

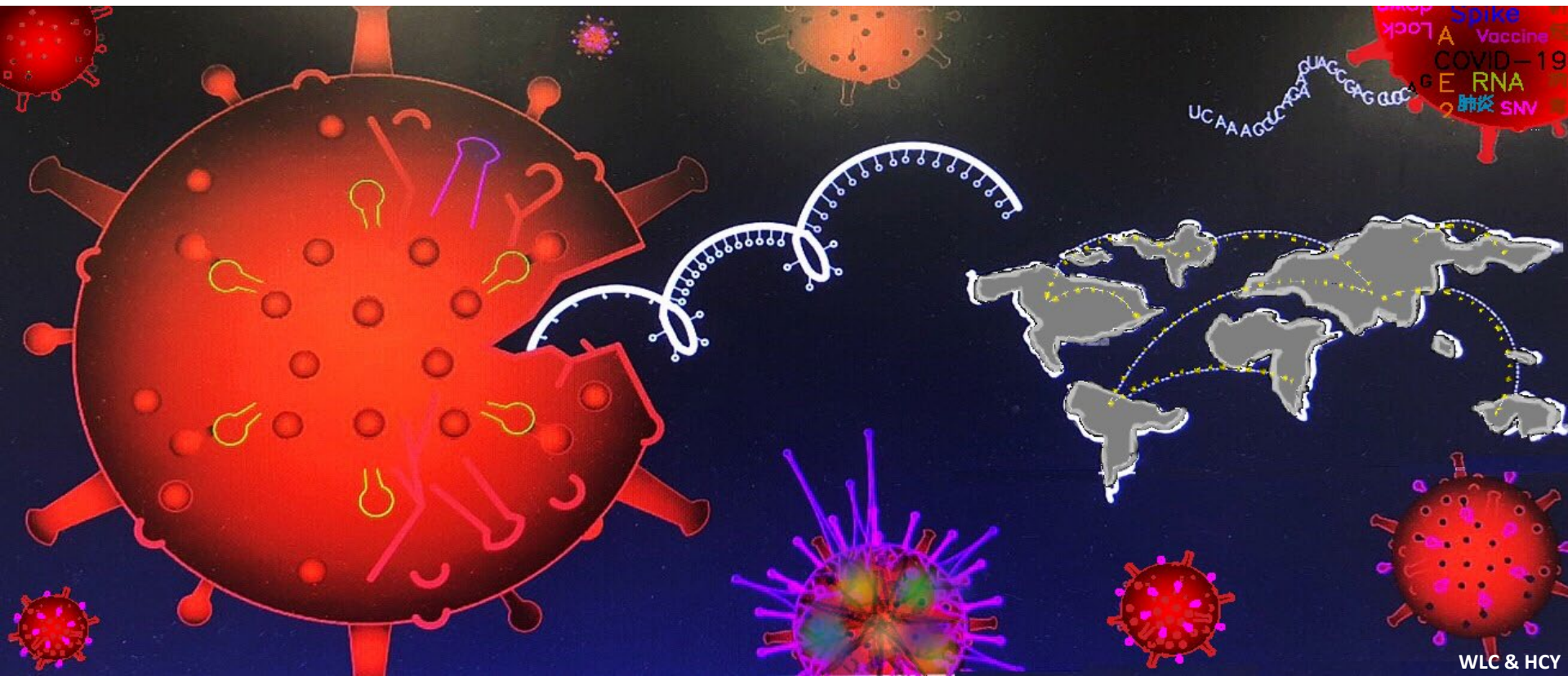
Lessons learned from the global COVID-19 pandemic

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2023-07-18



WLC & HCY

COVID-19: Coronavirus disease 2019

- Dec 2019: the **first case of the human-to-human transmission** of a novel coronavirus was reported in Wuhan, China.
- Feb 2020: **World Health Organization (WHO)** gave the disease a name, “**Coronavirus disease 2019**” (i.e., COVID-19).
- Feb 2020, the **International Committee on Taxonomy of Viruses (ICTV)** named the causal agent for COVID-19, “Severe Acute Respiratory Syndrome Coronavirus 2” (i.e., SARS-CoV-2).
- SARS-CoV-2 (Beta genera) as the **7th human coronavirus**: Other six are: **OC43 (Beta, 1960-1970)**, **229E (Alpha, 1960-1970)**, **SARS-CoV (Beta, 2003)**, **NL63 (Alpha, 2004)**, **HKU1 (Beta, 2005)**, and **MERS-CoV (Beta, 2012, 2015, 2018)**. Green color: common colds.

Symptoms of COVID-19

- **Incubation period: 2 to 14 days** after exposure
- **Pre-symptomatic transmission** (“before symptoms”)
- **Common symptoms** are similar to other **colds and flu**:
 - ✓ **Fever**
 - ✓ **Cough**
 - ✓ **Tiredness**
 - ✓ **Loss of taste or smell**
 - ✓ **Headache**
 - ✓ **Sore throat**
 - ✓ **Shortness of breath**

Statistics of COVID-19 Infection and Deaths



Search by Country, Territory, or Area



[Overview](#)

Measures

Table View

Data

More Resources

WHO Coronavirus (COVID-19) Dashboard

Back to top

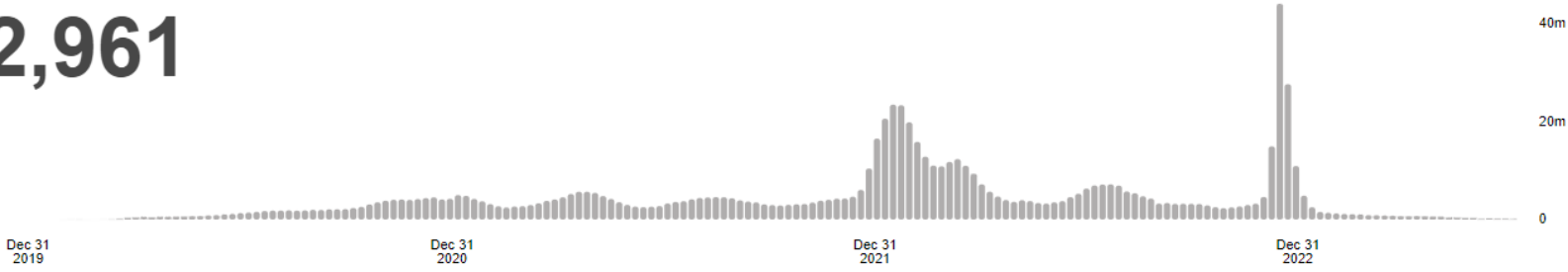
Globally, as of **12:14pm CEST, 12 July 2023**, there have been **767,972,961 confirmed cases** of COVID-19, including **6,950,655 deaths**, reported to WHO. As of **8 July 2023**, a total of **13,474,185,140 vaccine doses** have been administered.

