

SARS-CoV-2 Variant Timemaps

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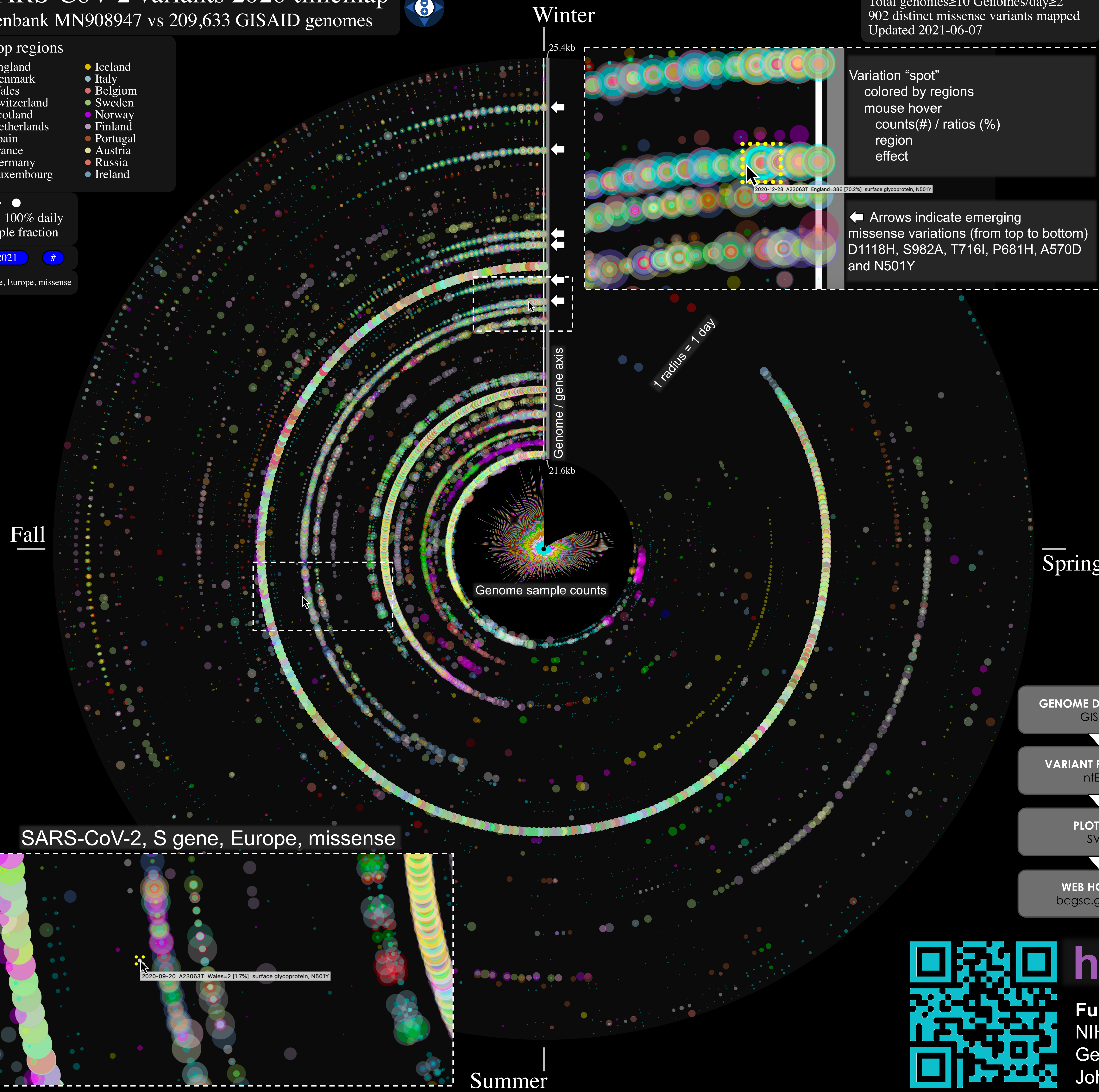
SARS-CoV-2 variants 2020 timemap
Genbank MN908947 vs 209,633 GISAID genomes

- Top regions
- England
 - Denmark
 - Wales
 - Switzerland
 - Scotland
 - Netherlands
 - Spain
 - France
 - Germany
 - Luxembourg
 - Iceland
 - Italy
 - Belgium
 - Sweden
 - Norway
 - Finland
 - Portugal
 - Austria
 - Russia
 - Ireland

