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# SARS-CoV-2 NCBI submission workflow + guidance for structuring and releasing metadata

## Ruth E Timme<sup>1</sup>, Emma Griffiths<sup>2</sup>

<sup>1</sup>US Food and Drug Administration; <sup>2</sup>Simon Fraser University

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GenomeTrakr StaPH-B 1 more workspace

Ruth Timme US Food and Drug Administration

**ABSTRACT** 

#### **PURPOSE:**

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

## Overview of NCBI's submission process and the metadata required:

Provides an overview of the submission process and includes a brief training video.

## SARS-CoV-2 NCBI submission protocol: SRA, BioSample, and BioProject

- Step-by-step instructions for establishing a new NCBI laboratory submission account and for creating and linking a new BioProject to an existing umbrella effort.
- SARS-CoV-2 raw data submission to SRA (Sequence Read Archive) and metadata to BioSample.

## SARS-CoV-2 NCBI consensus submission protocol: GenBank

Required: established BioProject and BioSamples

Submit SARS-CoV-2 assemblies to NCBI GenBank, linking to existing BioProject, BioSamples, and raw data.

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**KEYWORDS** 

SARS-CoV-2, NCBI, submission, genomic epidemiology, SRA, Biosample

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PHA4GE contextual metadata SOP

Version 1

by Ruth Timme, US Food and Drug Administration





Version 2

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SARS-CoV-2 NCBI submission protocol: SRA, BioSample, and BioProject

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