Fabrication: PA 2274 / 4-1

9 messages

Igson Negrin <provereag@gmail.com> Mon, Jun 27, 2022 at 09:51
To: Hüttemann, Kirsten <kirsten.huettemann@dfg.de>

Dear DFG and Kirsten,

This study in the field of virology was funded by the DFG.

https://www.nature.com/articles/s41467-022-28766-y

This work was funded by the Deutsche Forschungsgemeinschaft (DFG, German Research Foundation, grant number PA 2274 / 4-1).

I would like to inform (reinform) you that the fabrication has been carrying out in Germany (all sars-cov-2 studies), probably unintentionally and due to non-implementation and non-documentation of the necessary controls.

Fabrication: What virologists and virology-related scientists call viral genetic material actually comes from healthy human tissue. The genome of any "virus" can be constructed using the same technique that virologists use and using nucleic acids that do not come from supposedly infectious material but from healty human tissue.

Proof:

1. https://telegra.ph/Kontrollexperiment-Phase-1---Die-Widerlegung-der-Virologie-durch-den-cytopathsichen-Effekt-03-10

2. https://telegra.ph/Kontrollexperiment-Phase-2--Entlarvt-Wie-in-der-ma%C3%9Fgeblichen-Studie-zu-SARS-CoV-2-durch-die-chinesischen-Wissenschaftler-getrick-04-03

3. https://telegra.ph/Kontrollexperiment-Phase-3---Strukturelle-Analyse-von-Sequenzdaten--genetische-Untersuchungen-best%C3%A4tigen-Es-gibt-keine-krankmach-04-25

4. https://telegra.ph/Kontrollexperiment-Phase-3---Strukturelle-Analyse-von-Sequenzdaten--genetische-Untersuchungen-best%C3%A4tigen-Es-gibt-keine-krankmach-05-21

Dear DFG, I am going to repeat to you a quick solution to the problem in terms of facing reality quickly. You may now be interested in the quick solution to the problem and due to endangering the health of all citizens in Germany by anti-scientific experiments in virology. Please, ask the scientists and the institutions funded by DFG this (between the stars):

Did you or your colleagues try to extract RNA from uninfected supernatant and cells treated the same way as infected cells and supernatant but virus-free (without patient sample), and to generate the reads and implement" de novo "approach (de novo assembly) or to align (mapping assembly) the reads to sars-cov-2 genome? (extraction of RNA, library preparation, sequencing and assembly of sars-cov-2 genome). Do you have raw data (raw reads) published and documented for the control?

Dear DFG please react urgently! I am going to wait for your urgent response.

Institutions and scientists involved in fabrication and misinterpretation of experimental results:

- Institute of Virology, Freiburg University Medical Center, Faculty of Medicine, University of Freiburg, Freiburg, Germany

- Methodology and Research Infrastructure, Bioinformatics, Robert Koch Institute, Berlin, Germany

- Institute of Experimental and Clinical Pharmacology and Toxicology, Freiburg University Medical Center, Faculty of Medicine, University of Freiburg, Freiburg, Germany

- Hasso Plattner Institute, Digital Engineering Faculty, University of Potsdam, 14482 Potsdam, Germany

- Department of Infectious Disease Epidemiology, Robert Koch Institute, 13353 Berlin, Germany

- Epidemiology of Highly Pathogenic Microorganisms, Robert Koch Institute, 13353 Berlin, Germany

- The sientists: Tamara Kaleta, Lisa Kern, Martin Hölzer, Georg Kochs, Julius Beer, Daniel Schnepf, Martin Schwemmle, Philipp Kolb, Magdalena Huber, Svenja Ulferts, Sebastian Weigang, Alice Wittig, Lena Jaki, Stefan Kröger, Sébastien Calvignac-Spencer , Marcus Panning, Jonas Fuchs.