1 10 100% daily sample fraction
2021 #
S gene, Europe, missense

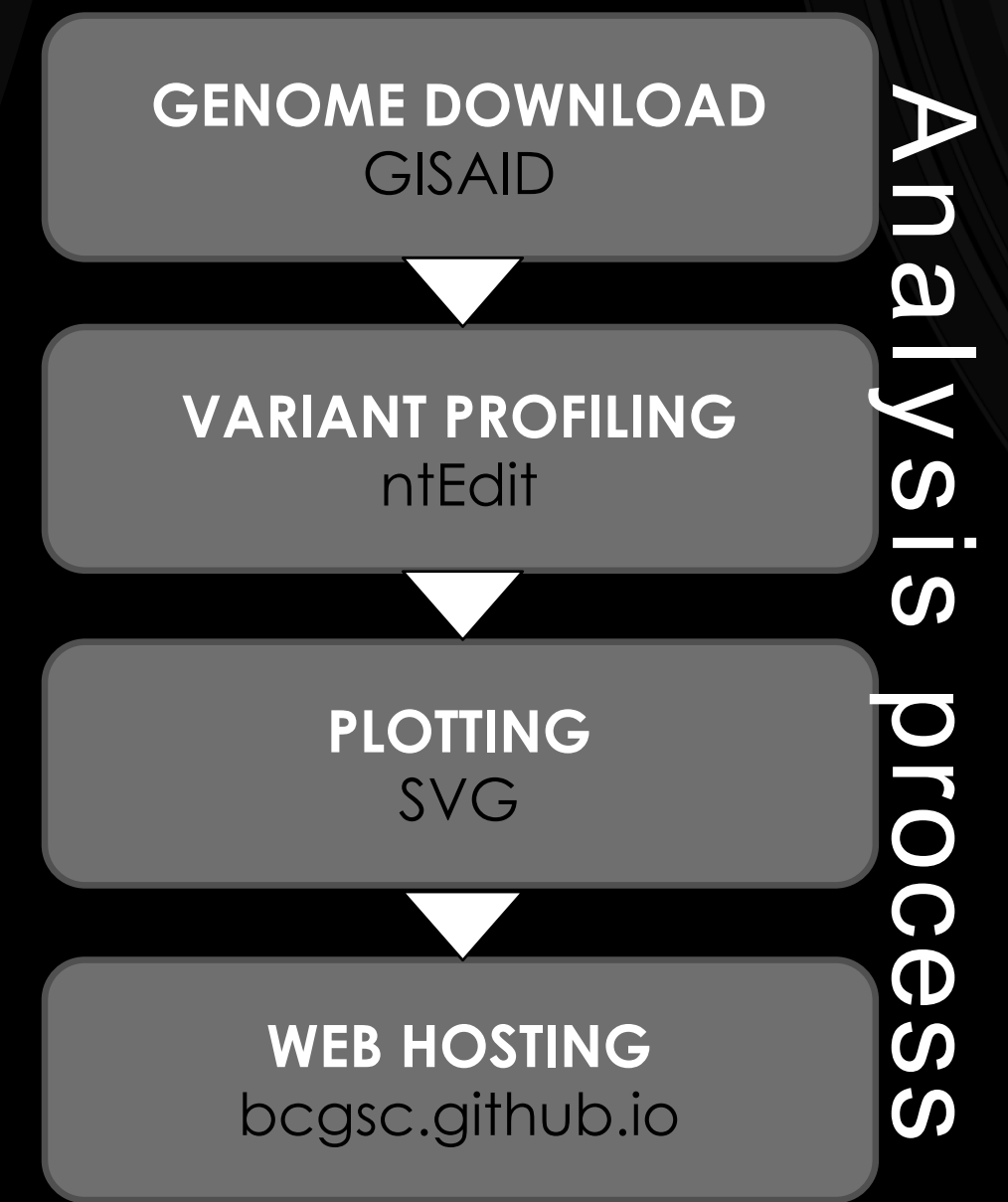
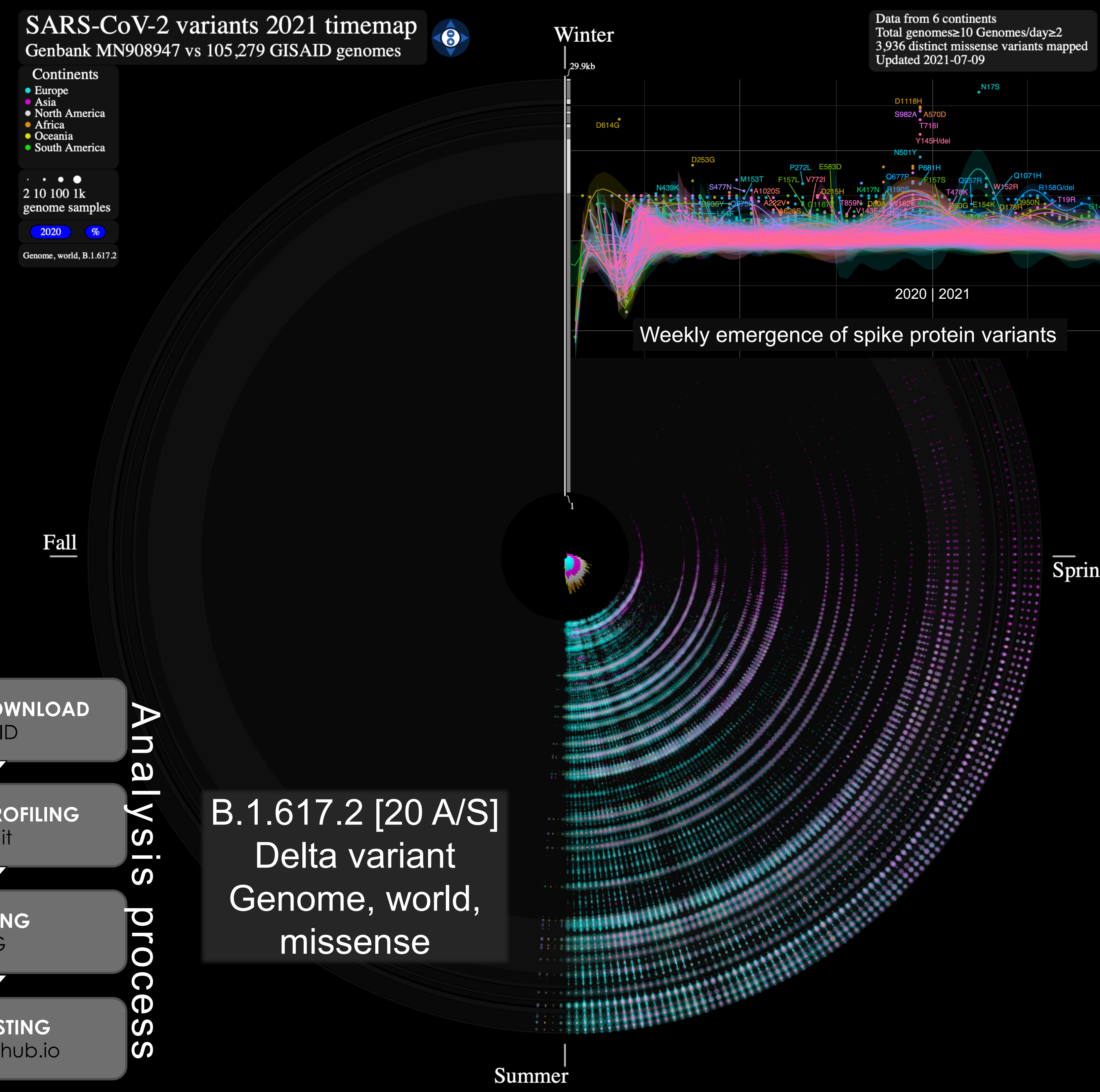
Data from 53 regions in Europe
Total genomes ≥ 10 Genomes/day ≥ 2
902 distinct missense variants mapped
Updated 2021-06-07

