge number of cases and deaths? I have answered these and other questions in great detail. The subject of this article is: have researchers proved SARS-CoV-2 exists?

The answer is no.

SOURCES:

[1] <https://www.andrewkaufmanmd.com/sovi/>

[2a] Isolation, characterization and analysis of bacteriophages from the haloalkaline lake Elmenteita, KenyaJuliah Khayeli Akhwale et al, PLOS One, Published: April

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<https://journals.plos.org/plosone/article?id=10.1371/journal.pone.0215734> – accessed 2/15/21

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<https://www.frontiersin.org/articles/10.3389/fcell.2020.573511/full> – accessed 2/15/21

[2c] “The Role of Extraellular Vesicles as Allies of HIV, HCV and SARS Viruses,” Flavia Giannessi, et al, Viruses, 2020 May

[3] <https://andrewkaufmanmd.com/>

[4] https://wwwnc.cdc.gov/eid/article/26/6/20-0516_article

[5] <https://www.thermofisher.com/us/en/home.html>

[6] <https://blog.nomorefakenews.com/category/covid/>

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