Best Regards,

Igson Negrin

Igson Negrin <provereag@gmail.com> Tue, Jun 28, 2022 at 11:40
To: Hüttemann, Kirsten <kirsten.huettemann@dfg.de>

Dear DFG and Kirsten,

More info (from the study):

"Whole genome sequencing

cDNA was produced from extracted RNA of oropharyngeal swab or cell culture supernatant samples using random hexamer primers and Superscript III (ThermoFisher) followed by a PCR tiling the entire SARS-CoV-2 genome (ARTIC V3 primer sets"

They should try the following control in order to be able to gain awareness of the fabrication of the alleged sars-cov-2 genome:

extraction of RNA from uninfected supernatant and cells treated the same way as infected cells and supernatant but virus-free (without patient sample), and to generate the reads and implement" de novo "approach (de novo assembly) or to align (mapping assembly) the reads to sars-cov-2 genome. (extraction of RNA, library preparation, sequencing and assembly of sars-cov-2 genome). Raw data (raw reads) should be published and documented for the control using:

A. amplicon-based WGS using specific primers (sars-cov-2 specific)

B. unbiased metagenomic sequencing using random primers (amplification with random hexamers)

Very important for A. and B.:

Try amplification with 12 cycles

Try amplification with 30 cycles

Try amplification with 45 cycles

If you need more info, please tell me.

Best regards,

Igson Negrin
[Quoted text hidden]

Igson Negrin <provereag@gmail.com> Mon, Jul 4, 2022 at 09:04
To: Hüttemann, Kirsten <kirsten.huettemann@dfg.de>

Reminder

On Mon, Jun 27, 2022, 09:51 Igson Negrin <provereag@gmail.com> wrote:
[Quoted text hidden]

Hüttemann, Kirsten <kirsten.huettemann@dfg.de> Mon, Jul 4, 2022 at 09:17
To: Igson Negrin <provereag@gmail.com>

Sehr geehrter Herr Negrin,

vielen Dank für Ihre Hinweise. Eine Kollegin oder ein Kollege aus dem Team Wissenschaftliche Integrität wird Ihre Hinweis prüfen und sich dann an Sie wenden.

Mit freundlichen Grüßen

Kirsten Hüttemann

Dr. Kirsten Hüttemann
Chancengleichheit, Wissenschaftliche Integrität und Verfahrensgestaltung

Deutsche Forschungsgemeinschaft (DFG)
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Kirsten.Huettemann@dfg.de
www.dfg.de

[Quoted text hidden]

Igson Negrin <provereag@gmail.com> Mon, Jul 4, 2022 at 09:21
To: Hüttemann, Kirsten <kirsten.huettemann@dfg.de>

Dear Kirsten

Thank you very much for quick response

OK. I am going to wait.

I will be very happy to share more information with DFG.

Best regards,

Igson
[Quoted text hidden]

Igson Negrin <provereag@gmail.com> Mon, Jul 11, 2022 at 09:15
To: Hüttemann, Kirsten <kirsten.huettemann@dfg.de>

Dear Kirsten,

Could you inform the colleague of your team (You didn't provide me with her/his email)?

Marcus Panning has been contacted! His team did not conduct the necessary control to be able to gain awareness about the fabrication of sars-cov-2 genome from a healthy human tissue (cell culture):

"Thank you for your interest in our article. Up to my knowledge we did not try this but I will check this."

"To get this straight we performed of course cell culture and RT-PCR of our negative controls but in these cases (no detection of viral RNA or positive cell culture) NGS seems to be a little odd."

Dear Kirsten could you (or anyone from DFG) ask Marcus Panning to conduct and documet the neccessary control in detail?

We could speed up this topic and find solution faster (I've been trying since April) because this is an urgent topic - the health of all citizens in Germany.

Dear Kirsten, I will help you and your colleagues from DFG as much as I can.

My colleague from Norway, Daniel Elden, will send the proof (PDF file) of the correspondence with Marcus Panning to you, tomorrow.