Global Situation

Daily Weekly

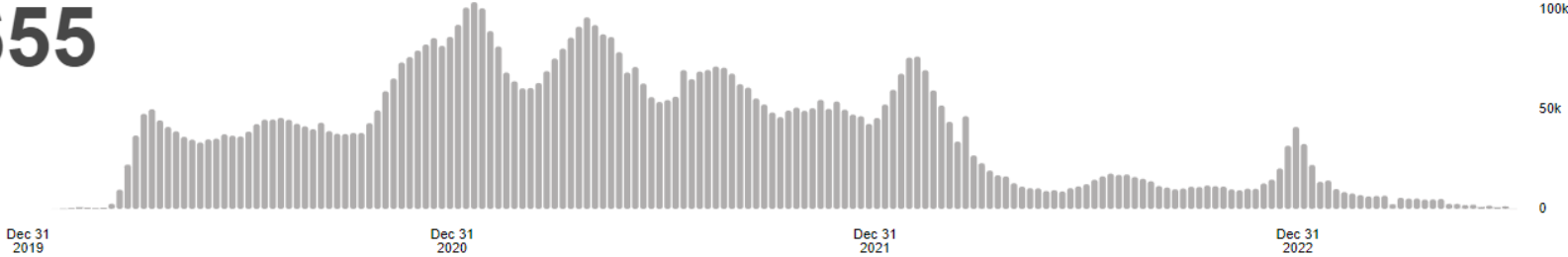
767,972,961

confirmed cases



6,950,655

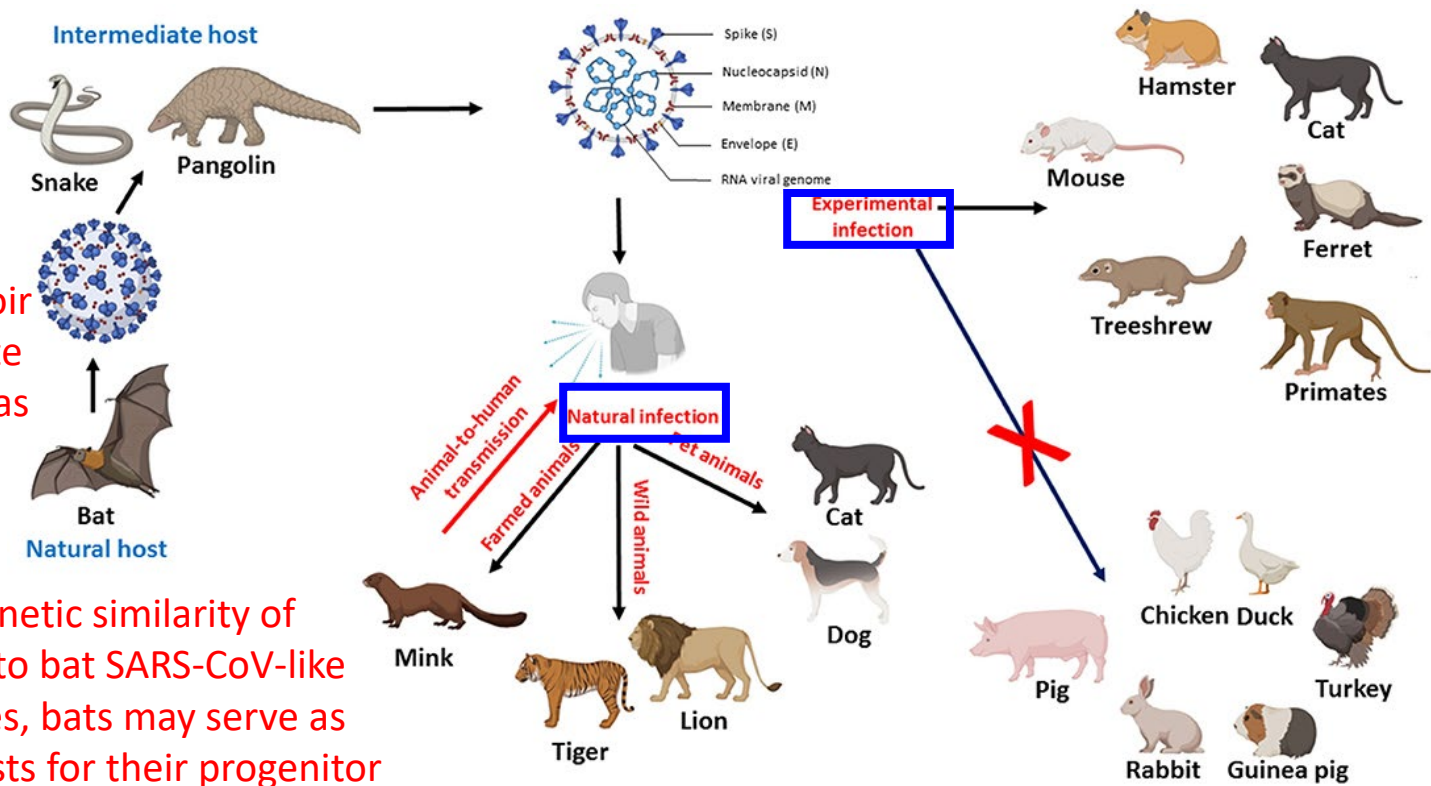
deaths



Source: World Health Organization
Data may be incomplete for the current day or week.

Origin of SARS-CoV-2

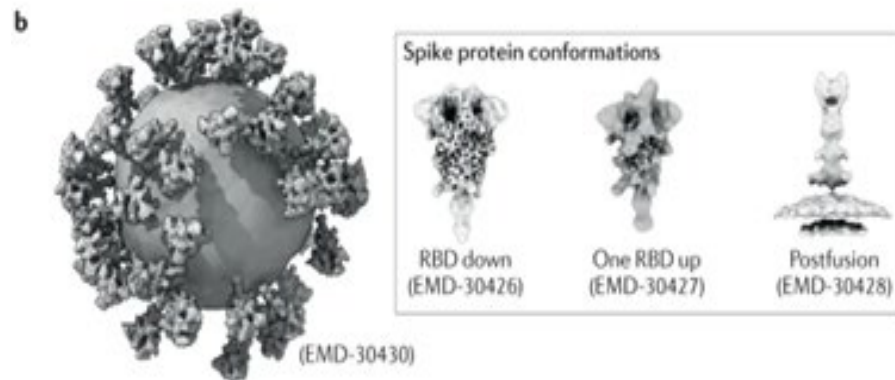
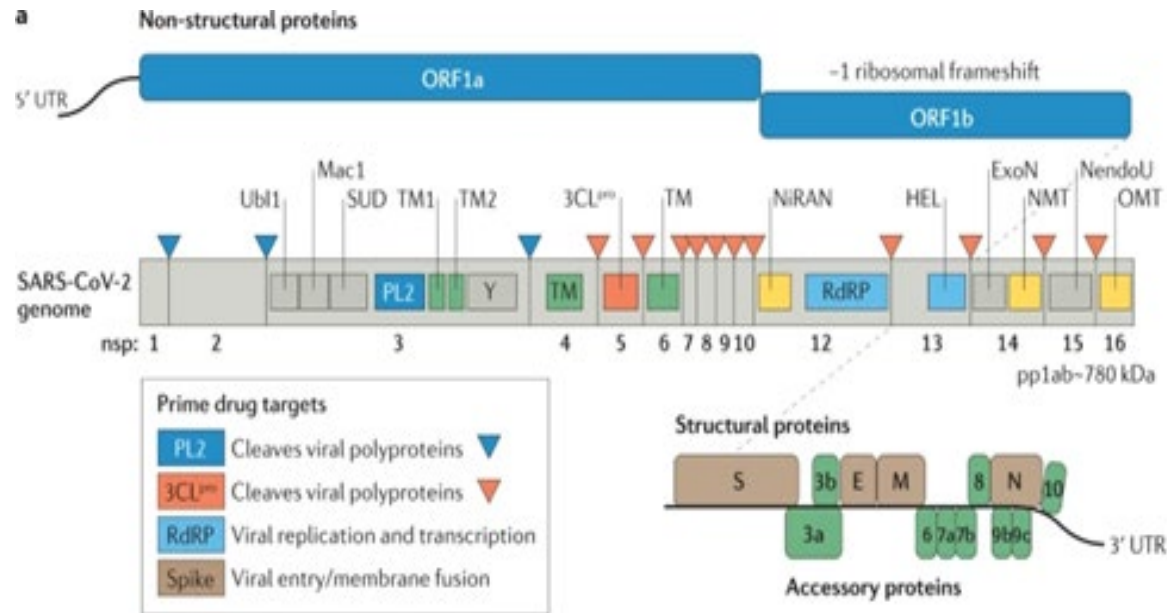
There are two competing ideas: a “zoonotic emergence” and a “laboratory escape” scenario.



