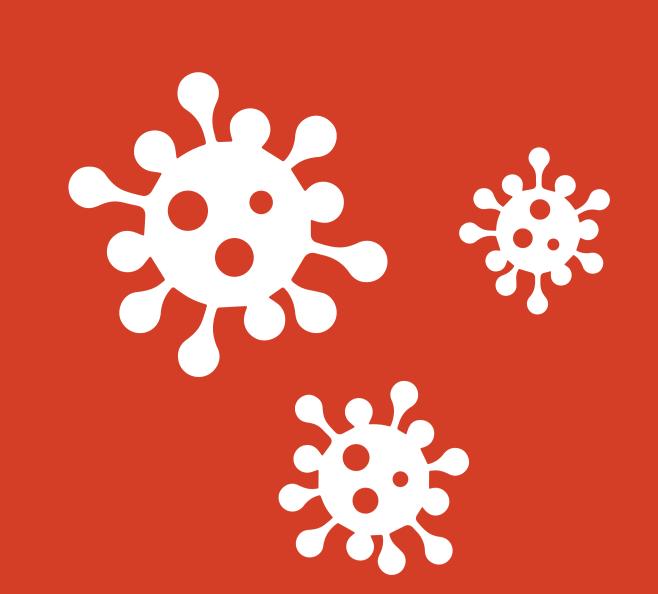


Autònoma de Barcelona

SARS-CoV-2. What's the infection origin in humans?

José Luis Orri Fabrellas

Date: June 2021



Introduction **SARS-CoV-2**

- it started in Wuhan, China.
- ** Evidence, the concept "ONE HEALTH" in a globalized world.
- Rhinolophus ferrumequinum reservoir.
- * Uncertainty about the intermediate host I.

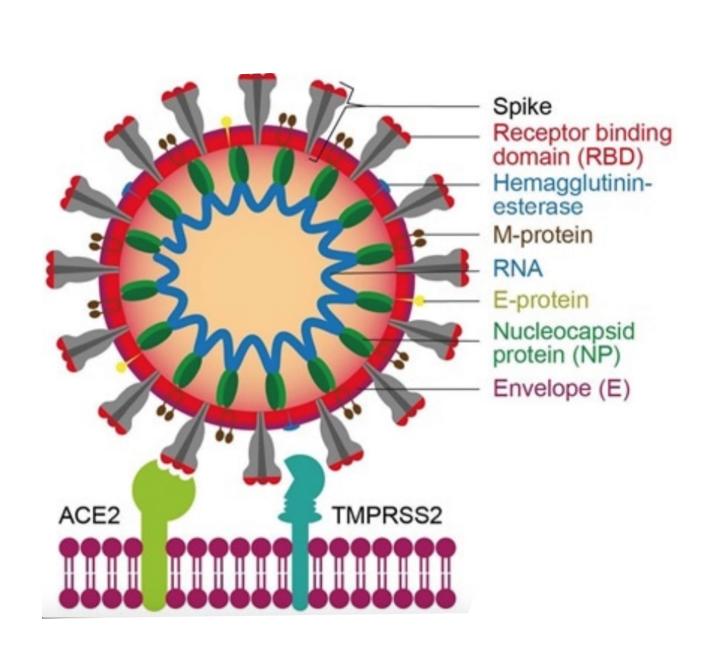
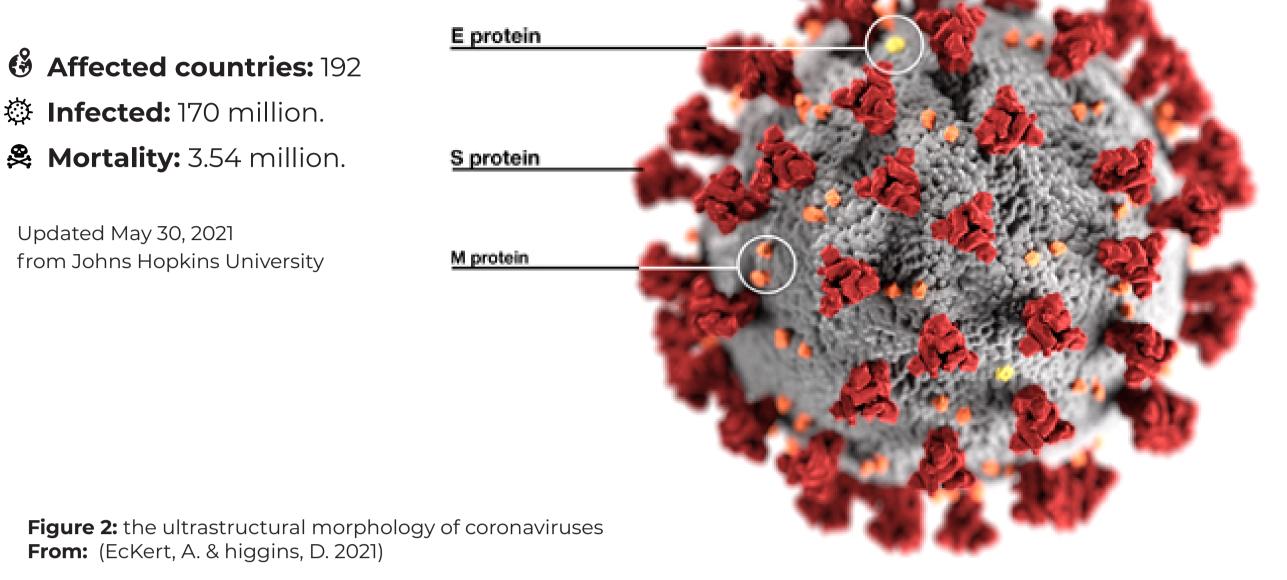


