



# SARS-CoV-2 sequence analysis: a worked example

Peter van Heusden <a href="mailto:pvh@sanbi.ac.za">pvh@sanbi.ac.za</a>,

South African National Bioinformatics Institute

## The UseGalaxy.Eu workflows

- Highly sensitive workflows for SARS-CoV-2 variant discovery
- Illumina & ONT platform
- Can be run on usegalaxy.eu or other Galaxy server
- Training material:
  - https://galaxyproject.eu/event/2021-06-21-sars-cov-2-data-analysis-monitoring-training/

### **Other Galaxy workflows**

- SANBI SARS-CoV-2 workflow:
  - Illumina: based on ivar toolkit
    - https://github.com/galaxyproject/iwc/tree/main/workflows/sars-cov-2-variant-calling/sars-cov-2-pe-illumina-artic-ivar-analysis

#### ARIES

- https://aries.iss.it/ Istituto Superiore di Sanità
- Illumina, Nanopore, Ion Torrent
- https://www.biorxiv.org/content/10.1101/2021.01.16.425365v2

#### **Demo**