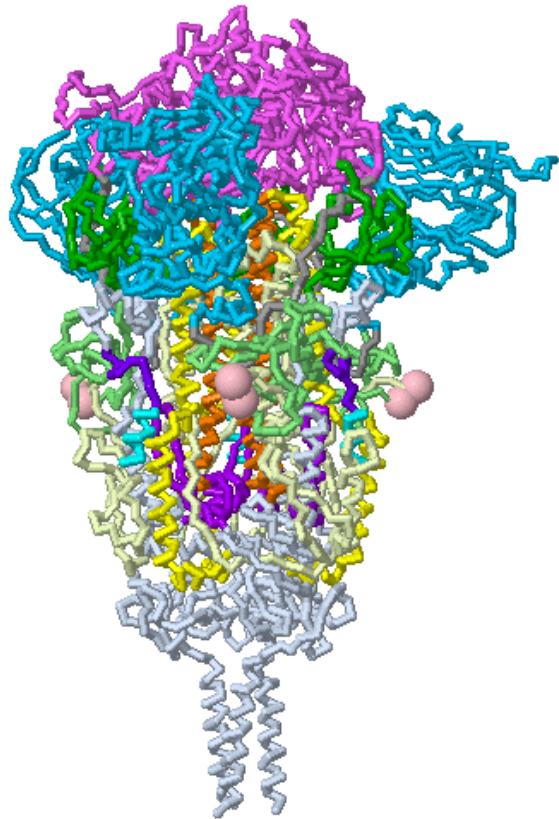
No bat reservoir or intermediate animal host was confirmed.

Given the genetic similarity of SARS-CoV-2 to bat SARS-CoV-like coronaviruses, bats may serve as reservoir hosts for their progenitor

Structure and Mechanisms of SARS-CoV-2



Spike protein of SARS-CoV-2

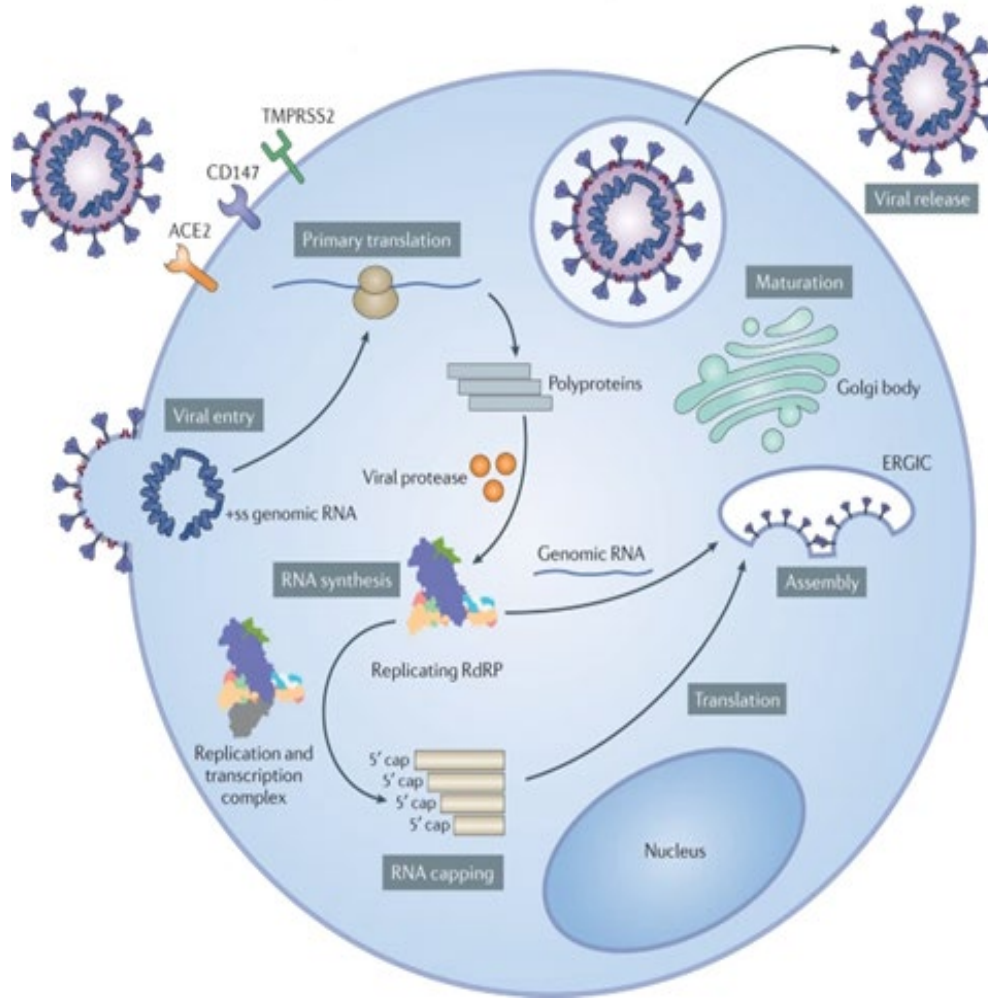


Jmol

S protein is responsible for the following:

- (1) Attachment at the host cell surface
- (2) Fusing the viral and host cell membranes
- (3) Viral entry (start of the infection)
- (4) Main target of neutralizing antibodies
- (5) Vaccine and therapeutic design

Entry pathways of SARS-CoV-2



SARS-CoV-2 variants in analyzed sequences

The number of analyzed sequences in the preceding two weeks that correspond to each variant group. This number may not reflect the complete breakdown of cases since only a fraction of all cases are sequenced.

