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In the military complaints process

of ...

and the...

not only for the justification of the hearing complaint, but also for the public and posterity, it is essential to state how dangerous the Covid-19 injections are, which the judges of the 1st Military Service Senate gave their judicial blessing on July 7th, 2022, and this, although the expert Prof. Dr. Ulrike Kämmerer also expressly referred to the bioweapon quality of the Covid-19 injections based on modRNA technology in her lecture on the 2nd day of the hearing on June 7th, 2022.

As a reminder, we're talking about modRNA here because this RNA is man-made.

Mrs. Prof. Dr. Kämmerer summarizes her statements in this regard in court on June 7, 2022, with which she justified the absolute unacceptability and irresponsibility of this Covid 19 "vaccination" obligation, as follows (quote):

"The website <https://www.stopgof.com/> describes in detail how the variant of SARS-CoV-2 was assembled in the laboratory as part of the so-called gain-of-function research. The operator of the site, together with other scientists, has now submitted a publication that proves in a technically complex and detailed manner that almost 100% of the SARS-CoV-2 virus must have come from a biological laboratory, available under the link:

<https://www.biorxiv.org/content/10.1101/2022.10.18.512756v1.full.pdf>

Furthermore, there are the following indications from the specialist literature:

A work published in 2018 from the laboratory of Christian Drosten (he is the last author and therefore the person responsible) (Muth et al; SciRep. 2018; 8: 15177) describes how in the SARS-1 strain Frankfurt 1 through targeted manipulation ("engineering ") different variants of the virus have been produced. So he knows exactly how such viruses are processed in the laboratory and made "armed".

These "engineered viruses" were then tested for their infectivity for human airway cell cultures and the "rank order" of the ability to replicate in these human cells was determined. This work could also have contributed to defining a variant that is more dangerous for humans.

With regard to the spike protein of SARS-CoV-2, it should be noted that the gene information, which is injected into the body in large numbers of copies by means of modRNA (Pfizer/Biontech or Moderna) or DNA (AstraZeneca or Janssen), the formation of a protein that corresponds exactly to the sequence of the original spike protein of the Wuhan-1 variant at the amino acid level (and in spatial protein structure).

This Wuhan-1 spike protein contains multiple sites that leave no doubt as to its origin in a gain-of-function laboratory variant. These obvious places have been described in detail in the publications by Mrs. Segreto (Segreto R, Bioessays. 2021 Mar; 43(3): 2000240; Bioessays. 2021 Jul; 43(7): 2100015; Environ Chem Lett. 2021; 19(4): 2743-2757) and also on the stopgof.com website

The following artificial properties that make the spike more toxic to humans have been explicitly discovered and described so far:

1. The so-called "codon usage", i.e. the bases in the nucleic acid that code for the amino acids, are artificially adapted (probably through cell culture or passage in humanized mouse models) to human genetic information, so that the protein can be produced particularly effectively in the cells .

2. In addition to the unusual receptor binding domains, which are obviously optimized for improved binding to the human ACE2 receptor, the Wuhan-1 spike (and thus also the "vaccination" spikes also has a receptor binding site for neuropilin-1) and thus, otherwise not found in corona viruses, a connection possibility to nerve cells, which can explain the increased neurotoxicity (loss of taste, paralysis, fatigue, neuropathies) of the original viruses, but above all of the "vaccines".

3. The "furin cleavage site" is undoubtedly and generally recognized as the most conspicuous feature of the virus spike, which was only observed in the Wuhan variant (and so far no other human pathogenic SARS viruses). This cleavage site was already described as conspicuous in one of the first publications (Wölfel et al, Nature. 2020 May;581(7809):465-469.). Firstly, this furin cleavage site allows a fragment of the spike protein to detach and circulate freely in the body (this spike fragment contains most of the toxic info), and secondly, cleavage of the spike protein at this site allows affected cells are fused with one another, which disrupts their cell integrity and can lead to massive malfunctions in the vessel walls and lung tissue, for example.

