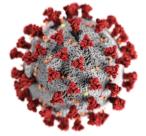


University of Baghdad College of Pharmacy





## Structural Basis of SARS-CoV-2 Receptor Binding and the Small Molecule Compounds Available as Potential Therapeutics

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Thursday, March 10, 2022

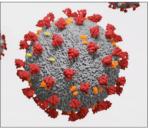
## Outline

- COVID-19 Pandemic
- Structure of SARS-CoV-2
- Entry and Lifecycle of SARS-CoV-2
- Potential small-molecule compounds available against SARS-CoV-2
- Conclusions

## **COVID-19 Pandemic**

CoVs caused three outbreaks :

Virus	SARS-CoV	MERS-CoV	SARS-CoV-2
Disease	Severe acute respiratory syndrome	Middle eastern respiratory syndrome	Coronavirus infectious disease 2019 (COVID-19)



- Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2)
- Declared a pandemic by the WHO in March 2020.
- Evidence for nonsymptomatic/presymptomatic spread.

https://doi.org/10.1016/j.it.2020.10.004

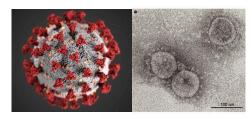
## **COVID-19 Pandemic**

- Since its emergence, it has infected millions of people, killed close to 3 million people worldwide, and costs world economies trillions of dollars.
- Several vaccines are being administered to people for protective immunity.
- Additional COVID-19 therapeutics are needed.
- Emergence of new sars-cov-2 variants.
- Many people are not becoming vaccinated.
- Repurposing approach with the small molecules to accelerate the development process.

https://doi.org/10.1155/2021/1828792

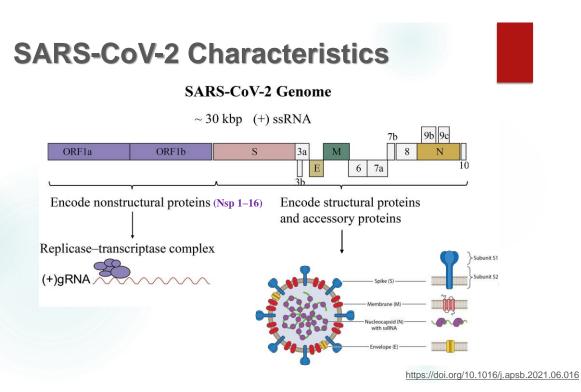
## **SARS-CoV-2** Characteristics

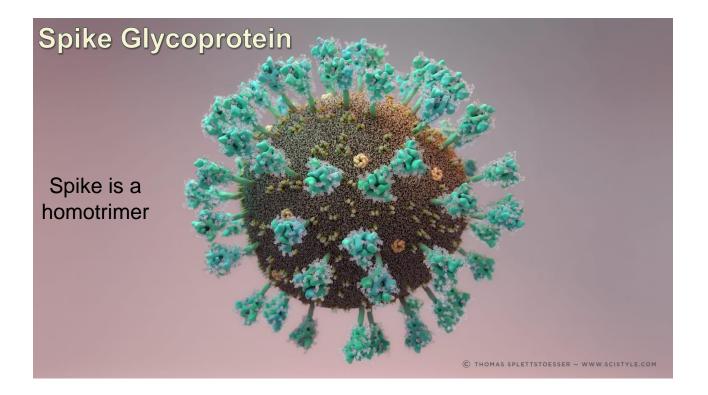
- Genus: Betacoronavirus.
- Enveloped, positive-sense singlestranded RNA virus.
- The outer surface peppered with 24– 40 spike proteins: attachment/entry.
- The receptor for the S protein is ACE2, expressed on : alveolar cells, esophageal cells, enterocytes, and others.

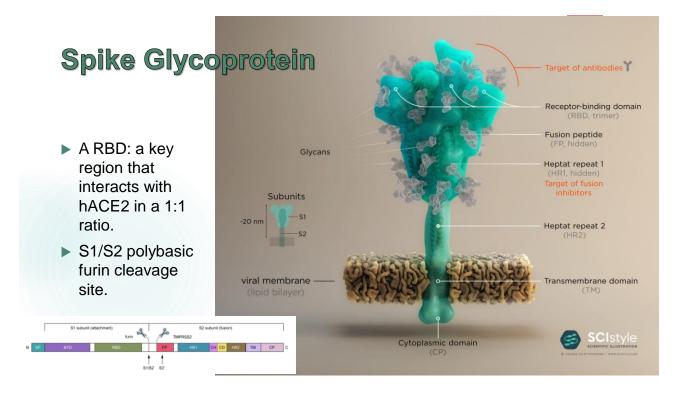


Corona=crown

https://doi.org/10.1016/j.it.2020.10.004 doi:10.3390/jcm9061885

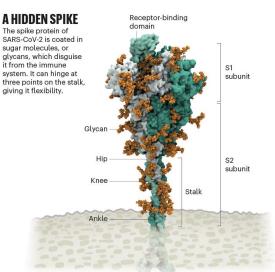






## Spike Glycoprotein

- Wildly flexible and hinge at three points.
- The S-glycoprotein has 22N-linked glycosylation sites that are spread throughout the NTD and RBD.
- ACE2 N-glycans interacts with SARS-CoV-2 spikes.