- Alpha
 Beta
 Gamma
 Delta
 Omicron (BA.2)
 Omicron (BA.1)
 Omicron (BA.5)
 Omicron (BA.4)
- Omicron (BA.2.12.1)
 Omicron (BA.2.75)
 Omicron (BQ.1)
 Omicron (XBB)
 Omicron (XBB.1.5)
- Recombinant
 Other



Source: GISAID, via CoVariants.org – Last updated 12 June 2023

OurWorldInData.org/coronavirus • CC BY

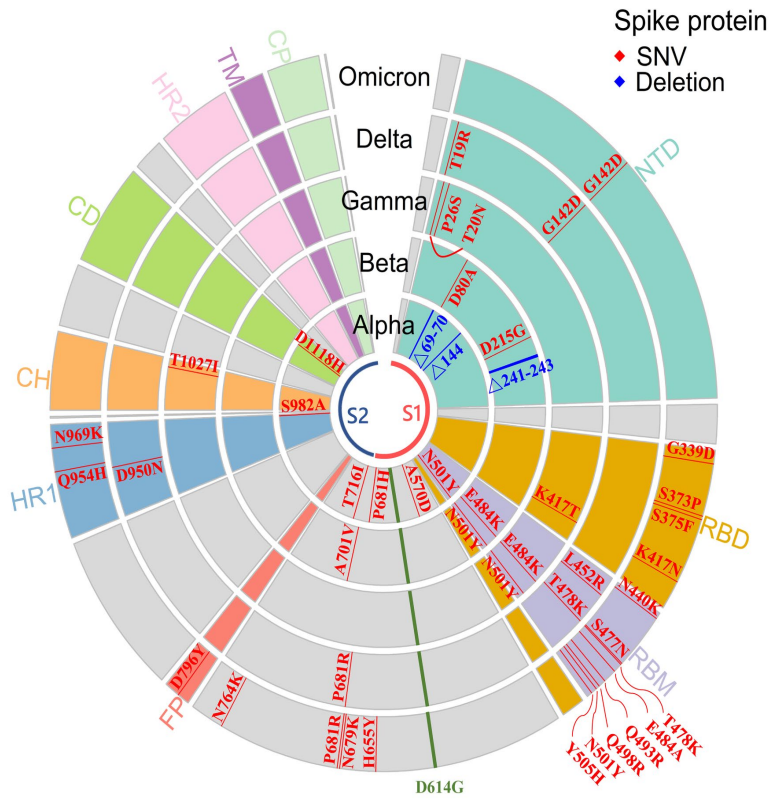
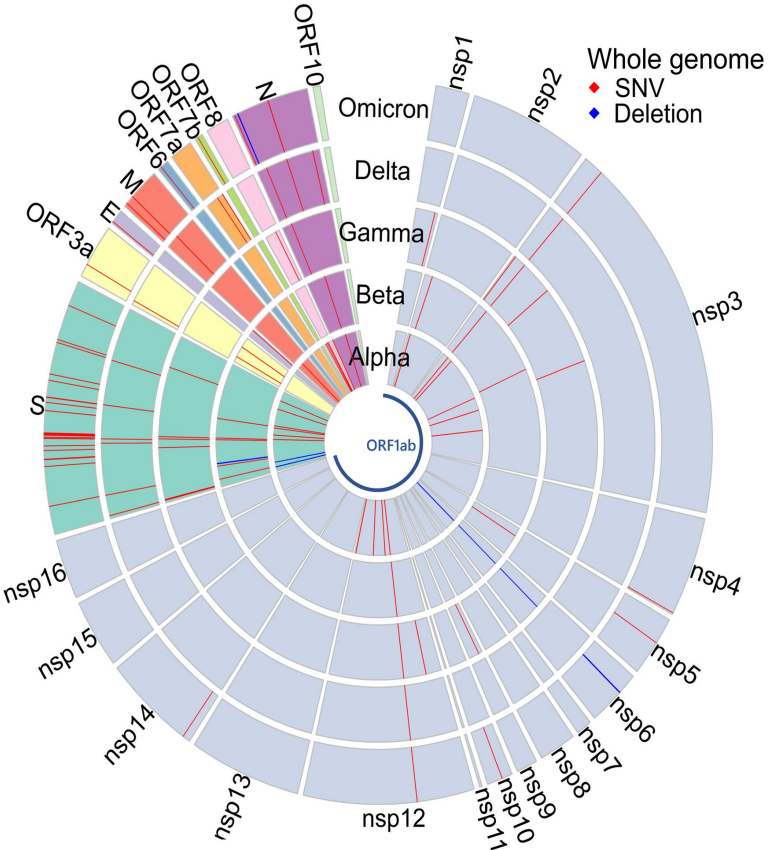
Note: Recently-discovered or actively-monitored variants may be overrepresented, as suspected cases of these variants are likely to be sequenced preferentially or faster than other cases.

XBB.1.5 is the most transmissible Omicron sub-variant currently.

Basic reproductive number R0

- Basic reproductive number **R0**:
 - Influenza **1.3**
 - 1918 Spanish flu **1.8**
 - Wu-Han variant **2.2 – 2.6**
 - D614G **3**
 - Alpha **4 – 5**
 - Delta **5 – 8**
 - Omicron **> Delta**
 - Early estimates of the R0 of BA.4 and BA.5 is **18.6**

Variants of SARS-CoV-2



COVID-19 diagnostic testing

- **A positive antigen test** that the person tested is **currently infected with COVID-19**, specifically during the **acute phase** of the infection.
- **A positive antibody test** suggests that the person tested **had a previous COVID-19 infection**; their immune system responded by producing antibodies (such as IgM and IgG) to combat the virus.
- **PCR tests** are considered the **“gold standard”** for COVID-19 testing because they are **highly accurate and sensitive in detecting the presence of the virus**; they are more reliable in identifying COVID-19 infections.

COVID-19 therapeutics

- **Antiviral Drugs** such as **Remdesivir** and **Molnupiravir** for inhibiting or interfering with **viral RNA replication**.
- **Immune Modulators** such as **Dexamethasone**, **Tocilizumab**, and **Baricitinib** for reducing **inflammation** and suppressing the **immune response**.
- **Monoclonal antibodies targeting SARS-CoV-2**, such as **Casirivimab and imdevimab (Regeneron)** and **Bamlanivimab and etesevimab (Eli Lilly)** for **neutralizing the virus and reducing viral replication**.
- **Convalescent plasma** for passively transferring antibodies from whom have recovered from COVID-19 to **help the recipient's immune response**.

COVID-19 vaccines

Vaccine platform	Advantages	Disadvantages
mRNA (Pfizer-BioNTech and Moderna)	Safe and well-tolerated; Highly adaptable to new pathogen; Native antigen expression	Lower immunogenicity; Requirement of low temperature storage and transportation; Potential risk of RNA-induced interferon response
Viral vector (Oxford-AstraZeneca and Johnson & Johnson/Janssen)	Stronger immune response; Preservation of native antigen; Mimicking natural infection	More complicated manufacturing process; Risk of genomic integration; Response dampened by pre-existing immunity against vector
Protein subunit (Novavax and Medigen Vaccine Biologics)	Safe and well-tolerated	Lower immunogenicity; Requirement of adjuvant or conjugate to increase immunogenicity
Inactivated virus (Sinovac and Bharat Biotech)	Stronger immune response; Safer than live attenuated virus	Potential epitope alteration by inactivation process

Non-Pharmaceutical Interventions (NPIs)

- NPIs, such as **social distancing**, **wearing masks**, **lockdowns**, and **quarantine measures**, are a range of measures and actions that can be taken to
 - **slow the transmission**
 - **protect vulnerable populations**
 - **ease the burden on healthcare systems**
 - **buy time for vaccine development and distribution**
 - **complement vaccine effectiveness**
- **Combining vaccination efforts with NPIs** offers the most effective approach to managing and mitigating the pandemic.

President James Liao's address on Mar 20, 2020

“..., Academia Sinica rapidly assembled a task force during the Lunar New Year period to initiate research work in **rapid viral detection, antibody screening, drug synthesis, and vaccine development**. In addition, we established working groups among different research organizations and academics in order to share information, materials, and methodologies for use in COVID-19 research ...”