Best regards,

Igson
[Quoted text hidden]

Hüttemann, Kirsten <kirsten.huettemann@dfg.de> Mon, Jul 11, 2022 at 09:36
To: Igson Negrin <provereag@gmail.com>

Sehr geehrter Herr Negrin,

ein Kollege aus dem Team Wissenschaftliche Integrität wird sich demnächst an Sie wenden.

[Quoted text hidden]
[Quoted text hidden]

Igson Negrin <provereag@gmail.com> Mon, Jul 11, 2022 at 09:39
To: Hüttemann, Kirsten <kirsten.huettemann@dfg.de>

Dear Kirsten,

OK. Thank you for quick response.

I am going to wait for the response of your colleague

Igson
[Quoted text hidden]

Igson Negrin <provereag@gmail.com> Tue, Jul 19, 2022 at 09:12
To: Hüttemann, Kirsten <kirsten.huettemann@dfg.de>

Dear Kirsten,

There are more useful informations about the fabrication.

The following informations are important in terms of the importance of the control I wrote to you about.

The informations can help DFG to pay attention to PCR-related fabrication in the case of the virology-related scientists funded by DFG (for example Marcus Panning).

I have news (results) from a country and Switzerland.

Scientists from a country did the necessary control, however, with the help of 8 PCR cycles only. They obtained sars-cov-2 sequences. These results have not been published but have been shared with my colleague Daniel Elden. I am sharing this information with the DFG in order to make the DFG informed about the possibility of generating sars-cov-2 sequences without the presence of any virus - Fabrication.

Also, I want to inform you about the control experiments carried out in Swiss laboratories. Scientists in Switzerland managed to generate the whole-genome of sars-cov-2 with the help of nucleic acids from healthy human tissues. With 30 pcr cycles they managed to generate 98% of sars-cov-2 genome with very good coverage. This indicates PCR-related fabrication.

I hope you can share this informations with your colleague from DFG?

If you or your colleague need more informations please tell me.

Best regards,

Igson
[Quoted text hidden]

Your allegation of scientific misconduct against Mr Marcus Panning

7 messages

Ridder, Philip <philip.ridder@dfg.de> Tue, Jul 19, 2022 at 16:08
To: provereag@gmail.com <provereag@gmail.com>

Dear Mr Negrin,

thank you for the allegations of scientific misconduct that you have sent to the German Research Foundation (Deutsche Forschungsgemeinschaft – DFG) since the end of June 2022. As you know, your request has reached the unit for Scientific Integrity at the DFG Head Office and it is currently processed at the DFG. I am a jurist in our unit and I will check your allegation. I will then get back to you and, if necessary, ask for further information.

Your request will be treated confidentially. We ask you to exercise equal confidentiality.

Yours sincerely,

Philip Ridder

Dr. Philip Ridder
Chancengleichheit, Wissenschaftliche Integrität und Verfahrensgestaltung

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philip.ridder@dfg.de
www.dfg.de

Igson Negrin <provereag@gmail.com> Tue, Jul 19, 2022 at 17:16
To: Ridder, Philip <philip.ridder@dfg.de>

Dear Philip,

Thank you for your message

OK.

I understand.

I will be happy to provide more information if neccessary. (There are some new informations sent to your colleague Kirsten)

I am going to wait.

Best regards,

Igson Negrin
[Quoted text hidden]

Igson Negrin <provereag@gmail.com> Fri, Jul 22, 2022 at 14:38
To: Ridder, Philip <philip.ridder@dfg.de>

Dear Philip,

I am sorry to bother you.

I am sending you examples of necessary controls in virology in pdf format in german language.

As you know, Marcus Panning confirmed that one of the necessary controls proving fabrication of the alleged sars-cov-2 genome was not carried out.

Best regards,

Igson Negrin

On Tue, Jul 19, 2022, 16:08 Ridder, Philip <philip.ridder@dfg.de> wrote:
[Quoted text hidden]

<postmaster@dfg.de> Fri, Jul 22, 2022 at 14:42
To: provereag@gmail.com

Ihre Nachricht wurde aufgrund ihrer Größe an keinen Benutzer zugestellt. Der Grenzwert liegt bei 22 MB. Die Größe Ihrer Nachricht beträgt 25 MB.