doi: 10.1038/d41586-021-02039-y. @nature

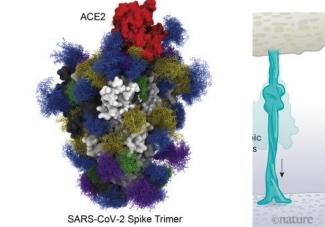
## SARS-CoV-2 Entry Mechanism

Spike protein

TMPR

- Activation by host proteases are required.
- SARS-CoV-2 binds to ACE2 on the target cell
- TMPRSS2 cuts a site on S2 subunit.
- That cut exposes hydrophobic amino acids that rapidly buries itself in the host cell membrane.

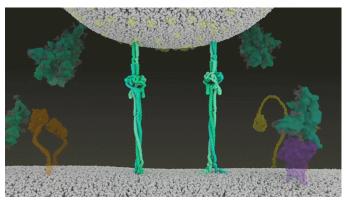
Glycomics-informed glycoproteomics & molecular dynamics simulations



https://doi.org/10.1146/annurev-pharmtox-061220- 093932 doi: 10.1038/d41586-021-02039-y.

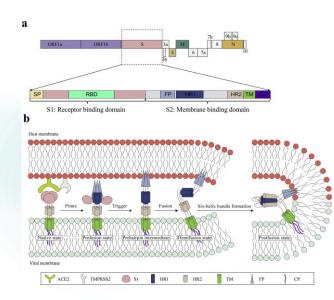
## SARS-CoV-2 Entry Mechanism

- The extended spike folds back onto itself, like a zipper, forcing the viral and cell membranes to fuse.
- The virus then ejects its genome into the cell.



doi: 10.1038/d41586-021-02039-y.

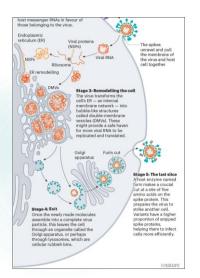
## SARS-CoV-2 Entry Mechanism



During fusion with the host, the S1 and S2 subunits separate. The S2 subunit forms a 6-HB, with a HR1 and HR2 region that fuses with the host cell membrane.

> https://doi.org/10.1146/annurev-pharmtox-061220- 093932 https://doi.org/10.1016/j.apsb.2021.06.016

## **SARS-CoV-2 Translation**



- ✓ On the inside of the cell, SARS-CoV-2 transforms the ER into DMVs providing a safe place for replication and translation.
- Proteins involved in making DMVs could be good drug targets.

doi: 10.1038/d41586-021-02039-y.

# <image>

- Ribosomes translate viral RNA.
- Papain-like protease (PLpro)
- 3C-like protease (3CLpro, Mpro)
- RNA- dependent RNA polymerase (RdRp)

<u>https://doi.org/10.1016/j.it.2020.10.004</u> <u>doi: 10.1038/d41586-021-02039-y.</u> <u>https://doi.org/10.1038/s41392-021-00733-x</u>

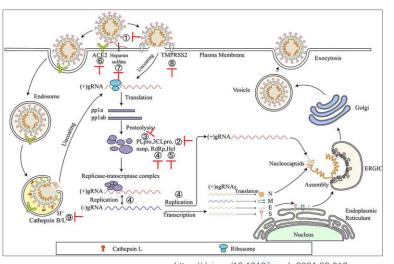
## **SARS-CoV-2** Translation

- Three mechanisms by which SARS-CoV-2 suppresses the translation of host mRNA in favour of its own:
- 1. 1. Nsp1 recruits host proteins to chop up all cellular mRNAs that don't have a viral tag.
- Reduces overall protein translation in the cell by 70%. Nsp1 physically block the entry channel of ribosomes so mRNA can't get inside.
- 3. 3. The virus shuts down the cell's alarm by preventing cellular mRNA from getting out of the nucleus.

doi: 10.1038/d41586-021-02039-y.

## **Therapeutic Targets**

- S-protein and Nsps.
- Protease inhibitors: prevent a virus from using TMPRSS2, cathepsin L or other proteases to enter host cells.
- Absence of a human homolog make Mpro an attractive target.



https://doi.org/10.1016/j.apsb.2021.06.016 https://doi.org/10.1146/annurev-pharmtox-061220- 093932

# Targeting Viral Entry via the Endosomal Pathway

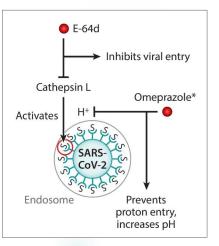
### Drugs targeting endosome acidification:

HCQ, CQ, and ammonium chloride: inhibit viral replication by increasing lysosome pH.

Questionable benefit.

