

Library File Management

The library file management tool in Partek® Flow® provides an easy way to create, process and manage reference sequences, cytoband files, annotation models, aligner indexes, gene sets, variant databases and microarray probe sequence files.

For a quick start, please watch the short videos below:

**Quickly setup library files for the hg19 human reference genome assembly.
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**Adding library files for a non-model organism
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You can watch the [full webinar on our website](#).

The process can be adapted to the different kinds of file types.

This user guide will cover the following topics:

- [Library File Management Settings](#)
- [Library File Management Page](#)
- [Selecting an Assembly](#)
- [Library Files](#)
- [Update Library Index](#)
- [Creating an Assembly on the Library File Management Page](#)
- [Adding Library Files on the Library File Management Page](#)
- [Adding a Reference Sequence](#)
- [Adding a Cytoband](#)
- [Adding Reference Aligner Indexes](#)
- [Adding a Gene Set](#)
- [Adding a Variant Annotation Database](#)
- [Adding a SNPEff Variant Database](#)
- [Adding a Variant Effect Predictor \(VEP\) Database](#)
- [Adding an Annotation Model](#)
- [Adding Aligner Indexes Based on an Annotation Model](#)
- [Adding Library Files from Within a Project](#)
- [Microarray Library Files](#)

Additional Assistance

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



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