

Thought piece - SCIENTIFIC MISCONDUCT - call to all people that believe in the scientific method!

Dear scientist, researcher, investigator,

We invite you to consider the following argumentation objectively and either disprove or adopt it. Please do not outsource this investigation to others, even if it will take some of your precious time. If you are a virologist or bioinformatician, fantastic! But you do not need to be an expert in order to **critically research and think logically to form your own opinion**. Please do not just listen to the “experts” or “trust the science”. Please do not just read one factchecker article or the defamation of a critical voice. Many people have given up their own ability to think for themselves, but not you, right? Do your own research, **convince yourself with facts** and uncover the truth for yourself.

Scientific misconduct

Research institutions define misconduct as “fabrication, falsification, or plagiarism in proposing, performing, or reviewing research, or in reporting research results.” “**Fabrication**” is defined as “**making up data and results**”.¹

Corona crisis - matters of fact

- a) Covid-19 has been associated with common symptoms of sickness and exclusively defined by the PCR test protocol, which has been used in an unscientific way.
- b) The PCR test protocol was created by Dr. Corman and Dr. Drosten and his team without having a virus isolate as reference.² His paper was not peer-reviewed and his role in the editorial board of the publishing journal represents a serious conflict of interest.
- c) The Corman-Drosten-PCR tests were recommended by the WHO, but were, against common laboratory practice, run at too high cycle thresholds, and led to the fabrication of alleged “Covid-19 cases”.
- d) The alleged SARS-CoV-2 virus genome was fabricated by assembling short genetic sequences from an unpurified sample of one Wuhan patient into a long hypothetical genome construct, instead of biochemically characterising the isolated and purified alleged virus particles.
- e) All electron micrograph pictures show common cell breakdown particles as alleged new viruses.

Additionally, **the scientific method** was and still is **violated** by the fact that **no control experiments**, to attempt to falsify the virus genome hypothesis of the new Coronavirus, have been performed. Self-control by possible and replicable experiments is a central requirement in natural sciences and a worldwide consensus at least since 1998. A lack thereof violates good research practice.³

¹ Book: On Being a Scientist: Third Edition, 2009, National Academy of Sciences
https://graduateschool.nd.edu/assets/21763/on_being_a_scientist.pdf

² Corman, Victor M et al. “Detection of 2019 novel coronavirus (2019-nCoV) by real-time RT-PCR” Euro surveillance: European communicable disease bulletin vol. 25,3 (2020) p. 2: “In the present case of 2019-nCoV, **virus isolates or samples from infected patients have so far not become available** [...]. [...] diagnostic workflow for 2019-nCoV screening and **specific confirmation, designed in absence of available virus isolates or original patient specimens.**”
<https://www.ncbi.nlm.nih.gov/labs/pmc/articles/PMC6988269/>

³ DFG - Code of Conduct: “Guidelines for Safeguarding Good Research Practice” (2019)
https://www.dfg.de/en/research_funding/principles_dfg_funding/good_scientific_practice/ p. 4: “The **recommendations** set out in the **1998** white paper initiated a system of self-monitoring and voluntary commitment [...] that has enjoyed **broad consensus** to this day. [...] serves as the **basis for the Code**, which also draws on international reference works, and describes appropriate standards for research in the form of guidelines.” & DFG - Memorandum: Safeguarding Good Scientific Practice (1998)
https://www.dfg.de/download/pdf/dfg_im_profil/reden_stellungnahmen/download/empfehlung_wiss_praxis_1310.pdf p. 74:



Applying the scientific method unveils consensus virus theory as flawed!

People are getting sick, but what is the cause? Let us hypothesise that a new virus is causing sickness (environmental, lifestyle, psychological factors excluded). An important part of the scientific method is subjecting hypotheses to rigorous tests. Experiments need to be designed to gather appropriate data to be able to conclude that the virus is indeed the cause.

Virologists have never done this and therefore violate the scientific method!

We believe virologists have formed a **consensus that assumes the existence of viruses** is proven by two methods. Firstly, **virus isolation**; and secondly, **virus genome assembly**. The problem is, both methods are applied unscientifically.

