

Twitter Thread by Coroldo1



Coroldo1

[@coroldo1](#)



#MN908947 #2019_nCov #GISAID #CHINACDC

GISAID Submission date dispute■

<https://t.co/yx666wb6dO>

Yong-Zhen Zhang's Genbank submission

[#CNIC](#) [#SARSCOV2](#) [#GISAID](#) [#CHINACDC](#) [#ISL402119](#) [#ISL402120](#) [#ISL402121](#)

Thread for evidence only,timestamp according to GMT+8, the timezone of China, where first 4 SARS-COV-2 genomes released to the world. Any further analysis subject to individual's discretion. So, Let's roll

— Coroldo1 (@coroldo1) [February 26, 2021](#)

[@R_H_Ebright](#) [@seeshawn](#) [@strom_m](#) [@edwardcholmes](#) [@arambaut](#) [@Ayjchan](#) This dispute on 'first open' is getting 'better'. GISAID formally wrote to authors who acknowledged Virological post as world's first release of 2019ncov, penalised authors of MN908947, didn't ID who..... 1/3

Supposedly from Fudan or UoS, ACCESSED and KNEW IVDC's 'initial' submission of IVDC-01/05 on GISAID on 10th Jan, 2020, via login records of his/her/theirs GISAID account credentials. Hence as a point to point rebuttal to 'pressure' & 'urge' to release the fastaqc file... 2/3

on 11th Jan(initiated very late on 10th Jan UTC) according to the well known narratives in global academic society and Nature/TIME stories. 3/3

<https://t.co/qJM9AkYzdj>

Imagining our life without covid-19 in 2020 for a while, then start to read

"Infectious diseases are like wildfires. You need to stomp them out when they are containable. In both cases, time is of the essence." – Anne Rimoin

Though many experts had been predicting a pandemic and worked with efforts and tactics in response for years before covid-19, here we all are, inevitably, entering the age of After Corona. Under the pressure of Covid-19 pandemic, outburst of human potential united scientists, doctors and nurses, biotech/pharmaceutical companies to developed vaccines in unprecedented speed. Did we cut some corners? Jumped into phase III first, used saline rather than vaccine matrix as control, started human trial to earlier without fully safety review, yes. But with the help of accumulated technology and manufacturing platforms, 'tolerance' from supervision, vaccine roll-out is on the track and we are closing our race against covid-19.

Are we going back to the routine and sometimes 'boring' life in Before Corona age, probably never will. The best case scenario would be stuck with Covid-19 for years to come if we couldn't catch up the upcoming variants. So isn't it justified to say we were not well prepared for pandemics, and will probably never be well prepared for potential pandemics? Some experts might argue by enhancing pathogen surveillance, creating innovative vaccine platforms, initiating novel global cooperation strategy, when in place, would help to put us in a better position when facing next pandemic. Hardly disagree, but at what cost? How to define "well", to define "better". There is no doubt that everyone's life was affected in certain degree and response measures were at the expense of freedom and sacrifice, mostly, in ordinary people.

Isn't it a plausible and logical angle to look into the details of this pandemic and seeking experience and lessons for our future preparation, at the lowest possible cost to the people? Thus back to the opening of this thread, quicker response.

This brings us to the discussion of the monumental actions of genome release. The importance and influence of the first genome release evolved during the course of pandemic. In the earlier days, it helps to bring the global wisdom into the virus research to identify it as 'potential' pandemic strain, and no doubt help for the ID and sequencing, as Ian Mackay commented immediately after release, "Start your engines assay designers!". With the progress of pandemic and encouraging news of vaccine trial, the date and person of the first release serve additional role as the beginning of global fight against virus, though we all understand it's the tireless efforts of numerous scientists after that day who contribute to our current 'little but huge' success.