In February 2020, Academia Sinica successfully synthesized Remdesivir at the “hundreds of milligrams level” with a purity of 97%, and further improved it to the “grams level” with a purity of 99%.

中研院 成功合成!

化學所陳榮傑研究團隊

百毫克級 瑞德西韋

- ✓ 從 0 開始
- ✓ 百毫克級
- ✓ 純度 97%
- ✓ 7 人團隊
- ✓ 日以繼夜
- ✓ 歷時 2 周

2020.02.20

f 中央研究院

In March 2020, Academia Sinica **discovered antibodies** capable of detecting viral antigens, which can be applied as **rapid screening reagents**.



中研院新冠病毒快篩檢測試劑 研發步驟

1. 新冠病毒核蛋白製備 → 2. 以噬菌體合成抗體庫進行抗體篩選



19天完成抗體篩選



3. 抗體特性分析及快篩抗體配對與原型製作

4. 快篩檢測裝置測試及驗證

- 廠商生產快篩試劑
- 食藥署驗證
- 衛福部許可



下一階段:快篩檢測裝置臨床應用



快篩步驟快速簡單



只需15至20分鐘

When the SARS outbreak occurred in Taiwan, Academia Sinica promptly **constructed two P3 negative-pressure laboratories** for developing COVID-19 rapid screening, viral testing, and vaccine and therapeutics.



In January 2021, Academia Sinica repurposed existing medicines (mefloquine, nelfinavir, and extracts of *Ganoderma lucidum*, *Perilla frutescens*, and *Mentha haplocalyx*) as anti-COVID-19 virus agents



In June 2023, Academia Sinica and Moderna jointly launch the Forward-looking Innovation Award to accelerate mRNA research in Taiwan





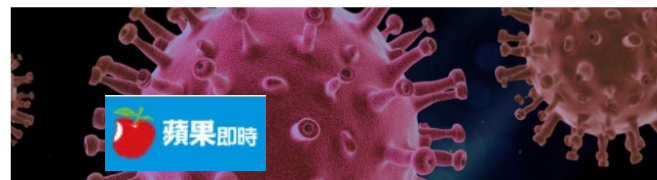
中研院發現全球流行同一株新冠病毒！第6型病毒適應性強

日期：2020-12-14 作者：王芊漢



英國變種病毒 中研院：第6型病毒株子型 可能將繼續變異

最新更新：2020/12/30 17:35



【武肺變種】台確診首例！中研院QA解疑 「現有疫苗還可有效對抗」

更新時間：2020/12/31 11:09

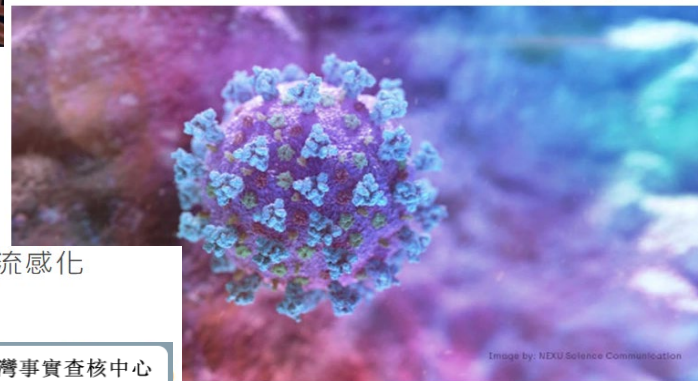


Image by: NEXU Science Communication

變種病毒傳播更快！中研院揭「源頭真相」：恐繼續變異

2020/12/31 06:00:00

追蹤三立:



中研院揪「至尊」病毒株 醫：武肺漸流感化

更新時間：2020/12/13 03:00



最新查核報告 政治與政策 生活 健康 科技資安 環境能源 國際 科學研究 研究動向

研究與動態

分享:

【中研院團隊解惑】10個Q&A 掌握變種病毒

更新日期：2021-02-19

記者馬麗昕、劉芮菁／報導

對新冠病毒的知識掌握越多，就越能幫助我們用「科學」來有效防疫。查核中心觀察社群媒體的「變種病毒」假訊息增多，為此查核中心訪問中研院「病毒即時監測跨領域團隊」成員，包括中研院統計科學研究所研究員楊欣洲，與生物醫學科學研究所研究員林宜玲，整理出10個問答，讓讀者可以簡單搞懂變種病毒。



GISAID (Global Initiative on Sharing Avian Influenza Data)



In Focus

Recommended composition of influenza virus vaccines for use in the 2023-2024 Northern Hemisphere Influenza Season

GISRS
GLOBAL INFLUENZA SURVEILLANCE & RESPONSE SYSTEM
Uniting the world in the fight against influenza

H5N1
A clade subsequence including... globe, 1.2.3.4.41... ever ca...
Events... link... no kno... partner... > for m

...nced
...An advisory group of experts taking part in a meeting organized by the Global Influenza Programme between 20-23 February 2023 analyzed influenza virus surveillance data generated by the Global Influenza Surveillance and Response System (GISRS) and issued on 24 February 2023, recommendations on the composition of the influenza vaccines for the upcoming influenza season.
...Recommendations are used by the national vaccine regulatory agencies and pharmaceutical companies to develop, produce and license influenza vaccines.

[More:](#)

中国新冠疫情 ▶ Lineage comparison ▶

hCoV-19 data sharing via GISAID

15,783,309
genome sequence submissions

Submission Tracker	Phylogenetics	Tracking Variants	Frequency Dashboards
<ul style="list-style-type: none"> hCoV-19 Global hCoV-19 USA hMpxV RSV 	<ul style="list-style-type: none"> hCoV-19 hMpxV RSV 	<ul style="list-style-type: none"> hCoV-19 Variants hMpxV Variants Influenza Subtypes RSV Subtypes 	<ul style="list-style-type: none"> hCoV-19 hMpxV Influenza RSV

GISAID Resources

Data Acknowledgement Locator

EPI_SET ID

Example ID

[Register for Access Credentials](#)

FASTA data → Data QC → Alignment

- **March 31, 2020: 1,932 strains:**
(For the initial classification and signature SNV analysis)
- **April 19, 2020: 6,228 strains:**
(For the main analysis)
- **June 8, 2020: 38,248 strains:**
(For a rapid strain typing)

Genome annotation

1	...	265	266	267	268	...	805	...	29675	...	29903	PP
5'UTR	...	5'UTR	nsp1	nsp1	nsp1	...	nsp1	...	3'END	...	3'END	Protein
...	...	G>T	A>T	T>G	T>G	...	G>A	...	G>A	...	C>T	NT
...	...	NA	M>L	M>R	M>R	...	G>G	...	NA	...	NA	AA
...	...	T	A>T	T>G	NA	...	G>G	...	NA?	...	NA?	
...	...	T	A>T	T>G	NA	...	-	...	NA?	...	NA?	
:	:	:	:	:	:	:	:	:	:	:	:	
...	...	G	A>A	T>T	NA	...	G>A	...	---	...	---	