- Nucleotide variation in >1.6M GISAID genomes accumulating since ground zero strain
- Interactive Scalable Vector Graphics (SVG) drag, pan, zoom, tilt, highlight, hover / tooltip
- Over 120 SVG timemaps, web-hosted world / regional views genes / genome / variation types variants of concerns, WHO convention, custom
- Panorama of longitudinal strain evolution in human hosts

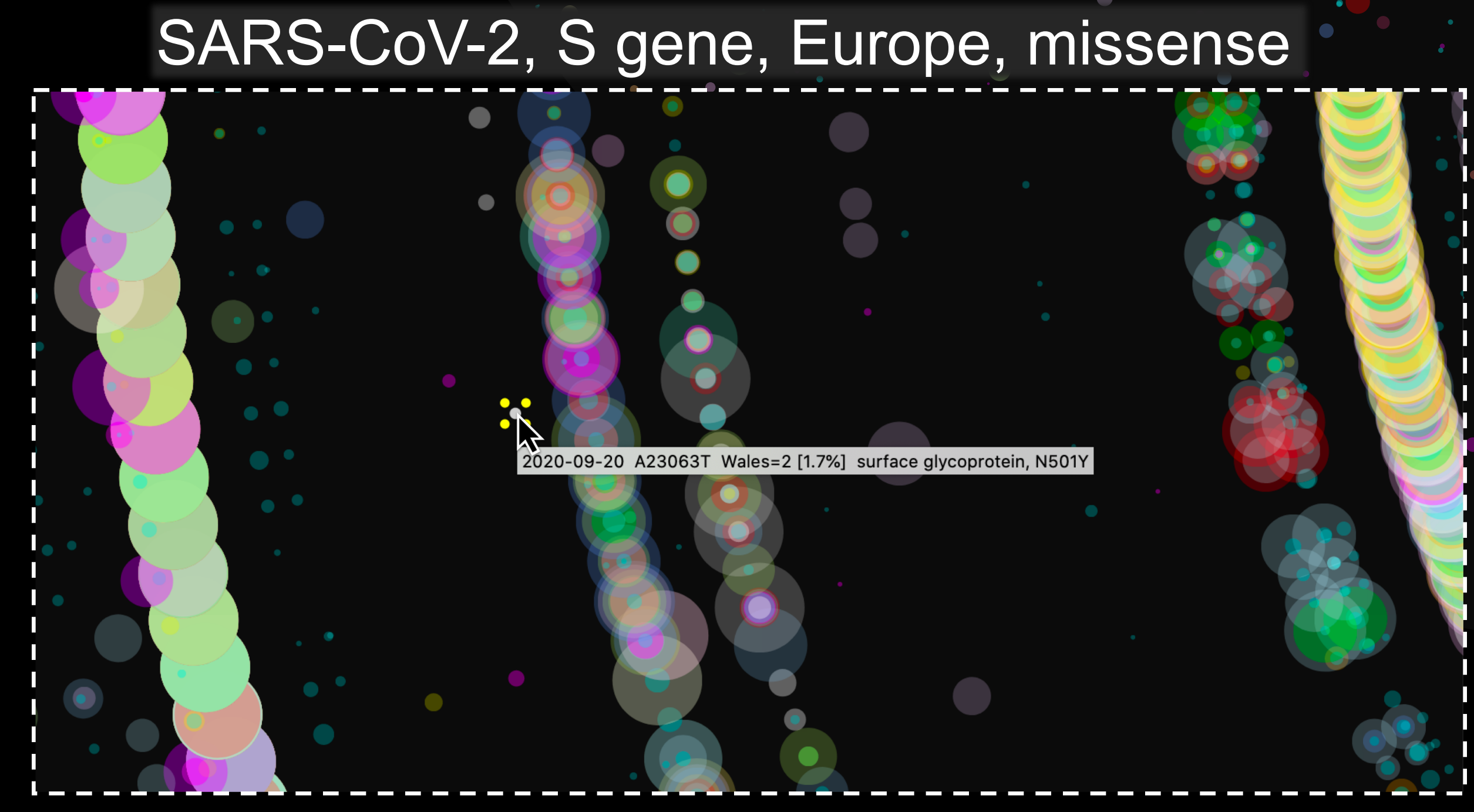


SARS-CoV-2 variants 2021 timemap
Genbank MN908947 vs 105,279 GISAID genomes

- Continents
- Europe
 - Asia
 - North America
 - Africa
 - Oceania
 - South America
- 2 10 100 1k genome samples
2021 #
Genome, world, B.1.617.2



B.1.617.2 [20 A/S]
Delta variant
Genome, world, missense



<https://bcgsc.github.io/SARS2>

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John Jambor Education Fund

<http://warrenlr.github.io>



<http://birol-lab.ca>