1. **Virus isolation:** A cell culture experiment where **genetic materials from different sources are mixed**. Contrary to the meaning of the word “isolation”, a mixture of cow RNA from calf serum, monkey RNA from Vero cells, human liver tumour RNA from HUH-7 cells, unspecified bacteria and/or antibiotics is used. **Virologists fail to use isolated and purified virus particles from sick patients!**
2. **Virus genome assembly:** **Short genetic sequences** from different origins (no isolate!) **are assembled into a long genome** (hypothetical construct!). This is claimed to be viral based on the observation of the **cytopathic effect** (dying of cells), but in reality, **caused by the conditions of the experiment** itself (starvation and poisoning). Genome alignments to previous virus genomes (hypothetical constructs!) lead to **circular reasoning**, i.e., if enough random genetic material is collected, it can be aligned to any previously created virus genome.

The lack of control experiments in both “virus proof” methods is unscientific!

For details and to convince yourself (or not!) from these astonishing conclusions, please review **Dr. Lanka’s argumentation** in relation to virus isolation, sequencing and control experiments:

- [The Virus Misconception. Part 1](#)⁴
- [The Virus Misconception. Part 2](#)⁵
- [The Virus Misconception. Part 3](#)⁶
- [The Initiators of the Corona Crisis Have Been Clearly Identified: VIROLOGISTS](#)⁷

The universal hypothesis that **viruses cause disease can no longer be upheld**, as clearly laid out by Dr. Lanka. His conclusions are the following:

- No virus has ever been isolated⁸ according to proper scientific methods⁹
- The observed ‘cytopathic effect’ is not caused by a virus but the design of the virus isolation¹⁰ experiment itself; Dr Lanka has proven this twice now: Firstly, in 2016 with the independent expert

“Findings, wherever factually possible, **must be controlled and replicated** before being submitted for publication.” & p. 94:

“**Systematically doubting one’s own findings, however, is at the core of scientific method. Repetition of experiments** – if possible, independently – is **particularly important** [...]”

⁴ <http://wissenschaftplus.de/uploads/article/wissenschaftplus-the-virus-misconception-part-1.pdf>

⁵ <http://wissenschaftplus.de/uploads/article/wissenschaftplus-the-virus-misconception-part-2.pdf>

⁶ <http://wissenschaftplus.de/uploads/article/wissenschaftplus-the-virus-misconception-part-3.pdf>

⁷ <http://wissenschaftplus.de/uploads/article/The-Initiators-of-the-Corona-Crisis-Have-Been-Clearly-Identified.pdf>

⁸ FOIs reveal that health/science institutions around the world (165 and counting!) have no record of SARS-COV-2 isolation/purification, anywhere, ever <http://tinyurl.com/norecordfound>

⁹ Statement On Virus Isolation (SOVI) <https://andrewkaufmanmd.com/sovi/>

¹⁰ <https://www.britannica.com/science/separation-and-purification> “separation of a substance into its components and the removal of impurities”



opinion no. 5 brought forward in the Lanka-Measles trial on 10th Feb. 2016.¹¹ Secondly, in 2020 when performing several SARS-CoV-2 control experiments single-handedly.¹²

- The observed and photographed particles are not viruses but common human and experiment-induced deceasing cell fragments
- The calculated genome of viruses is not real but rather invented by assembling small pieces of genetic sequences of multiple sources (human, animal, drugs) by utilising simulation software
- The sequence alignment of viruses is always done with reference to a given virus genome that has previously also been computationally generated
- The invented 'in silico' (in a computer) virus genome is always derived from a mixed cell culture or an unpurified patient sample, and never directly from an isolated and purified sick-patient sample (violating 'Koch's postulates' 1. and 2.)
- Control experiments with 'uninfected' saliva/blood are leading to the same 'cytopathic effect', but are never performed (falsifiability cannot be perpetuated!)
- Control experiments for electron micrographs of "viruses" are never done, i.e., shown particles are never biochemically characterised by filtration, ultracentrifugation and gel electrophoresis to determine if the whole genome of the visualised particles matches the virus (replicability cannot be upheld!)
- Control experiments with different 'short read metagenomic assembler software' are leading to different results for the virus genome when using the same samples (replicability cannot be upheld!)