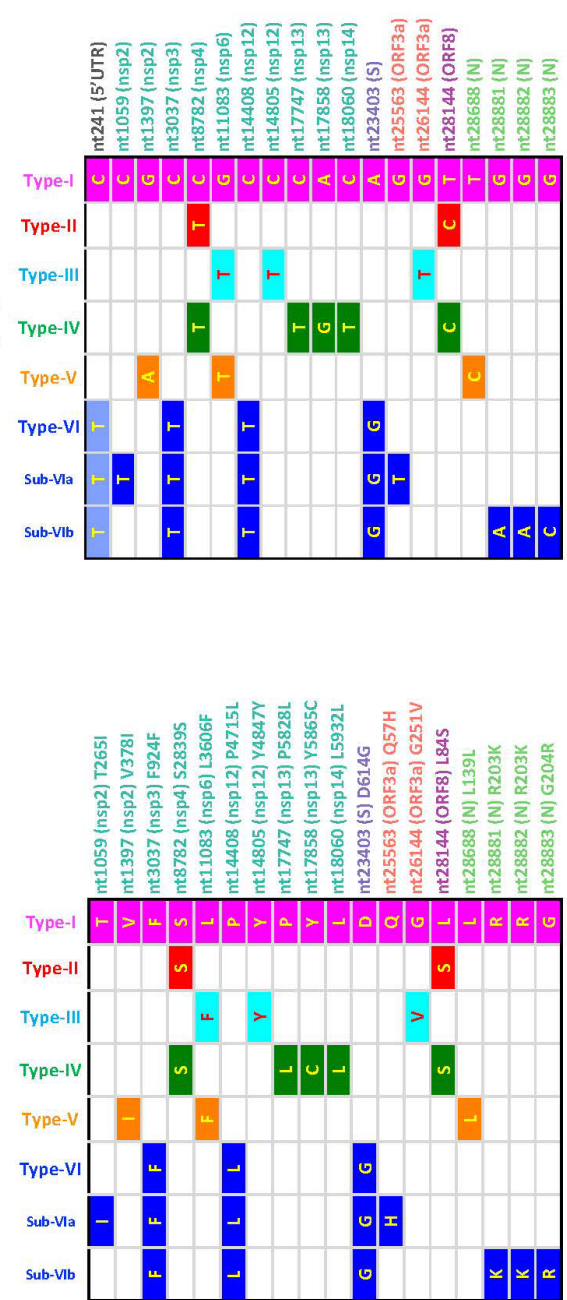
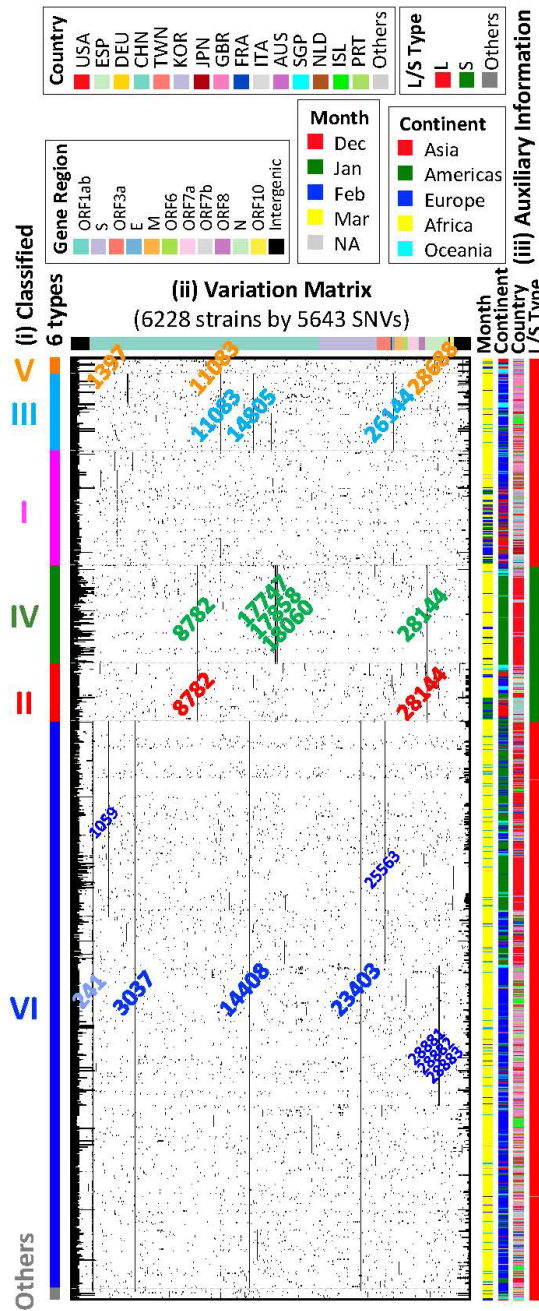
No.	Strain	Country	Date
1	EPI_ISL_581117	Europe/UK /England	2020/09/21
2	EPI_ISL_601443	Europe/UK /England	2020/09/20
:	:	:	:
>1M	EPI_ISL_414380	Asia/Singapore	2021/03/05

Strain metadata

Nucleotide sequence

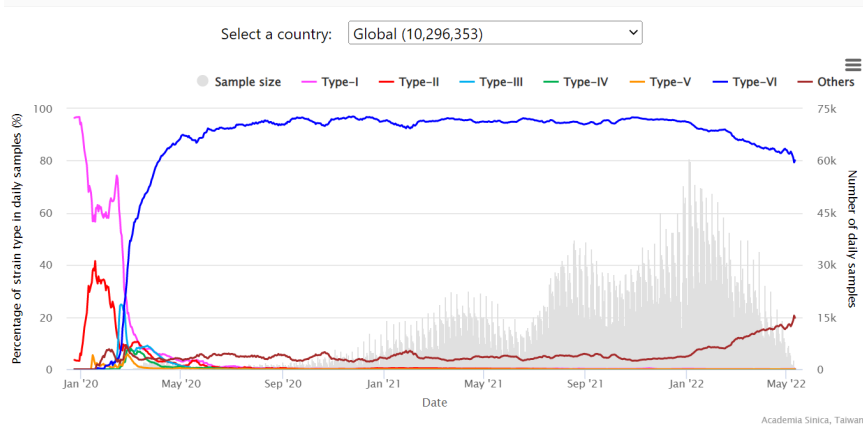
Remark: Variations were detected by comparing to the reference genome of **Wuhan-Hu-1** (Wu et al., Nature 579, 265-269 (2020)).

In April 2020, Academia Sinica identified six major types of COVID-19 virus and Type VI becomes the dominant strain in the world.



