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# SARS-CoV-2 ENA submission workflow + guidance for structuring and releasing metadata V.1

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**ABSTRACT** 

#### PURPOSE:

This workflow provides an overview on the metadata specification recommended for SARS-CoV-2 sequence data and a series of protocols outlining the steps for ENA submission.

## **PHA4GE Contextual Metadata SOP**

- This protocol provides step-by-step instructions for populating the template, and also addresses a number of
  ethical, privacy and practical considerations that should be discussed with your data steward prior to any type
  of data sharing.
- The appendices provide additional instructions and examples of how to curate sample type descriptions, and how to identify additional standardized terms should you need them.

## SOP for populating EBI submission templates (ENA)

Guidance for populating the ENA metadata template using PHA4GE fields and terms.

## SARS-CoV-2 EBI submission protocol: ENA, BioSample, and BioProject

- Step-by-step instructions for establishing a new EBI (Webin) submission account and for creating and linking a new BioProject to an existing umbrella effort.
- SARS-CoV-2 raw data submission to ENA (European Nucleotide Archive) and metadata to BioSample.

## SARS-CoV2 EBI assembly submission protocol

Required: established BioProject and BioSamples

 Submit SARS-CoV-2 assemblies (consensus sequences) to ENA linking to existing BioProject, BioSamples, and raw data.

THIS COLLECTION ACCOMPANIES THE FOLLOWING PUBLICATION

Griffiths, Emma J., Ruth E. Timme, Andrew J. Page, Nabil-Fareed Alikhan, Dan Fornika, Finlay Maguire, Catarina Inês Mendes, et al. "The PHA4GE SARS-CoV-2 Contextual Data Specification for Open Genomic Epidemiology," August 9, 2020. https://doi.org/10.20944/preprints202008.0220.v1.

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ENA

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Version 1

by Nabil-Fareed Alikhan, Quadram Institute Bioscience

ENA

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