

Submission Type

★ How do you want to submit your data?

☒ Single genome ← **click here**

Manually complete a web form to describe one genome assembly and to upload its sequences.

☐ Batch/multiple genomes

Upload the "Genome Info" file ([download template ahead of time](#) or during submission), a tab-delimited text file that describes each of your genome assemblies and their attributes/metadata, plus the genome sequences (one file per genome).

Information that must be common to all genomes in the batch are:

- BioProject
- (initial) release date
- assembly type (either WGS or non-wgs, not a mix of both types)
- file type (FASTA or SQN)
- gap/Ns details
- publication information (for FASTA submissions only)
- PGAP request status (Yes/No; for prokaryotic genomes only)
- [See more details here](#)

Continue

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