

SRA

☒ Update my contact information in profile

## General Information

Required fields are marked with \* asterisk

### BioProject

i BioProject describes the goal of your research effort.

\* Did you already register a BioProject for this research, e.g. for the submission of the reads to SRA and/or of the genome to GenBank?

☒ Yes ☐ No

\* Existing BioProject

PRJNAXXXXXX **Input BioProject ID**

### BioSample

i The BioSample records the detailed biological and physical properties of the sample that was sequenced. A BioSample can be used in more than one BioProject since it should be used for all the data that were obtained from that sample. Usually SRA data sets are generated from more than one sample.

\* Did you already register a BioSample for this sample, e.g. for the submission of the reads to SRA and/or of the genome to GenBank?

☒ Yes ☐ No

**Select**

### Release date

i **Note:** Release of BioProject or BioSample is also triggered by the release of linked data.

**Select**

\* When should this submission be released to the public?

☒ Release immediately following processing  
☐ Release on specified date or upon publication, whichever is first

i Please allow 24-48 hours for propagation of the data to the NCBI SRA public site.

Continue

Click

## SRA Metadata

Required fields are marked with \* asterisk

i For more detailed help with SRA submission please read the [SRA Submission Wizard Help](#).

\* How do you want to provide your metadata?

☐ Use built-in table editor  
☒ Upload a file using Excel or text format (tab-delimited)

**1. Select**

\* Metadata file **2. Click and Upload**

or drag and drop it here

i [Download Excel spreadsheet](#) (designed to make it easier to select the correct metadata values), edit, save and then upload the modified Excel file.

Continue

3. Click

## Files

Required fields are marked with \* asterisk

- i
- Each file must be listed in the [SRA metadata table you uploaded](#). If you are uploading a tar archive, list each file name, not the archive name.
  - Unique file names that do not contain any sensitive information should be used for all files. File names as submitted appear publicly when data is retrieved from the cloud.
  - Files can be compressed using **gzip** or **bzip2**, and may be submitted in a **tar** archive, but archiving or compressing your files is not required. **Do not use zip!**

**1. Select**

\* How do you want to provide files for this submission?

☒ Web browser upload via HTTP or Aspera Connect plugin  
Do not use web browser HTTP upload if you are uploading files over 10 GB or more than 300 files.  
☐ FTP or Aspera Command Line file preload  
All files for a submission must be uploaded into a single folder.  
☐ AWS or GCP bucket

\* Files **2. Click and Upload**

or drag and drop them here

☐ Autofinish submission

i Try to complete the submission automatically. Basic checks will be done on the files in the background. If there are no errors, this submission will be submitted for you. If the checks fail, you will get email notification to come back to fix the problems.

EN

Continue

3. Click and Review  
the following page