

Recursos del Bibliotecario 2/8/21

- Identification of a novel coronavirus causing severe pneumonia in human: a descriptive study (Ren, et.al.):
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- A Novel Coronavirus from Patients with Pneumonia in China, 2019 (Zhu, et.al.):
<https://www.nejm.org/doi/full/10.1056/nejmoa2001017>
- Genomic characterisation and epidemiology of 2019 novel coronavirus: implications for virus origins and receptor binding (Lu, et.al.):
<https://www.thelancet.com/pdfs/journals/lancet/PIIS0140-6736%2820%2930251-8.pdf>
- Evolution and epidemic spread of SARS-CoV-2 in Brazil (Candido, et.al.):
<https://science.sciencemag.org/content/369/6508/1255.abstract>
- Genomic epidemiology of novel coronavirus - Global subsampling (The Nexstrain Team):
https://nextstrain.org/ncov/global?c=pangolin_lineage&l=clock
- Preliminary genomic characterisation of an emergent SARS-CoV-2 lineage in the UK defined by a novel set of spike mutations (Rambaut, et.al.):
<https://virological.org/t/preliminary-genomic-characterisation-of-an-emergent-sars-cov-2-lineage-in-the-uk-defined-by-a-novel-set-of-spike-mutations/563>
- Emergence and rapid spread of a new severe acute respiratory syndrome-related coronavirus 2 (SARS-CoV-2) lineage with multiple spike mutations in South Africa (Tegally, et.al.):
<https://doi.org/10.1101/2020.12.21.20248640>
- Genomic characterisation of an emergent SARS-CoV-2 lineage in Manaus: preliminary findings (Faria, et.al.):
<https://virological.org/t/genomic-characterisation-of-an-emergent-sars-cov-2-lineage-in-manaus-preliminary-findings/586>
- Early Transmission Dynamics, Spread, and Genomic Characterization of SARS-CoV-2 in Panama (Franco, et.al.):
https://wwwnc.cdc.gov/eid/article/27/2/20-3767_article
- Robust spike antibody responses and increased reactogenicity in seropositive individuals after a single dose of SARS-CoV-2 mRNA vaccine (Krammer, et.al.):
<https://www.medrxiv.org/content/10.1101/2021.01.29.21250653v1>