Ridder, Philip (philip.ridder@dfg.de)

Ihre Nachricht konnte nicht gesendet werden, weil sie zu groß ist.

Diagnoseinformationen für Administratoren:

I am sorry to bother you.

I am sending you examples of necessary controls in virology in pdf format in german language.

As you know, Marcus Panning confirmed that one of the necessary controls proving fabrication of the alleged sars-cov-2 genome was not carried out.

Best regards,

Igson Negrin

On Tue, Jul 19, 2022, 16:08 Ridder, Philip <philip.ridder@dfg.de> wrote:
[Quoted text hidden]

Ridder, Philip <philip.ridder@dfg.de> Mon, Jul 25, 2022 at 10:16
To: Igson Negrin <provereag@gmail.com>

Dear Mr Negrin,

thank you for providing additional information. We will consider it in the further process.

Yours sincerely,

Philip Ridder

Dr. Philip Ridder
Chancengleichheit, Wissenschaftliche Integrität und Verfahrensgestaltung

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[Quoted text hidden]

Igson Negrin <provereag@gmail.com> Wed, Aug 10, 2022 at 14:14
To: Ridder, Philip <philip.ridder@dfg.de>

Dear Philip,

Is there any progress in terms of implementing and documenting the controls? (Marcus Panning)

I read rules of DFG. DFG can contact an expert in the field of virology. However, no expert has ever carried out and documented the necessary controls. There is no such publication.

There is about 278000 publications (LitCovid Data) and no one tried to conduct and document the controls.

Kind regards,

Igson Negrin

[Quoted text hidden]

Von: Ridder, Philip
Gesendet: Donnerstag, 18. August 2022 16:46
An: 'Igson Negrin' <provereag@gmail.com>
Betreff: Confidential: Your allegation of scientific misconduct against Prof Marcus Panning
Vertraulichkeit: Vertraulich

Dear Mr. Negrin,

thank you for your messages starting on 25 May 2022, which have reached the DFG Head Office unit for Research Integrity. We handle your request confidentially. We would like to address your allegation against Professor Panning (University of Freiburg) today.

First of all, we would like to inform you that our activity in the investigation of cases of alleged scientific misconduct is based on codified Rules of Procedure (hereinafter referred to as "Rules"), which we attach to this e-mail (in English and German language). The Rules outline the course of proceedings, but they also determine what constitutes scientific misconduct which the DFG is capable to investigate – and what does not. You can find additional information about the proceedings on the DFG website.

The DFG takes allegations of potential scientific misconduct very seriously. However, we can only initiate an investigation if there is a sufficient and substantial suspicion of a scientific misconduct in the field of the DFG's funding activity. The reported (potential) misconduct needs to fulfill one of the matters contained in section II of our Rules. Not all questionable research practices or violations of good scientific practice can be prosecuted by the DFG.

So far, we do not see that the behavior you have reported constitutes a sufficient falsification or fabrication of data. Technical mistakes, such as leaving out control experiments which would, from a scientific point of view, normally be expected from researchers in this area, do not for themselves constitute a falsification or fabrication of data according to our Rules of Procedure.

Therefore, the information at hand does not justify opening an investigation.

Yours sincerely,

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www.dfg.de

Igson Negrin <provereag@gmail.com> Sat, Aug 20, 2022 at 12:06
To: Ridder, Philip <philip.ridder@dfg.de>

Dear Philip,

Thank you for the explanation.

I do not think that the absence of necessary and logical controls constitutes a fabrication.

The experiment setup used by Marcus Panning and all other virologists generates artifacts. Genomes in virology do not exist in reality but represent mental and software constructs (invention, fabrication).

I would not like to repeat all the information I have already sent to you.