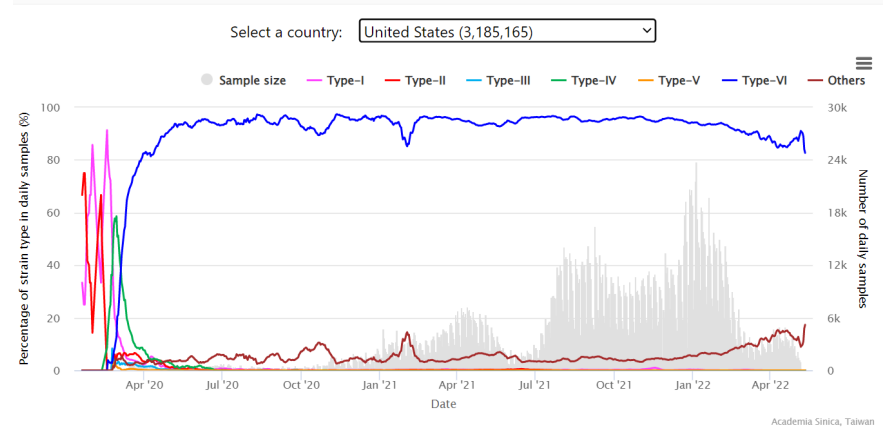
Type VI became the dominant strain in the world

Temporal Distributions of Six Strain Types



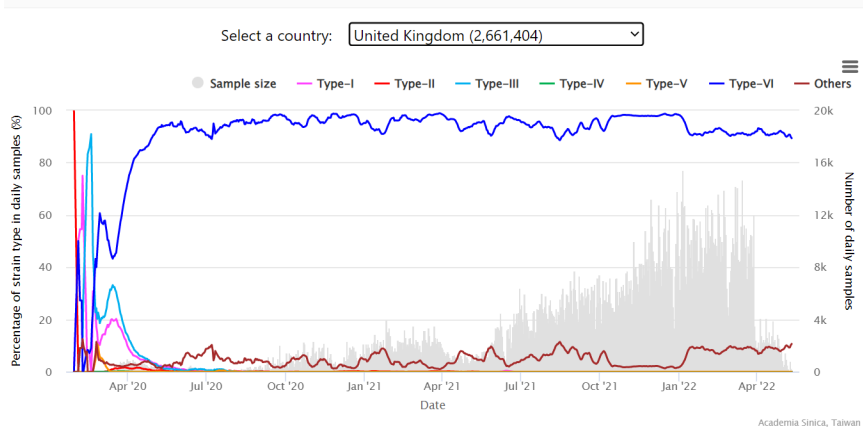
Distribution of Strain Types over time: Global

Temporal Distributions of Six Strain Types



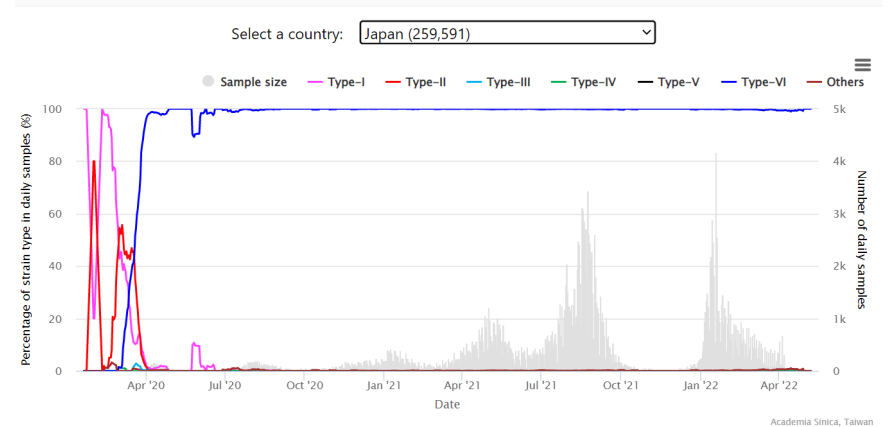
Distribution of Strain Types over time: United-States

Temporal Distributions of Six Strain Types



Distribution of Strain Types over time: United-Kingdom

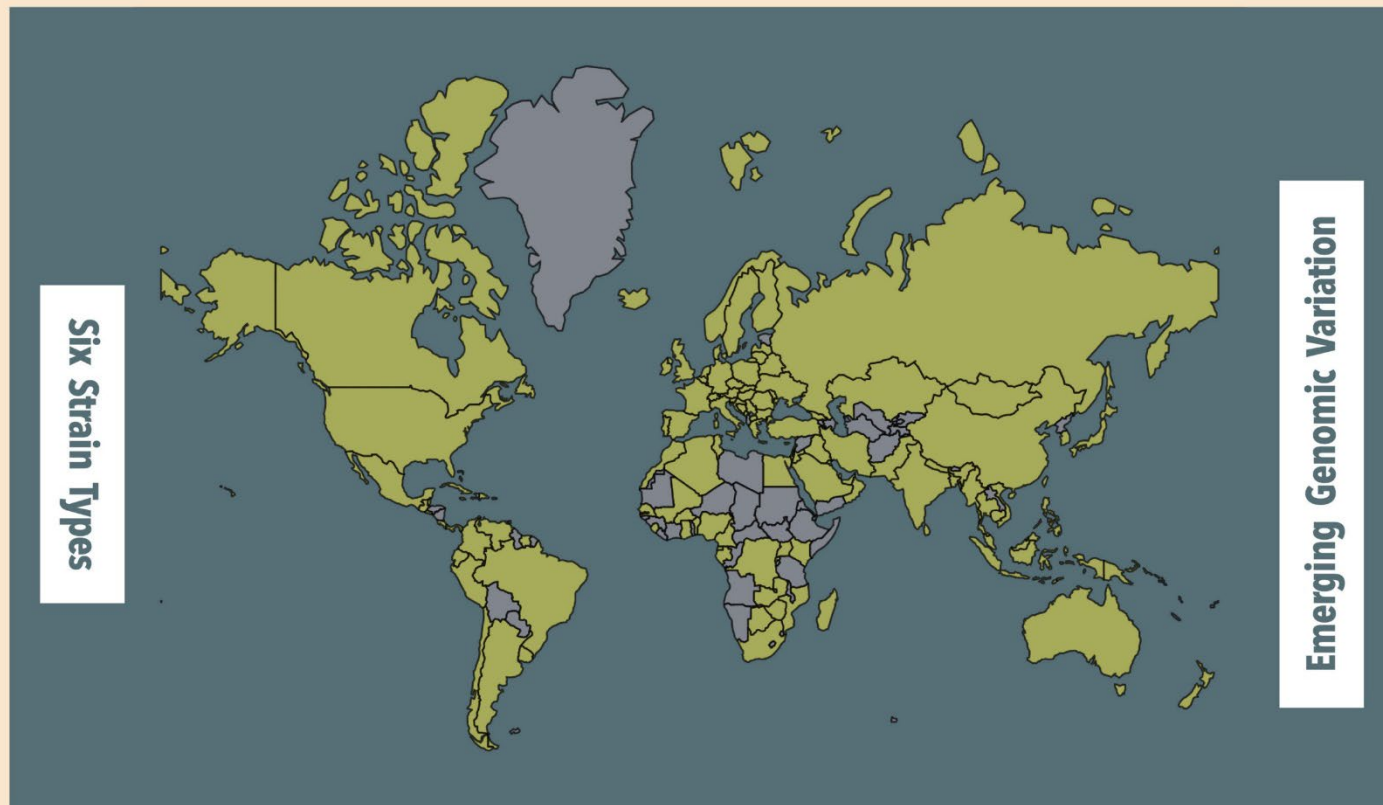
Temporal Distributions of Six Strain Types



Distribution of Strain Types over time: Japan

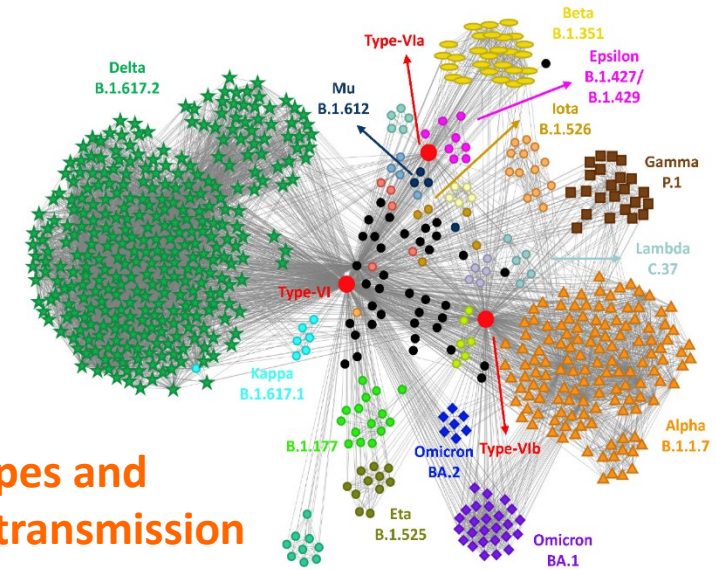
In December 2020, Academia Sinica established the **SARS-CoV-2 Variation Monitoring Network** to track the transmission dynamics and evolution of SARS-CoV-2 variants.

