

# SARS-CoV-2 (COVID-19) by the numbers

Yinon M. Bar-On<sup>1</sup>, Avi Flamholz<sup>2</sup>, Rob Phillips<sup>3,4</sup>, and Ron Milo<sup>1†</sup>

<sup>1</sup>Weizmann Institute of Science, Rehovot 7610001, Israel

<sup>2</sup>University of California, Berkeley, CA 94720, USA

<sup>3</sup>California Institute of Technology, Pasadena, CA 91125, USA

<sup>4</sup>Chan Zuckerberg Biohub, 499 Illinois Street, SF CA 94158, USA

†Corresponding author, send comments to [ron.milo@weizmann.ac.il](mailto:ron.milo@weizmann.ac.il); see updated version at: <https://bit.ly/2UBtcQ5>

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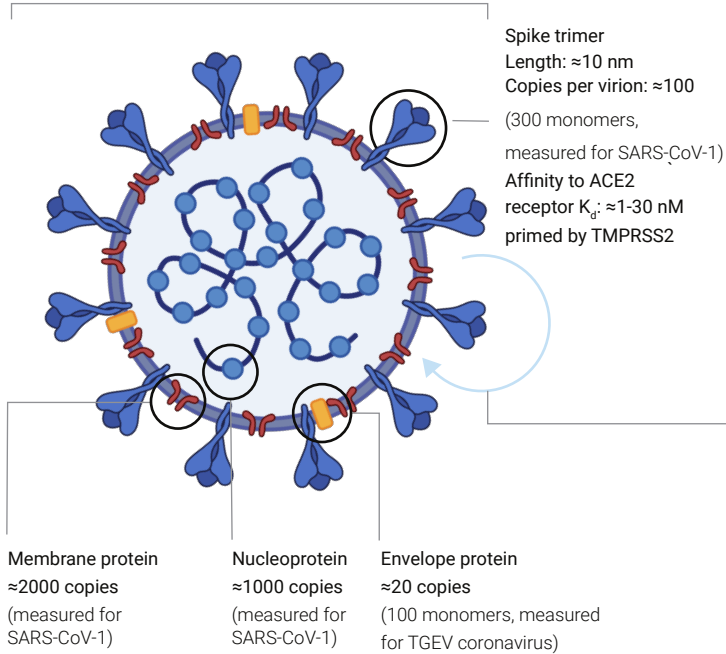
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## Size

Diameter:  $\approx 100$  nm

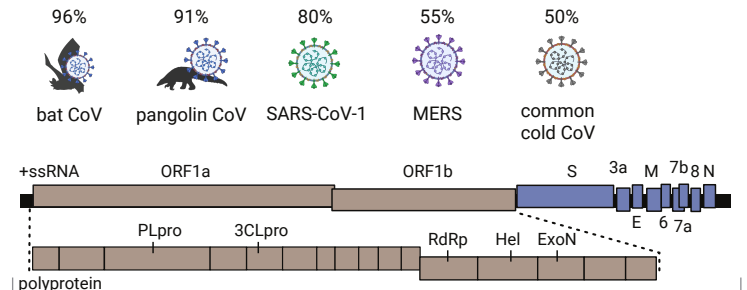
Volume:  $\sim 10^6 \text{ nm}^3 = 10^{-3} \text{ fL}$

Mass:  $\sim 10^3 \text{ MDa} = 1 \text{ fg}$



## Genome

Nucleotide identity to SARS-CoV-2



Length:  $\approx 30$  kb;  $\beta$ -coronavirus with 10-14 ORFs (24-27 proteins)

Evolution rate:  $\sim 10^{-3} \text{ nt}^{-1} \text{ yr}^{-1}$  (measured for SARS-CoV-1)

Mutation rate:  $\sim 10^{-6} \text{ nt}^{-1} \text{ cycle}^{-1}$  (measured for MHV coronavirus)

## Replication Timescales

in tissue-culture

Virion entry into cell:  $\sim 10$  min (measured for SARS-CoV-1)

Eclipse period:  $\sim 10$  hrs (time to make intracellular virions)

Burst size:  $\sim 10^3$  virions (measured for MHV coronavirus)

## Host Cells

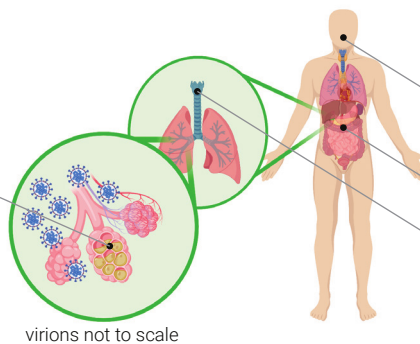
(tentative list; number of cells per person)

Type I & II pneumocytes ( $\sim 10^{11}$  cells)

Alveolar macrophage ( $\sim 10^{10}$  cells)

Mucous cell in nasal cavity ( $\sim 10^9$  cells)

Host cell volume:  $\sim 10^3 \mu\text{m}^3 = 10^3 \text{ fL}$



## Concentration

maximal observed values following diagnosis (ref, ref, ref)

Nasopharynx:  $10^6$  -  $10^9$  RNAs/swab

Throat:  $10^4$  -  $10^{11}$  RNAs/swab

Stool:  $10^4$  -  $10^8$  RNAs/g

Sputum:  $10^6$  -  $10^{11}$  RNAs/mL

RNA counts can markedly overestimate infectious virions

## Antibody Response - Seroconversion

Antibodies appear in blood after:  $\approx 10$ -20 days

Maintenance of antibody response:

$\approx 2$ -3 years (measured for SARS-CoV-1)

## Virus Environmental Stability

Relevance to personal safety unclear

half-life (ref)

time to decay 1000-fold

Aerosols:  $\approx 1$  hr  $\approx 4$ -24 hr

Surfaces:  $\approx 1$ -7 hr  $\approx 4$ -96 hr

e.g. plastic, cardboard and metals

Based on quantifying infectious virions. Tested at 21-23°C and 40-65% relative humidity. Numbers will vary between conditions and surface types (ref).

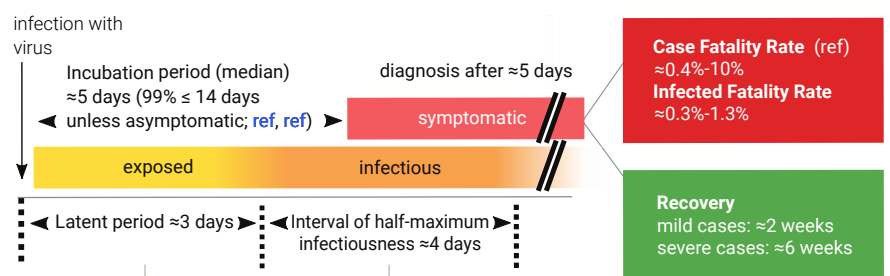
Viral RNA observed on surfaces even after a few weeks (ref)

## "Characteristic" Infection Progression in a Single Patient

Basic reproductive number  $R_0$  (ref, ref):

typically 2-4, but varies further across space and time

(number of new cases directly generated from a single case)



Inter-individual variability is substantial and not well characterized. The estimates are parameter fits for population median in China and do not describe this variability (ref, ref).