I would like to remonstrate. There are new facts:

Whole "genome" fabrication – sars-cov-2

Marcus Panning reports the existence of alleged whole-genome sequencing and documents alleged whole-genome sequencing in the publication. However, in reality, the team of Marcus Panning has never conducted such experiments. The alleged whole genome of sars-cov-2 never appears in reality as a whole.

Professor Panning always use whole-construct assembly (without neccessary controls). Also, it should not be forgotten that the consensus sequences presented to the public by the professor Panning are not a reality:

"It is surprising, then, that scientists forget this and treat the consensus as reality" - Consensus Sequence Zen; Thomas D. Schneider.

This (alleged whole-genome sequencing of sars-cov-2) is a typical example for research misconduct. This is fabrication.

A nucleic acid is a single biomacromolecule. This biomolecule is made up of nucleotides. Nucleotides are interconnected by covalent bonds. The first nucleotide is connected to the second, the second to the third, the third to the fourth, etc. When sequencing an entire

(whole) biomacromolecule, the first nucleotide is detected first, then the second nucleotide, then the third, then the fourth, and so on.

Marcus Panning did not sequence the whole molecule of 29903 nucleotides but only short PCR products up to 400 nucleotides in length (the lenght of the reads is about 150bp). These PCR products are called amplicons.

Amplicons are fabricated

Fabricated amplicons do not exist in reality. Thanks to the anti-scientific use of PCR technology, sequences that do not exist in the starting material are obtained. (How to investigate this? With controls.)

Of particular concern is that no scientific publication regarding the sars-cov-2 construct in the world documents gel electrophoresis of the alleged whole genome. There are electrophoregrams of fabricated amplicons, but there are no electrophoregrams of the "whole genomes". It should not be forgotten that gel electrophoresis is the gold standard for determining the length of a genome.

Fabrication of isolation of "viral" particles and biomolecules

Experimental data are fabricated by reporting experiments that were never conducted. Marcus Panning but also other organizations and institutions around the world are involved in misleading the public regarding the existence of isolation, purification and biochemical characterization of alleged specific biomolecules and alleged specific particles built by these biomolecules.

Regardless of the use of anti-scientific terms such as "isolation in cell cultures" and “viral isolate in cell culture supernatant”, but also the use of microscopy, virology-related scientists have a scientific obligation to examine alleged specific particles and alleged specific biomolecules according to the rules of experimental biochemistry and molecular biology. For example: in science, isolation of small biochemically uncharacterized particles is performed with density gradient centrifugation (rate zonal and isopycnic). Detailed illustration (without controls) is available here: https://youtu.be/Yc7CLOviOPk

However, in virology, these basic rules of scientific work are always violated. In virology, nucleic acids are not extracted from "viruses" but from a mixed-sample containing mixed nucleic acids orginating from bovine (FBS), human, human microbiome, animal, cancer, bacteria, fungi and the like. It should not be forgotten that no media is free from bovine RNA (publication: Extracellular small non-coding RNA contaminants in fetal bovine serum and serum-free media).

Virologists most commonly use VTM, FBS, and cell culture media. In this mixed sample, it is not possible to determine the origin of nucleic acids, and for that reason, bioinformatics use reference constructs called templates for alignment.

However, these templates were also created from a mixed sample, which makes the use of templates and processes such as mapping-assembly and de novo assembly meaningless in virology.

Conclusion for "Human Virology":

1. There is no sars-cov-2 whole genome in reality.
2. There is no "viral" particle in reality.
3. There is no "viral" biomolecule in reality.
4. There is no sars-cov-2 in reality.
5. There is no Covid-19 in reality.
6. Sars-cov-2 is complete artefact because it is fabricated.

If you need more informations please tell me.

Kind regards,

Igson Negrin
[Quoted text hidden]

Igson Negrin <provereag@gmail.com> Tue, Aug 23, 2022 at 10:23
To: Ridder, Philip <philip.ridder@dfg.de>

Reminder
[Quoted text hidden]

Igson Negrin <provereag@gmail.com> Tue, Aug 23, 2022 at 12:49
To: Ridder, Philip <philip.ridder@dfg.de>

Dear Philip,

Sorry to bother you again.