SARS-CoV-2 Variation Monitoring Network

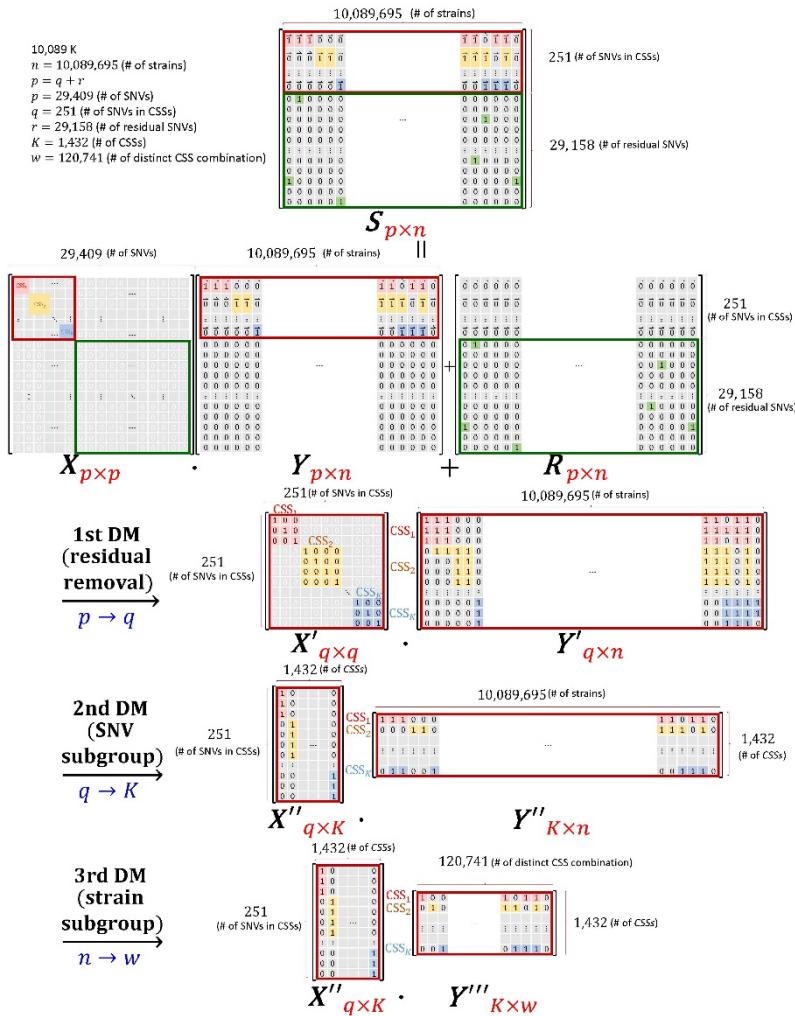
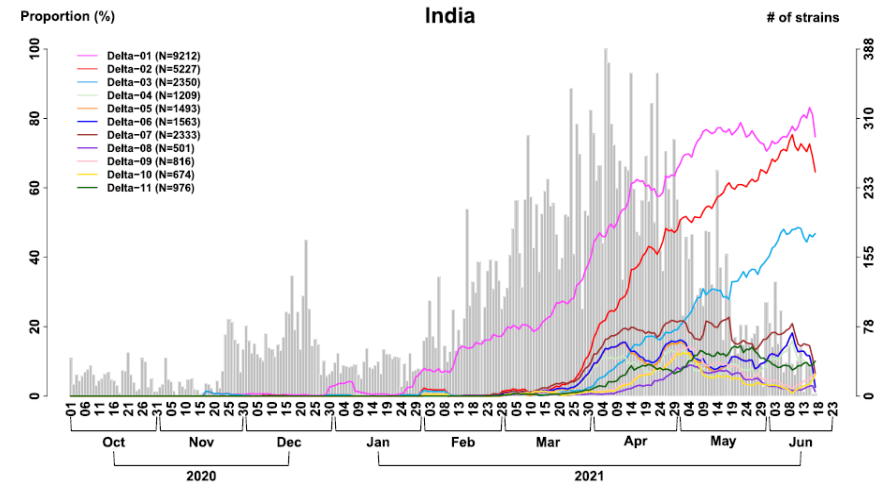


Subtyping of major SARS-CoV-2 variants reveals different transmission dynamics based on 10 million genomes

(B) Virus network analysis



(C) Subtypes and dynamic transmission of Delta variant



(A) An efficient data dimensionality reduction

Taiwan's experience with the COVID-19 pandemic (1)

Lessons from SARS in 2003:

- Regular reviews for **infection control** and **drills** in the medical care system have been held periodically. The hospitals could **activate immediate measures** such as triage, quick isolation, strictly controlled crowd flow, medical task force, and epidemic prevention materials to **maintain the normal operation of the medical care system**.
- Residents remained **self-alert** during the disease pandemic, voluntarily put on **face masks**, **wash hands** properly, and practiced **social distancing**.

Taiwan's experience with the COVID-19 pandemic (2)

- In January 2020, when news of a “new and unknown respiratory disease emerging in China” was first disclosed, an inter-ministerial **Central Epidemic Command Center** was activated, providing four important principles — **Rapid measures**, **Early deployment**, **Prudent actions**, and **Transparency** for blocking virus spread into Taiwan.
- **Taiwan CDC** began to implement relevant prevention strategies, including **Surveillance and laboratory diagnosis**, **Border control**, **Control of community transmission**, **Medical system response and preparedness**, and **Health education and fighting disinformation**.
- When the supply of face masks met the domestic demand, Taiwan began to **donate medical supplies to more than 80 countries worldwide** to assist them in combating COVID-19 and to realize the spirit of “**Taiwan Can Help, and Taiwan is Helping!**”

Lessons Learned from the Global COVID-19 Pandemic

- Preparedness and Early Response
- Importance of Global Cooperation
- Healthcare System Strengthening
- Vaccine Development and Distribution
- Importance of Public Health Measures
- Health Inequities and Vulnerable Populations
- Digital Transformation and Telemedicine
- Risk Communication and Misinformation
- Economic Impact and Social Resilience

COVID-19 is not the final global health challenge we will face, highlighting the ongoing need for vigilance, preparedness, and investment in public health infrastructure to effectively respond to future infectious diseases.

Thank you for your attention!