

https://www.ncbi.nlm.nih.gov/sra/docs/submitbio/

An official website of the United States government [Here's how you know](#)

NIH National Library of Medicine
National Center for Biotechnology Information

SRA SRA Advanced

Getting Started Submission Quick Start Search and Download SRA in the Cloud Archive Documentation

Submit your project and biological samples

- [Introduction](#)
- [BioProject submissions guidelines](#)
- [BioSample submissions guidelines](#)
- [Updating BioProject and BioSample data](#)
- [Contact SRA staff](#)

Introduction

Go to

Go to:

Registering project and biological samples at the NCBI BioProject and the BioSample databases is a prerequisite for any public SRA submission. The BioProject and BioSample databases store data that relate to organizational and biological aspects of sequencing experiments.

Getting Started

- [Getting Started](#)

Submitting Data

General

- [Quick Start](#)
- [BioProject](#)
- [File Format](#)
- [SRA Metadata](#)
- [SRA File Upload](#)
- [Frequently Asked Questions](#)

SRA Submission

- [Submitting](#)

NCBI Requires Bioproject and Sample Submissions for High Throughput Sequencing submissions (SRA)

Submission Portal

Home **My submissions** Manage data Groups Templates My profile

BioProject submission: SUB9611187

Delete submission

polar bear Metagenome

1 SUBMITTER 2 PROJECT TYPE 3 TARGET 4 GENERAL INFO 5 BIOSAMPLE 6 PUBLICATIONS 7 REVIEW & SUBMIT

General Info

Required fields are marked with * asterisk

Release date

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

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

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

*** Project title** ?

polar bear Metagenome

*** Public description** ?

polar bears have recently been discovered in the vicinity of Lilliput Hydrothermal Vent Field. We collected samples with the aim of exploring evolutionary divergence.