

Pipelines

A pipeline is a series of tasks used to process and analyze genomic data. Pipelines are built and managed under the *Analyses* tab of any project in Partek Flow. Additional information on pipelines is given in the [Creating and Analyzing a Project](#) documentation and there is further guidance on how to build analysis pipelines for different kinds of data on the [tutorials](#) page.

The combination of circular data nodes and rectangular task nodes displayed under the *Analyses* tab are herein collectively referred to as the *task graph*.

This user guide will cover the following topics:

- [Making a Pipeline](#)
- [Running a Pipeline](#)
- [Downloading and Sharing a Pipeline](#)
- [Previewing a Pipeline](#)
- [Deleting a Pipeline](#)
- [Importing a Pipeline](#)

Pipelines can also imported and managed on the [Pipeline managment](#) page.

Additional Assistance

If you need additional assistance, please visit [our support page](#) to submit a help ticket or find phone numbers for regional support.



Your Rating: ☆☆☆☆☆ Results: ★★★★★ 38 rates