Well, here comes that But. Though in the beginning some scientists questioned the possible embargo date, none of the authors came to clarify the doubt and Genbank was taking the blame. Here quoted again from Ian Mackay simply because his post was closeby, "Genbank is not helpful for rapid response. Or any timely sequence release in my lengthy experience." All this would probably quietly slide away(not quoting Milley Cyrus) even no clarification in any of the occasions genome authors publicly commented on the release, and Genbank records obtained via FOIA. We all had better things to do except been brought into attention by the announcement from GISAID on their version of the genome release. GISAID pointed out the embargo thus the MN908947 had not yet been released at the time of the virological post though did not indicate the embargo date was set on 20200712. And, credentials of some of the persons listed as authors of Wuhan-hu-1 were among the access credentials used on Jan 10, 2020 to access the first genomes available on GISAID.

Such a dispute of the first genomes release carries out away even after one year into the covid-19, the details will provide valuable information and lessons to our next preparation, how to do it quicker, or avoid potential delay, share of the crucial information, hence we can 'constrain' the outbreak in relatively smaller scale. In terms of peer review confidentiality, in this particular case, in this particular scale of pandemic, clarity clearly weighs more.

All reasonable people understand that in terms of decision making, it should only be weighted in its unique circumstance. And reasoning the decision making isn't like piece together a 2D jigsaw puzzle or simply connecting the dots, it contains multiple parameters and act as whole. The post on the Virological.org marks the start of global cooperation without question, and can we learn to do quicker if we could find space for micro-improvements for next time? That's what people are doing now.

To be clear, the below was NOT a secret, to my■■■ colleagues since 2020. Speak on behalf of most of us, the reason of silence varied: occupied hours fighting against ■; didn't want to engage in the ■ twisted power play between various parties; been there, done that, didn't care.

our reason to post here, still varied: ridiculous fictions by various influential agency and person(we wish you all have the dignity to retract your stories and books); outrageous suppression by private database to acknowledge the fact; Urgency for global data sharing consensus;

To remove the tumors of public health system; most important of all, the magnitude of COIVD made us thinking, what went wrong, again?

To be crystal clear AGAIN, this pandemic demonstrated the failure of our public health responses in every country, no exception. The future way out is to make ourselves better, stronger, together, and have a clear weight in mind of general people's well being over personal gains.

Spare us with your excuses such as grand interest, economics.....and particularly 'science without politics', if we rephrase politics as compromise or quid-pro-quo, they are as common in our research as your daily pint.....

COI: worked with SARS,HIV,EV71,H1N1,Ebola,MERS,H7N9 etc outbreaks in CDC.

And calling for authors of first genomes, including MN908947 to publicly disclose their COI with companies. Welcome any challenge against the authenticity of any information posted here.

For convenience, the timestamp here were all GMT+8 for your own analysis, though it involved communications in timezone of United States, Singapore.

Decision regarding the genomes submission in IVDC (Other two institutes also preparing), date 2020-1-12 GMT

病毒所拟投GISAID的3个序列



maxj2004 <maxj2004@aliyun.com>
2020/1/12 1:47

收件人: 谭文杰; Dayan Wang; 王大燕; 赵翔 抄送: wenbo_xu1@aliyun.com; 武桂珍

[保存所有附件](#)



wh004_G2_consensus_29897...
22.37 KB



wh005_G3_consensus_29788...
29.85 KB



wh008_G3_consensus_29849...
29.91 KB

各位领导 and 专家, 这是按照高主任的要求和许所的建议, 中心实验室提供的 3 个基因组序列 (含样品来源, 测序方法和长度) 其中样品 004 就是 3 个实验室 (中心, 流感和应急中心) 最后核实的病毒所最终序列。妥否, 请指正。 马

Xuejun Ma Ph.D., Professor,
Director, Department of Core Facility,
PI, Key Laboratory of Medical Virology, Ministry of Health,
National Institute for Viral Disease Control and Prevention,
Chinese Center for Disease Control and Prevention,
Changbai Rd 155, Changping district, Beijing, 102206, China.
Tel.: +86 10 58900810; Fax: +86 10 58900810

PS: As yahoo China ceases the mail service, my previous mailbox (maxj2004@yahoo.com.cn) will be invalid on Dec.30, 2013. Please email to my current mailbox (maxj2004@aliyun.com), Thanks.