The following unusual properties of the spike protein (which can only be explained by laboratory optimization) were also found by means of sequence analyses

1. The most important docking sites (receptor binding domains) of HIV (Pradhan P, doi:<https://doi.org/10.1101/2020.01.30.927871>)

2. An amino acid sequence corresponding to the neurotoxin of poison snake (king cobra) (Segreto; Environ Chem Lett. 2021;19(4):2743-2757)

3. A heparin binding site, which not only enables the spike protein to open up other cells as host cells for the virus, but above all causes a massive negative effect on blood coagulation (Segreto; Environ Chem Lett. 2021; 19(4): 2743-2757)

4. A region of the spike protein is designed to correspond to the most important region of prions (Alzheimer's, BSE, scrapie) and is believed by the discoverers to be responsible for very rapid acute deaths from Jakob-Kreutzfeld disease (Moret-Chalim C ; DOI:10.13140/RG.2.2.14427.03366)

5. A so-called "superantigen motif", which occurs exclusively in SARS-CoV-2 spike and could trigger the "cytokine storm", which was observed in some very severe courses of the virus infection but also as a result of the "vaccination".

In a synopsis of the information published so far, it can be stated that

that the spike protein of the SARS-CoV-2 virus is obviously adapted for maximum damage in humans - and due to the clear pattern, this can only have happened in the context of laboratory work.

Thus, the spike protein from the Wuhan-1 variant clearly corresponds to a "gain-of-function" product, which is transported 1:1 with the "vaccinations" in humans and must therefore be classified as a biological weapon." (Quote end)

Evidence for the entire preceding lecture:

Expert testimony from Prof. Dr. re. hum. biol. Ulrike Kämmerer, to be loaded via the women's clinic and polyclinic of the University Hospital Würzburg (contact details are already available to the court)

Sources for this:

As Attachment 1:

Copy of the publication "Superantigenic character of an insert unique to SARS-CoV-2 spike supported by skewed TCR repertoire in patients with hyperinflammation" with hyperinflammation) by Mary Hongying Cheng et al.

As attachment 2

Copy of the publication "Attenuation of replication by a 29 nucleotide deletion in SARS-coronavirus acquired during the early stages of human-to-human transmission" early stages of human-to-human transmission) by Doreen Muth et al. "& Christian Drosten".

As Attachment 3:

Copy of the publication "Should we discount the laboratory origin of Covid-19?" (Translated: Should we rule out the origin of Covid-19 in the laboratory?" by Segreto et al.

As Attachment 4:

Copy of the publication "Uncanny similarity of unique inserts in the 2019-nCoV spike protein to HIV-1 gp120 and Gag"

The expert Prof. Dr. As part of her lecture on June 7th, 2022, Kämmerer pointed out in particular that Colonel Prof. Dr. Roman Wölfel, Head of the Institute for Microbiology of the Bundeswehr since October 2019, must also have recognized these connections due to his special qualifications and his above-mentioned work from 2020 as well as his other specialist publications.

She also mentioned that precisely this institute for microbiology, headed by Prof. Wölfel, has the task of identifying biological dangers (biowarfare agents) at an early stage in order to ward off dangers for the soldiers and the population.

The website of the Bundeswehr Institute for Microbiology states, among other things:

"The Bundeswehr Institute for Microbiology is a departmental research facility of the Bundeswehr for **protection against biological warfare** agents and other dangerous infectious diseases. It deals scientifically with a large number of bacteria, viruses and biotoxins, which could also potentially be misused as warfare agents. These are pathogens or toxins that rarely occur in nature, which can usually cause serious, sometimes fatal diseases that can be easily transmitted from person to person and are difficult to treat. **Being**

able to diagnose these quickly and unequivocally is one of the research goals at the Bundeswehr Institute of Microbiology.

The institute has laboratory capabilities that can be deployed quickly anywhere in the world, as well as a diagnostics department accredited according to international standards. The test procedures and laboratory skills set up in Munich offer a wide range of possible applications, including in the diagnosis of natural infections and outbreaks. In January 2020, the Bundeswehr Institute for Microbiology diagnosed the first cases of COVID-19 in people in Germany, isolated the SARS-Coronavirus-2 and decoded its genome..." (end of quote)

Source:

<https://www.instmikrobiobw.de>

Especially in view of his early participation in the diagnosis, isolation and decoding of the SARS-CoV-2 virus, his special expertise and his specialist tasks, it is an incredible process that senior physician Prof. Dr. Roman Wölfel did not vehemently object to the obligation to tolerate Covid-19 injections.

Schmitz
Lawyer