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Version 4

Populating NCBI template for submissions using BioNumerics v7.6 V.4

DOI

<https://dx.doi.org/10.17504/protocols.io.81wgbp5bovpk/v4>



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Protocol status: Working

We use this protocol and it's working

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Last Modified: March 21, 2024

Protocol Integer ID: 97004

Keywords: NCBI submission, BioNumerics, biosample, SRA, metadata, bioproject, bionumerics for submission, submission of genometrakr, genometrakr submission, environmental isolates to ncbi, ncbi template for submission, ncbi-update submission sheet, using bionumeric, populating ncbi template, genometrakr, bionumeric, collecting isolate metadata, records to ncbi, genometrakr network management, familiar with the mandatory metadata field, isolate metadata, ncbi, environmental isolate, sequencing record, mandatory metadata field, sequencing, standardized procedure,

Abstract

PURPOSE: to define the standard operating procedure for collecting isolate metadata using BioNumerics for submission of food/environmental isolates to NCBI.

SCOPE: to provide a standardized procedure to collect isolate metadata using BioNumerics for submission of food/environmental isolates to NCBI.

RESPONSIBILITIES- SOP Responsible Officials: Ruth Timme, Maria Balkey

The GenomeTrakr Network Management will be responsible to monitor GenomeTrakr submissions processed through Bionumerics and ensure that all GT labs are familiar with the mandatory metadata fields required for submission of GenomeTrakr sequencing records to NCBI.

V3: Dropdown menus from controlled vocabulary added to the ncbi_update submission sheet



New Version for this protocol

- 1 Visit the new version of this protocol by clicking the following link:
<https://www.protocols.io/view/populating-ncbi-template-for-submissions-using-bio-3b9l4qn4ovo5/v1>