I hope you will take the information I have provided to you very seriously.

It would be very useful to inform the president of the DFG (Professor Dr. Katja Becker). She is a member of the Interdisciplinary Commission for Pandemic Research. She has an obligation to identify research gaps and to close them.

There is a major gap in the alleged sars-cov-2 pandemic: the absence of the necessary control experiments. These controls prove in detail: sars-cov-2 is a complete artifact of experiment setup. It is fabricated. It doesn't exist in reality.

All informations are available. I hope someone from DFG in interested for the controls. I hope DFG will not ignore the controls.

Can you inform Professor Katja Becker about the mentioned gaps in the research?

Kind regards,

Your allegation of scientific misconduct against Prof Marcus Panning

8 messages

Ridder, Philip <philip.ridder@dfg.de> Thu, Aug 25, 2022 at 13:35
To: Igson Negrin <provereag@gmail.com>

Dear Mr. Negrin,

our assessment of the allegations at hand remains unchanged.

A formal right to “remonstrate” does not apply in this case as no proceedings have been initiated which could be discontinued. We have not opened an investigation for the following reasons (see below):

“So far, we do not see that the behavior you have reported constitutes a sufficient falsification or fabrication of data. Technical mistakes, such as leaving out control experiments which would, from a scientific point of view, normally be expected from researchers in this area, do not for themselves constitute a falsification or fabrication of data according to our Rules of Procedure.”

Yours sincerely,

Philip Ridder

Dr. Philip Ridder
Chancengleichheit, Wissenschaftliche Integrität und Verfahrensgestaltung

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www.dfg.de

Igson Negrin <provereag@gmail.com> Thu, Aug 25, 2022 at 13:41
To: Ridder, Philip <philip.ridder@dfg.de>

Dear Philip,

Thank you for quick response

You didn't answer me about president of DFG: Katja Becker

Could you contact professor Katja or could you provide me with her email or email of the comission?

Thank you for your help

Kind regards,

Igson Negrin
[Quoted text hidden]

Igson Negrin <provereag@gmail.com> Thu, Aug 25, 2022 at 17:18
To: Ridder, Philip <philip.ridder@dfg.de>

Dear Philip,

“So far, we do not see that the behavior you have reported constitutes a sufficient falsification or fabrication of data. Technical mistakes, such as leaving out control experiments which would, from a scientific point of view, normally be expected from researchers in this area, do not for themselves constitute a falsification or fabrication of data according to our Rules of Procedure.”

Could your team check my new informations (not in the form of remonstration)? Does this constitute enough fabrication so you could be able to open investigation:

I do not think that the absence of necessary and logical controls constitutes a fabrication.

The experiment setup used by Marcus Panning and all other virologists generates artifacts. Genomes in virology do not exist in reality but represent mental and software constructs (invention, fabrication).

Whole "genome" fabrication – sars-cov-2

Marcus Panning reports the existence of alleged whole-genome sequencing and documents alleged whole-genome sequencing in the publication. However, in reality, the team of Marcus Panning has never conducted such experiments. The alleged whole genome of sars-cov-2 never appears in reality as a whole.

Professor Panning always use whole-construct assembly (without neccessary controls). Also, it should not be forgotten that the consensus sequences presented to the public by the professor Panning are not a reality:

"It is surprising, then, that scientists forget this and treat the consensus as reality" - Consensus Sequence Zen; Thomas D. Schneider.

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Of particular concern is that no scientific publication regarding the sars-cov-2 construct in the world documents gel electrophoresis of the alleged whole genome. There are electrophoregrams of fabricated amplicons, but there are no electrophoregrams of the "whole genomes". It should not be forgotten that gel electrophoresis is the gold standard for determining the length of a genome.