The above points are valid for all alleged viruses from the Adeno-associated virus, to the Polio and the Zika virus. **Thanks to Corona, obvious proofs of fabrication can be found in the original SARS-CoV-2 paper by Fan Wu et al** that established the SARS-CoV-2 virus genome.¹³

- Alleged virus isolation from one single patient, but no virus isolation or purification (filtration, centrifugation) of particles from the patient sample was performed (no repeatability) → unscientific!
- Researchers apply meta-transcriptomic sequencing and randomly select the longest contig sequence → unscientific fabrication!
- Two different software algorithms lead to drastically different genome outcomes (30,474 nt in Megahit versus 11,760 nt in Trinity, i.e., 62% shorter!) → unscientific fabrication!
- Alignment to a bats coronavirus ensures similarity of SARS-CoV-2 to a bats-virus, i.e., circular reasoning (no controls!) → unscientific fabrication!
- PCR method with very high cycle thresholds of 40 was used for determination and confirmation (PCR > 30 is meaningless = "dirty"!) → unscientific fabrication!

Relevant citations:

"The viral genome organization of WHCV was determined by **sequence alignment to [...] a coronavirus associated with bats** (bat SL-CoVZC45, GenBank accession number MG772933) [...]." (p.266)

"[...] its **whole genome** sequence (29,903 nt) has been assigned GenBank accession number MN908947." (p.266)

¹¹ Sequencing report by an independent laboratory in Germany (and confirmed by a second laboratory) "Opinion on the nucleic acid sequence of the 'Measles virus' filed as evidence in the Lanka-Measles trial that Dr. Lanka won: Federal Court of Justice (BGH) file number: I ZR 62-16 as of 1.12.2016 http://wissenshaftplus.de/uploads/article/BGH_Beschluss_I_ZR_62-16.pdf & Higher Regional Court Stuttgart file number: 12 U 63/15 as of 16.02.2016 http://lrbw.juris.de/cgi-bin/laender_rechtsprechung/document.py?Gericht=bw&GerichtAuswahl=Oberlandesgerichte&Art=en&sid=46bf3db2df690aba6e4874acaf45b6&nr=20705&pos=0&anz=1

¹² WISSENSCHAFTPLUS magazine 2/2021 (in German only) https://www.wplus-verlag.ch/de_DE & explanatory video [watch from min 42:20] DR. TOM COWAN - SESSION 1: THE CONTAGION MYTH (westonaprice) <https://www.bitchute.com/video/iL8IUrarkOg0/>

¹³ Wu, Fan et al. "A new coronavirus associated with human respiratory disease in China." Nature vol. 579,7798 (2020): 265-269. <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC7094943/>



“[...] **isolation** of the virus **from only a single patient** is not sufficient to conclude that it caused these respiratory symptoms [...].” (p.268)

“the **longer sequence** (30,474 nt)—which covered almost the whole virus genome—**was used for primer** design for PCR confirmation and **determination of the genome termini.**” (p. 270)

“The viral loads of WHCV in BALF were determined by quantitative **real-time RT–PCR** [...] **by 40 cycles** [...].” (p. 270)

Conclusions

The above points highlight that **common virus theory contradicts the scientific method** and **violates the Code of Conduct for every scientist**. Virologists fail to design, conduct and report the necessary experiments to either prove or disprove their hypotheses.

This might be one of the most significant aberrations of our time. To err is human, but to disregard critical arguments is ignorant and unscientific. We know that many questions remain unanswered, but to reject a flawed hypothesis, it is not necessary to provide a complete explanation.

We appeal to your honour – please address this potential scientific misconduct

Please support Dr. Lanka in his efforts to disseminate this information widely. Please report this knowledge to your superiors, colleagues and personal contacts. Please demand from responsible virologists and institutions that the strictly necessary experiments are conducted immediately.

We hope for your pure conscience, moral awareness and a sense of right and wrong.

Yours sincerely,
Corona_fakta_DK

P.S. If you would like to follow all information around this topic, we have created the following Danish information channel on the **Telegram App** (available on Android, iOS and Windows): **t.me/Corona_fakta**

Alternatively, you can find many immensely well-researched **information articles** here (feel free to use translation services for your own language):

In German <https://telegra.ph/Corona-Fakten-Liste-der-wichtigsten-Artikel-09-08>

In English <https://telegra.ph/Corona-Investigative-09-26>

