



Novel Coronavirus Host Genomic study - South Africa COVIGen-SA

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Outline

- COVID-19 globally and in Africa
- COVIGen-SA update
- Challenges & Next steps

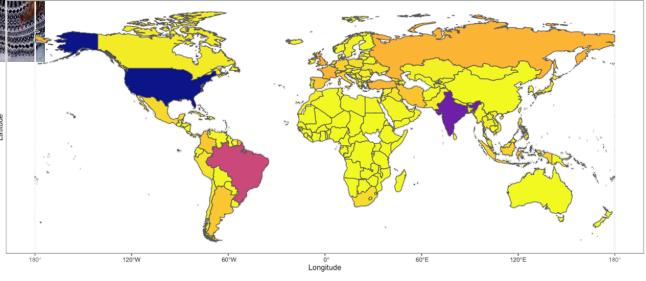


COVIGEN-SA

Coronavirus Host Genomics Study – South Africa

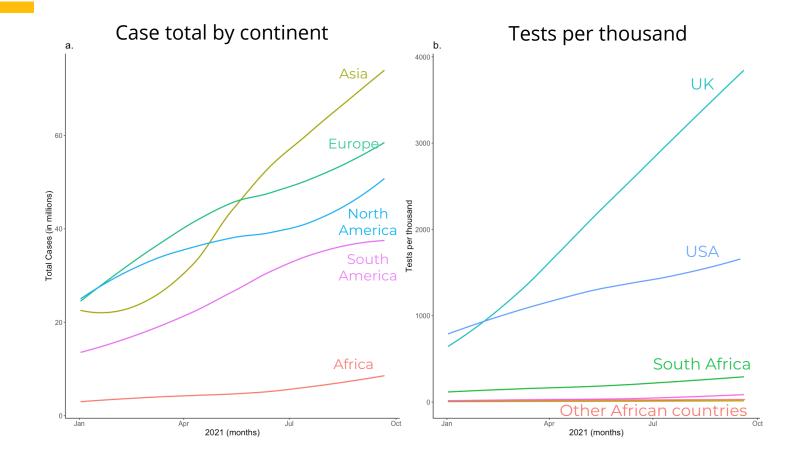
Total cases per country (as of 21 September 2021)

South Africa (28 October 2021) 2,921,114 cases 89,104 deaths



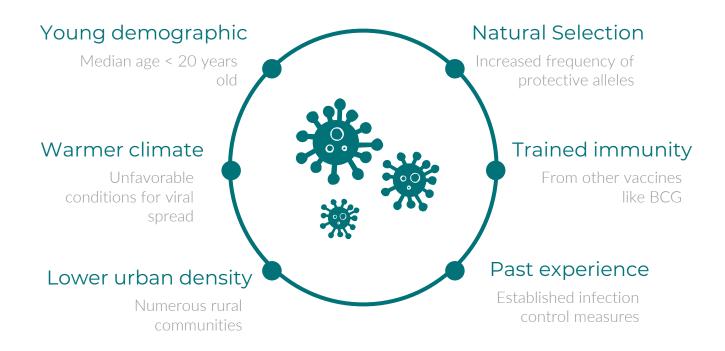
10 million







Africa's low burden





COVIGen - SA

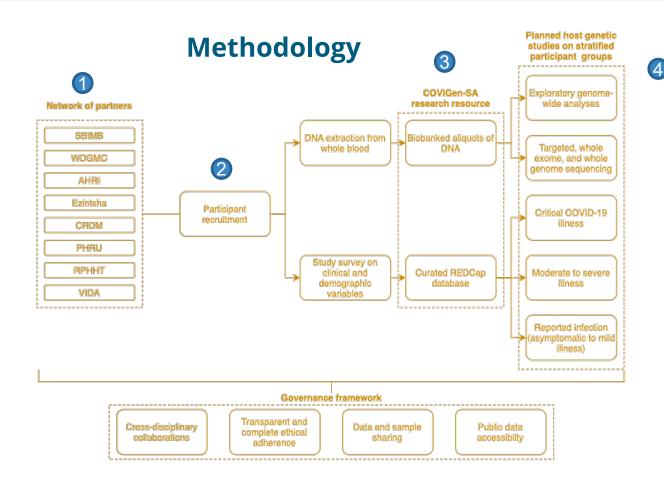
Aim

A platform for COVID-19 research that accounts for individual variability in the genetics and health backgrounds of Black South Africans, in line with a precision medicine approach.