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Experimental data are fabricated by reporting experiments that were never conducted. Marcus Panning but also other organizations and institutions around the world are involved in misleading the public regarding the existence of isolation, purification and biochemical characterization of alleged specific biomolecules and alleged specific particles built by these biomolecules.

Regardless of the use of anti-scientific terms such as "isolation in cell cultures" and “viral isolate in cell culture supernatant”, but also the use of microscopy, virology-related scientists have a scientific obligation to examine alleged specific particles and alleged specific biomolecules according to the rules of experimental biochemistry and molecular biology. For example: in science, isolation of small biochemically uncharacterized particles is performed with density gradient centrifugation (rate zonal and isopycnic). Detailed illustration (without controls) is available here: https://youtu.be/Yc7CLOviOPk

However, in virology, these basic rules of scientific work are always violated. In virology, nucleic acids are not extracted from "viruses" but from a mixed-sample containing mixed nucleic acids orginating from bovine (FBS), human, human microbiome, animal, cancer, bacteria, fungi and the like. It should not be forgotten that no media is free from bovine RNA (publication: Extracellular small non-coding RNA contaminants in fetal bovine serum and serum-free media).

Virologists most commonly use VTM, FBS, and cell culture media. In this mixed sample, it is not possible to determine the origin of nucleic acids, and for that reason, bioinformatics use reference constructs called templates for alignment.

However, these templates were also created from a mixed sample, which makes the use of templates and processes such as mapping-assembly and de novo assembly meaningless in virology.

Conclusion for "Human Virology":

1. There is no sars-cov-2 whole genome in reality.
2. There is no "viral" particle in reality.
3. There is no "viral" biomolecule in reality.
4. There is no sars-cov-2 in reality.
5. There is no Covid-19 in reality.
6. Sars-cov-2 is complete artefact because it is fabricated.

If you need more informations please tell me.

Kind regards,

Igson Negrin

Fabrication: more info

5 messages

Igson Negrin <provereag@gmail.com> Sun, Aug 28, 2022 at 12:56
To: wi@dfg.de

Dear Research Integrity Team (Ms. Liane Badiane, Dr. Kirsten Hüttemann, Ms. Lydia Llaga, Dr. Philip Ridder, Mr. Martin Steinberger),

My point is not that the absence of necessary and logical never-conducted and documented control experiments constitutes fabrication. I contacted you because the experiment setup used by Professor Marcus Panning generates constructs (invention, fabrication) that do not exist in reality. In science this can only be investigated through control experiments.

"Negative controls in epidemiological studies are analogous to negative controls in laboratory experiments, in which investigators test for problems with the experimental method by leaving out an essential ingredient..."

https://www.ncbi.nlm.nih.gov/pmc/articles/PMC5428075/

"The second general approach is to perform negative controls: to repeat the experiment under conditions in which it is expected to produce a null result and verify that it does indeed produce a null result. Several strategies are employed to design negative controls, such as:

• Leave out an essential ingredient."

https://www.ncbi.nlm.nih.gov/pmc/articles/PMC3053408/

All virologists in the world and professors Marcus Panning have never tried to leave out patient sample and to conduct and document experiment from start to finish (the finish is Next Generation Sequencing and Whole "Genome" assembly).

In short:

1. Marcus Panning and all virologists (human virology) fabricate the alleged sars-cov-2 by experiment setup. Sars-cov-2 is an artifact of the experiment setup used by Marcus Panning. Sars-cov-2 doesn't exist in reality.

2. The fabrication used by Marcus Panning can only be investigated through control experiments as I have explained in previous messages. These controls have already been carried out as I informed you, and the result is: sars-cov-2 does not exists.

In the previous message I have shared new informations so you could start with investigation.

Kind regards,

Igson Negrin

Igson Negrin <provereag@gmail.com> Wed, Aug 31, 2022 at 11:55
To: wi@dfg.de

Dear Research Integrity Team,

I will contact you on September 5 if there is no response by then.