因雅虎中国停止服务, 我以前的电子邮箱 (maxj2004@yahoo.com.cn) 将于 2013 年 12 月 30 日无效, 以后的电子邮件请发送至我现在的邮箱 (maxj2004@aliyun.com) 谢谢。

Submission to GISAID

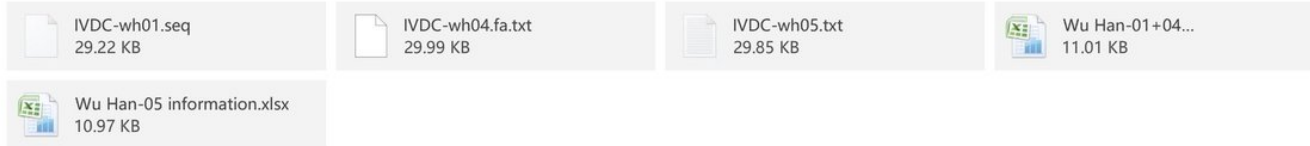
Three CoV Sequences from IVDC, China CDC



Wang Dayan <wangdayan@ivdc.chinacdc.cn>
2020/1/12 4:29

收件人: Peter Bogner 抄送: tanwj28@163.com; ZX; wenbo_xu1@aliyun.com; wugz@ivdc.chinacdc.cn; wangwl@ivdc.chinacdc.cn; maxj2004@aliyun.com; George Fu Gao (高福)

保存所有附件



Dear Peter,

Please find attached three sequences and related information files. Please be noted that sequence 01 and 04 are in one Excel file, and 05 in a separated one. The differences are labeled in Red, for your convenience.

Please help us to register (Could please let me know when registration is done) and please help upload to GISAID. Please hold it until we tell you to release (we were told we need to release with two other labs at the same time).

You may need below information for registration:

Name: Wenjie Tan

Phone: 8610-58900878

Email: tanwj28@163.com

Institute: National Institute for Viral Disease Control and Prevention, China CDC

Address: 155 Changbai Road, Changping District, Beijing 102206 China

If you need more information, please feel free to let us know.

The age and the sex of the patients are not fully confirmed now, so please just leave it blank and will add when ready.

Best regards,

Dayan

Here was referring to A*STAR in Singapore.

Re: Three CoV Sequences from IVDC, China CDC



Peter Bogner <peter@gisaid.org>
2020/1/12 9:44

收件人: Wang Dayan 抄送: tanwj28@163.com; ZX; wenbo_xu1@aliyun.com; wugz@ivdc.chinacdc.cn; wangwl@ivdc.chinacdc.cn; maxj2004@aliyun.com; George Fu Gao (高福)

Brilliant

We forward these updates to our team and trust they too will get up momentarily. We plan on writing to all of you, once all three lab data are uploaded for a final review, before you and your colleagues give us the green light to make public the data of all 3 labs at the same time.

We are still very keen to receive an image of the virus from an electron microscope if it can be arranged. It could really help China's standing and have a wider media effect if it come from China.

Xie Xie

Homepage text drafted by IVDC

China releases genetic sequence of newly discovered coronavirus from Wuhan

The Chinese Center for Disease Control and Prevention, Chinese Academy of Science and Chinese Academy of Medical Science released the genome sequence of a newly discovered coronavirus. The virus was identified during an outbreak in the city of Wuhan, where patients were suffering from respiratory illnesses such as pneumonia since December 2019.

The genome sequence of this betacoronavirus is crucial to develop specific diagnostic tests and to identify potential intervention options.

Viruses image submission by IVDC

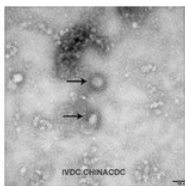
Re: Text for GISAID Homepage



Wang Dayan <dayanwang@cnic.org.cn>

2020/1/12 11:17

收件人: peter@gisaid.org 抄送: tanwj28@163.com; ZX; wenbo_xu1@aliyun.com; wugz@ivdc.chinacdc.cn; wangwl@ivdc.chinacdc.cn; maxj2004@aliyun.com; George Fu Gao (高福)



Dear Peter,

As attached! Please embedded with "by China CDC"

Thank you!

Best regards,

Dayan

Now, 'proof reading' against any existing announcements, stories, legal letters etc al.

<https://t.co/q5eBeJn3V7>

h/t archived by [@Biol4Ever](#)