Figure 1: SARS-CoV-19. Structure From: (Robertson, 2021)

Current state of the pandemic situation

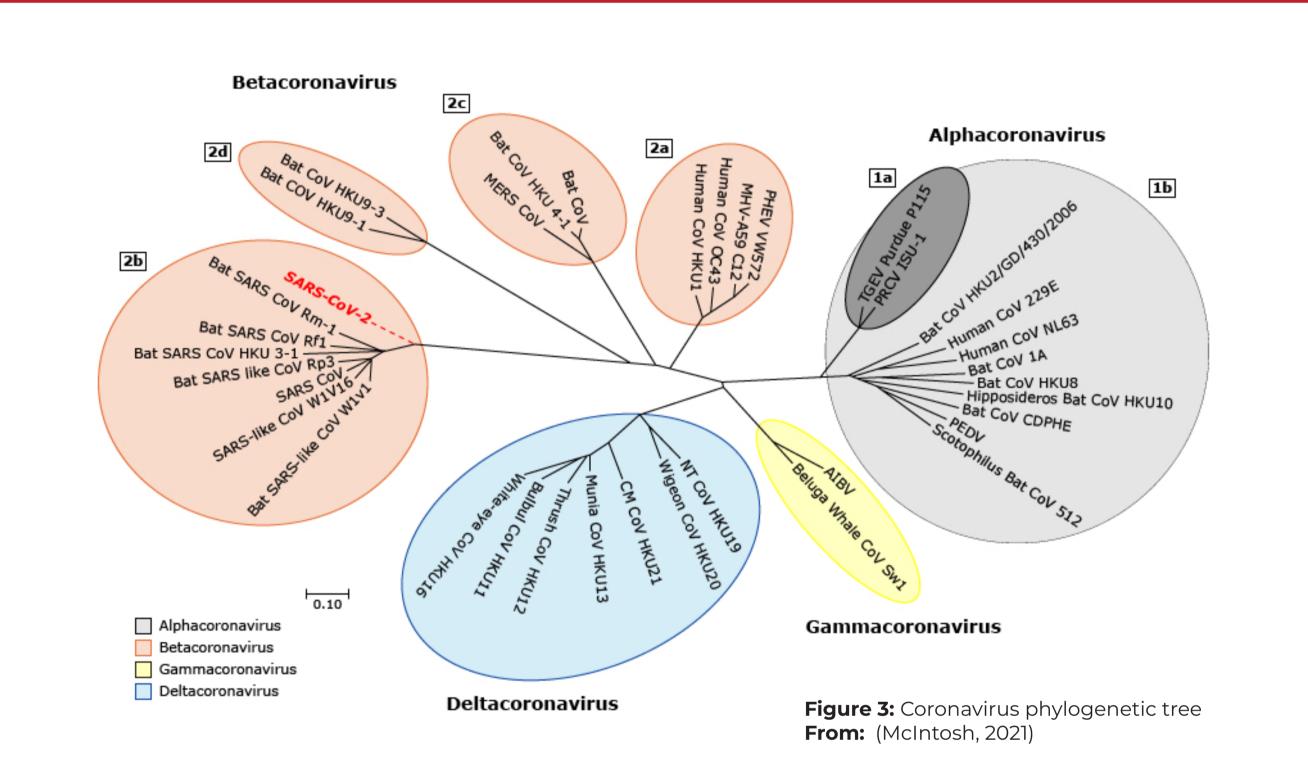
- Affected countries: 192 infected: 170 million.