I hope that you can take my new messages into account for the start of the investigation.

Kind regards,

Igson Negrin
[Quoted text hidden]

Igson Negrin <provereag@gmail.com> Mon, Sep 5, 2022 at 10:06
To: wi@dfg.de

Reminder

On Sun, Aug 28, 2022, 12:56 Igson Negrin <provereag@gmail.com> wrote:
[Quoted text hidden]

More info about fabrication

4 messages

Igson Negrin <provereag@gmail.com> Mon, Sep 5, 2022 at 11:28
To: Hüttemann, Kirsten <kirsten.huettemann@dfg.de>

Dear Kirsten,

I am sharing with you additional information so that you could begin your investigation. A lot of fabrication and manipulation was applied by Professor Marcus Panning.

"The raw reads were pre-processed with fastp v.0.20.173 and mapped to the SARS-CoV-2 Wuhan-Hu-1 reference genome (Genbank: NC_045512) using BWA-MEM v.0.7.1774"

https://www.ncbi.nlm.nih.gov/pmc/articles/PMC8894356/

Professor Marcus Panning is involved in a severe manipulation and fabrication due to use of construct NC_045512 whose invention is based on severe manipulation.

Manipulation: Using raw reads of NC_045512, it is not possible to reproduce the longest contig (30,474 nucleotides).

More info: https://www.ncbi.nlm.nih.gov/pmc/articles/PMC5428075/

You can also read new messages sent to your team about fabrication conducted by professor Marcus Panning and his team.

I am waiting for a response from you or your team.

Kind regards,

Igson Negrin

Igson Negrin <provereag@gmail.com> Thu, Sep 8, 2022 at 10:46
To: wi@dfg.de

Reminder 2

On Sun, Aug 28, 2022, 12:56 Igson Negrin <provereag@gmail.com> wrote:
[Quoted text hidden]

Igson Negrin <provereag@gmail.com> Mon, Sep 12, 2022 at 10:04
To: Hüttemann, Kirsten <kirsten.huettemann@dfg.de>

Dear Kirsten,

I hope you or your team were able to check (maybe with the help of a bioinformatician) the construct NC_045512 used by Marcus Panning (and all virologist in the world)?

It would be very important.

Marcus Panning used https://usegalaxy.eu/ in his study. On the website you can get the result of the manipulation in a few hours: the longest contig is 29802 and not 30474. It is not possible to reproduce the longest contig because of severe manipulation used for the generation of the construct NC_045512. The construct is used by Marcus Panning.

I will send you a tutorial with the help of which it is possible to understand the manipulation with just a few clicks (even a child can click several times like in a game).

I will be happy to provide you with more informations.

Alleged "Raw" reads of the construct MN908947 (NC_045512) are available here:

https://www.ncbi.nlm.nih.gov/bioproject/?term=PRJNA603194

Kind regards,

Igson Negrin
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Hüttemann, Kirsten <kirsten.huettemann@dfg.de> Mon, Sep 12, 2022 at 10:42
To: Igson Negrin <provereag@gmail.com>
Cc: Ridder, Philip <philip.ridder@dfg.de>

Sehr geehrter Herr Negrin,

mein Kollege, Herr Dr. Ridder, hat die Bearbeitung Ihres Anliegens übernommen. Ich werde daher Ihre Mail an meinen Kollegen weiterleiten und darf Sie bitten, den weiteren Austausch in dieser Angelegenheit direkt mit Herrn Ridder vorzunehmen.

Mit freundlichen Grüßen

Kirsten Hüttemann

Dr. Kirsten Hüttemann
Chancengleichheit, Wissenschaftliche Integrität und Verfahrensgestaltung

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Igson Negrin <provereag@gmail.com> Sun, Jan 22, 2023 at 08:52
To: wi@dfg.de

Reminder 3 (the last reminder)

On Sun, Aug 28, 2022, 12:56 Igson Negrin <provereag@gmail.com> wrote:
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