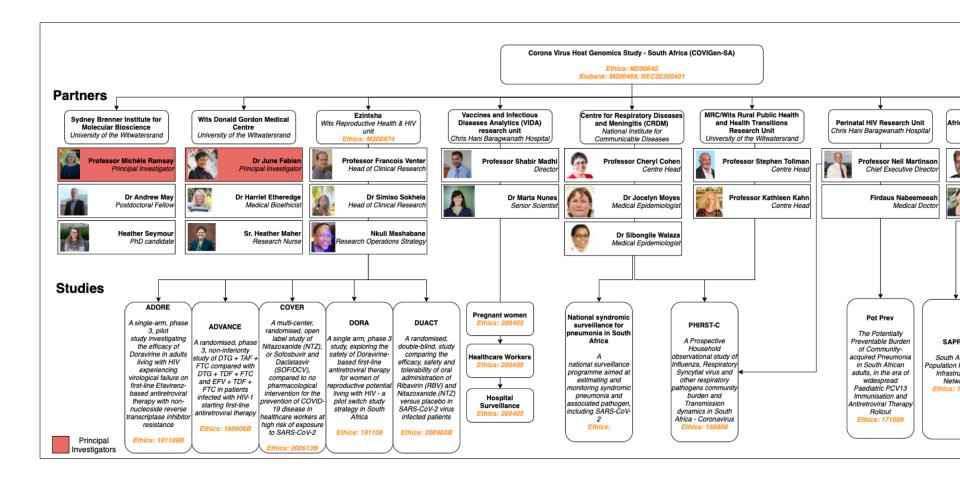
Objectives

- Establish a research resource of harmonised clinical and genetic data for a large sample of SARS-CoV-2 positive Black South Africans
- Conduct a multi-approach investigation into host genetic factors affecting COVID-19 susceptibility and severity



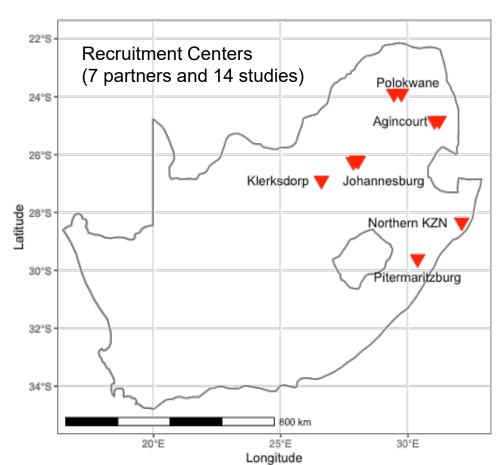






COVIGen - SA

- Total records: 1707
- SARS-CoV-2 positive: **1080** (63%)
- Severe COVID-19: 221 (20%) (supplemental oxygen, ICU admission and/or ventilation)
- Asymptomatic or mild symptoms: 859
- Genotyped: 576
- Additional funded genotyping: 576





Genotyping: Principal Component Analysis

572 individuals

1 517 137 SNPs following QC cutoffs:

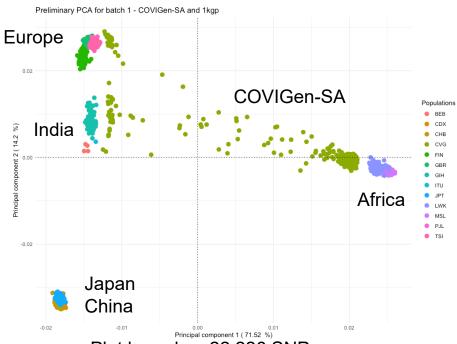
• Missingness per SNP: 0.1

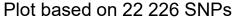
Missingness per individual: 0.1

Minor allele frequency: 0.05

 Hardy-Weinberg threshold: 0.0000001

Autosomal markers only









Challenges and Next Steps

- Integrating phenotype data
- Increasing number of severe COVID-19 cases
- Resources for further genotyping



- Focus on partnerships and recruitment
- Publish cohort paper
- Analyze preliminary data
- Partner with other host genomics studies