Updated May 30, 2021 from Johns Hopkins University



What are coronaviruses?

- -Taxonomically: Viruses belonging to the order *Nidovirales*, family Coronaviridae and subfamily Orthocoroviringe.
- -Divided into 4 genres: Alphacoronavirus, Betacoronavirus, Gammacoronavirus and Deltacoronavirus.



Seven infectious **Coronaviruses in**

humans:

- Alphacoronavirus: HCoV-229E i HCoV-NL63.
- Betacoronavirus: HCoV-OC43. HCoV-HKU1, SARS-CoV, MERS-CoV i SARS-CoV-2.

SARS and MERS

-SARS-CoV: Start in China in 2003. Horseshoe bat reservoir. Intermediate civeta host.

-MERS-CoV: Start in Saudi Arabia 2012. Egyptian Sheath-tailed Bat Reservoir (Taphozous perforatus). Intermediate dromedary host.

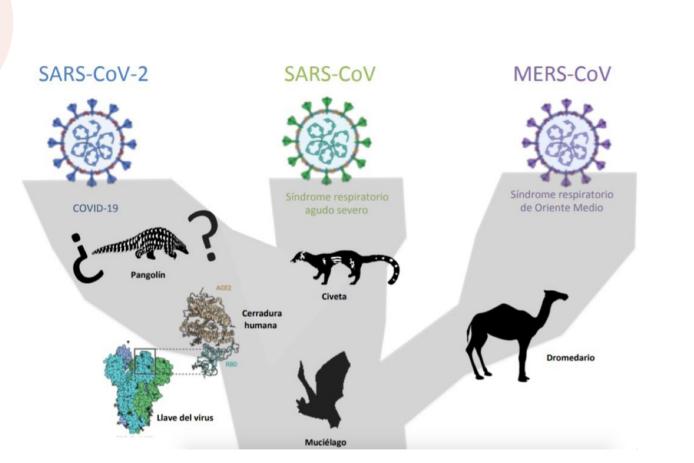


Figure 4: SARS-CoV, MERS-CoV and SARS-CoV-2 **From:** (Díez *et al.*, 2020)

SARS Cov-2 in animal species

- Cats, ferrets, hamsters, big cats, bats and non-human primates. Dogs limited susceptibility.
- Other domestic species, including pigs and poultry, do not appear susceptible.
- Susceptibility given by the sequence of 22 amino acids (ACE-2) directly involved in the ACE-2 and RBD interaction of the virus.

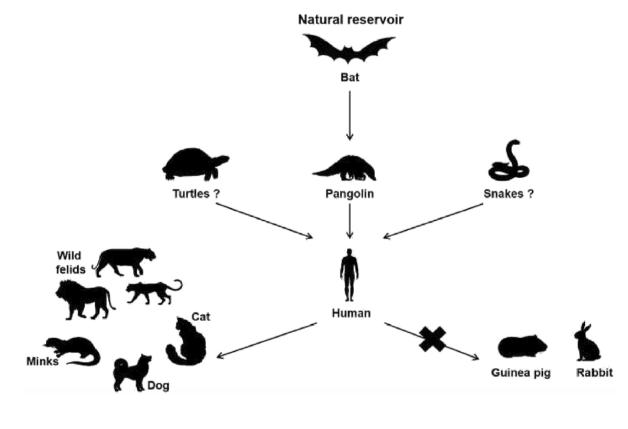


Figure 5: Susceptibility of a range of animals to natural infection by SARS-CoV-2.

