

Library File Management Page

To access the library file management page click the avatar in the top right corner and choose **Settings**. Then under **Components** on the left, click **Library files**.

The library files page has three tabs - *Genomic library files*, *Prep kits files*, and *Other library files*. This section of the user guide will focus on the *Genomic library files* tab, which is relevant for next-generation sequencing analysis (Figure 1).

Genomic library files **Prep kit files** **Other library files**

Assembly

Homo sapiens (human) - hg38



▼ **Reference files**

Library file	Owner	Ignore	Actions
Reference sequence	Administrator	☐	
Cytoband	Administrator	☐	

▼ **Reference aligner indexes**

[+ Add reference aligner indexes](#)

Library file	Owner	Ignore	Actions
Bowtie index	Administrator	☐	
Bowtie 2 index	Administrator	☐	
HISAT2 2.1.0 index	Administrator	☐	
BWA index	Administrator	☐	
STAR 2.7.8a index	Administrator	☐	

> **Annotation models**

> **Gene set databases**

> **Variant annotations**

> **SnEff variant databases**

> **microRNA targets maps**

Figure 1. Genomic library files tab

The **arrows** (> /) expand/collapse each section. Associated library files are shown in a table in each section. In the *Actions* column of each table, the **view**  icon displays additional library file details, the **red bin** () icon dissociates a library file. The hourglass () icon indicates a library file is being created.

Additional Assistance

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

[« Library File Management Settings](#) [Selecting an Assembly »](#)



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