

Mar 20, 2024

Version 3

Populating NCBI template for submissions using BioNumerics V.3

DOI

<https://dx.doi.org/10.17504/protocols.io.36wgg3ejylk5/v3>



Ruth Timme¹, Maria Balkey¹, Julie Haendiges¹, Brian Sauders², Tina Lusk Pfefer²

¹US Food and Drug Administration; ²New York State Department of Agriculture & Markets

GenomeTrakr

Tech. support email: genomeTrakr@fda.hhs.gov



Maria Balkey

US Food and Drug Administration

Create & collaborate more with a free account

Edit and publish protocols, collaborate in communities, share insights through comments, and track progress with run records.

Create free account

OPEN  ACCESS



DOI: <https://dx.doi.org/10.17504/protocols.io.36wgg3ejylk5/v3>

Protocol Citation: Ruth Timme, Maria Balkey, Julie Haendiges, Brian Sauders, Tina Lusk Pfefer 2024. Populating NCBI template for submissions using BioNumerics . **protocols.io** <https://dx.doi.org/10.17504/protocols.io.36wgg3ejylk5/v3> Version created by **Maria Balkey**



License: This is an open access protocol distributed under the terms of the **Creative Commons Attribution License**, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited

Protocol status: Working

We use this protocol and it's working

Created: March 20, 2024

Last Modified: March 20, 2024

Protocol Integer ID: 97005

Keywords: NCBI submission, BioNumerics, biosample, SRA, metadata, bioproject, bionumerics for submission, submission of genometrakr, environmental isolates to ncbi, ncbi template for submission, genometrakr submission, ncbi-update submission sheet, using bionumeric, populating ncbi template, collecting isolate metadata, genometrakr, bionumeric, records to ncbi, genometrakr network management, isolate metadata, familiar with the mandatory metadata field, ncbi, sequencing record, environmental isolate, 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-3byl4qn4ovo5/v1>