7.

Hypothesis of the origin of SARS-Cov-2

- -November 2020, joint study between China and WHO.
- "Extremely unlikely" No evidence in genome sequences that occurred in a laboratory.
- "Probable to very probable" a zoonotic effect of an intermediate host.

8.

Natural origin of SARS-CoV-2

-It has circulated among the population and evolved to SARS-CoV-2.

-Zoonosis from an intermediate host.

- It probably existed in other countries such as France, India and Holland before Wuhan outbreak.

References

Figure 1: Robertson, S. (2021). Protein-based vaccines against SARS-CoV-2 may provide advantages over mRNA-based vaccines. News-Medical. Figure 2: Eckert, A., & Higgins, D. (2021). SARS-CoV-2_illustration. CDC. https://www.cdc.gov/coronavirus/2019-ncov/images/2019-nCoV-virus-illustration.jpgRS-CoV-2_illustration.png

Figure 3: McIntosh, K. (2021). Coronaviruses. UpToDate, Inc. https://www.uptodate.com/contents/coronaviruses/print

Figure 4: Díez, F., Plaza, J., Coiras, M., Primo, E., Bojo, C., Pérez-Gómez, B., Rodriguez, F., García-Carpintero, E., Sánchez, L., Álvarez, D., & Alcamí, J. (2020). Origen del

SARS-Cov-2. 1–7. https://www.conprueba.es/sites/default/files/informes/2020-05/origen-sars-cov-2.pdf

Figure 6: AVMA (2021). SARS-CoV-2 in animals. (2021). American Veterinary Medical Association All Rights Reserved. https://www.avma.org/resources-tools/ani-

SARS-CoV-2 infections in animals

SARS-CoV-2 in nonhuman animals, in addition to in people. This is a global snapshot of the eported cases of naturally acquired SARS-CoV-2 infections in animals.







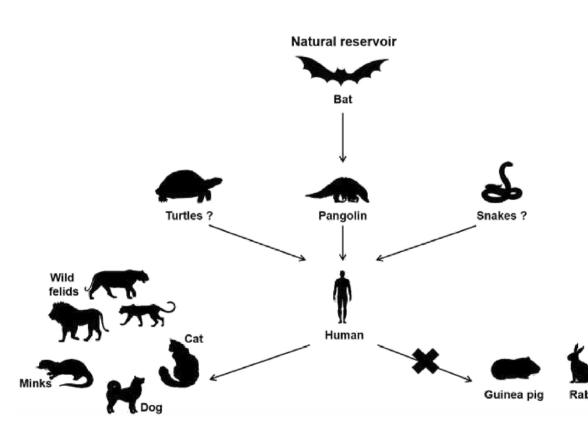




Figure 6: Inefective animals **From:** (AVNA, 2021)

Conclusions

- It is essential to know the "Source to report on global strategies to mitigate the risk of future
- 🔅 Human and animal health is interdependent and closely linked to the ecosystems in which they coexist (One Health).
- 🔅 Zoonotic origin has the bat as a reservoir because the genetic sequence has a coincidence of 96%.
- * Likeliest hypotheses of infection: - Non-pathogenic zoonotic infection that mutated in humans to SARS-Cov-2. - Pathogenic zoonotic infection already
- like SARS-CoV-2. * Androgenic SARS-CoV-2 infection, but not infected from animals to humans.
- * Nationwide News reported on May 8 that the possibility of genetic manipulation is being considered.



From: (do Vale et al., 2021)

Figure 5: do Vale, B., Lopes, A. P., Fontes, M. da C., Silvestre, M., Cardoso, L., & Coelho, A. C. (2021). Bats, pangolins, minks and other animals - villains or victims of SARS-CoV-2? Veterinary Research Communications, 45(1), 1–19. https://doi.org/10.1007/s11259-021-09787-2

mal-health-and-welfare/covid-19/sars-cov-2-animals-including-pets