Omeprazole: interferes with lysosomal activity and inhibits double-stranded RNA formation.

Combinations.

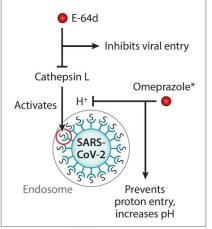


https://doi.org/10.1146/annurev-pharmtox-061220- 093932

# Targeting Viral Entry via the Endosomal Pathway

#### Drugs targeting cathepsin:

- ► E64d: a broad-spectrum inhibitor, targets lysosomal cathepsin.
- In combination with camostat mesylate could fully block viral entry.



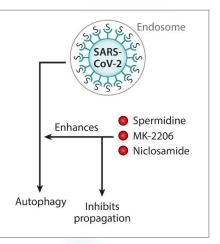
# Targeting Viral Entry via the Endosomal Pathway

## Drugs targeting the autophagy pathway:

**Spermidine**: induces autophagy.

Questionable

- ► MK-2206: Inhibit of AKT1→upregulates Beclin-1 and promotes autophagy. MK-2206 reduced SARS-CoV-2 propagation by 88%.
- Niclosamide: an orally bioavailable, chlorinated salicylanilide. It inhibits SKP2, stabilizes Beclin-1, and enhances autophagy. It also blocks endosomal acidification.

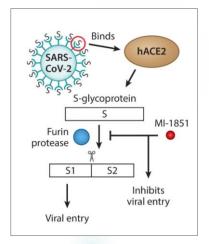


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## Targeting Viral Entry Through the Plasma Membrane Fusion Pathway

# Drugs targeting furin and TMPRSS2 proteases:

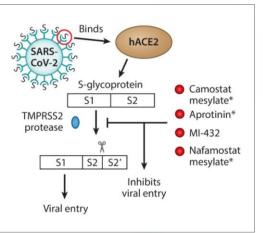
- MI-1851: a synthetic furin inhibitor potently inhibits S-glycoprotein cleavage.
- Reduced viral titers by 30- to 75-fold.



## Targeting Viral Entry Through the Plasma Membrane Fusion Pathway

#### Drugs targeting furin and TMPRSS2 proteases:

- Camostat mesylate/Nafamostat mesylate : approved TMPRSS2 inhibitors.
- Aprotinin: shows a high antiviral effect. Also interferes with formation of dsRNA.
- MI-432: a synthetic peptide mimetic inhibitor of TMPRSS2
- Combination of MI-432 and MI-1851 has an increased anti-SARS-CoV-2 activity.

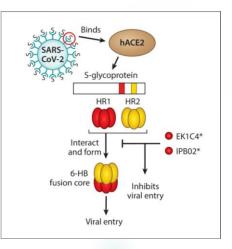


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# Targeting Viral Entry via the Endosomal Pathway

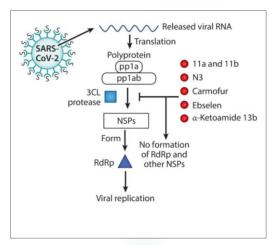
#### Drugs targeting HR1 of S2 subunit of S-glycoprotein:

- EK1C4: disrupt formation of the 6-HB fusion core by binding to the HR1 domain.
- Modification of EK1 with a cholesterol moiety, EK1C4 forms a more stable complex with HR1, which enhances its antiviral activity (149-fold).
- IPB02: lipopeptide fusion inhibitor that targets the HR1 region.



## **Targeting the Main Protease (Mpro)**

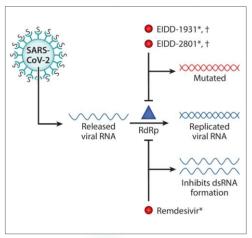
- 11a and 11b: covalently binds to the catalytic site Cys145 and blocks enzymatic activity.
- N3: binds irreversibly in the Mpro substrate recognition pocket.
- Ebselen and carmofur: Covalently bind to the catalytic Cys145.
- α-Ketoamide 13b: block Mpro by 2 hydrogen bonding interactions.



https://doi.org/10.1146/annurev-pharmtox-061220- 093932



- Remdesivir: potently stops viral replication activity via RNA chain termination.
- NHC (EIDD-1931): potent antiviral activities against RNA virus replication: Ebola virus, SARS-CoV, and MERS-CoV.
- NHC exhibits only a low level of resistance with multiple viruses.
- EIDD-2801: an isopropyl ester prodrug of NHC with improved oral bioavailability and pharmacokinetics in vivo.



## Conclusions

- Despite global efforts, COVID-19 remains a serious concern. Although many clinical trials of the repurposed drugs, immune-based therapies and investigational antivirals have been conducted, there is still no highly effective therapeutic available. The mutation of SARS-CoV-2 make vaccine and drug discovery more uncertain. Accordingly, developing specific or broad-spectrum inhibitors is urgently needed.
- The outbreak of COVID-19 has highlighted the importance for the development of broad-spectrum antiviral agents to combat future coronaviruses.

