

Navigation

The File System:

+ New

- Create a new folder
- Upload a new file from your device
- Upload a new file from a url

Home

- For files you own

Shared with me

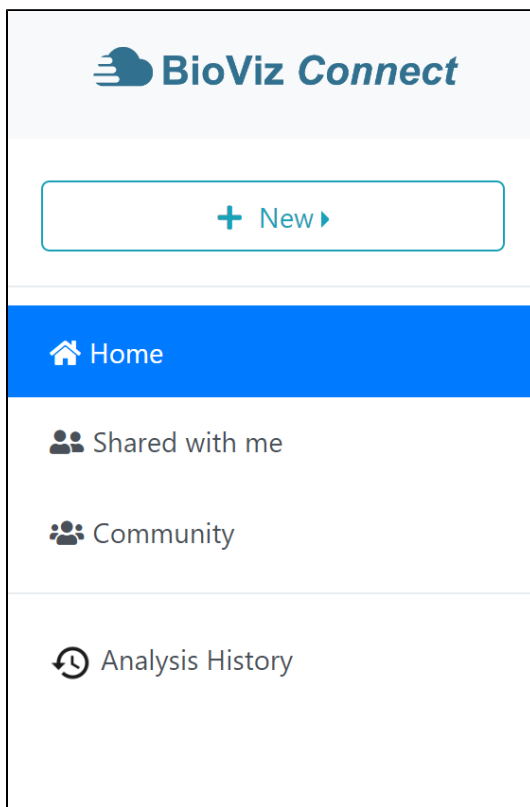
- Files made public to you

Community

- Files public to everyone

Analysis History

- Files created from the analyses you have run



Main panel:

Navigate folders by double clicking on a folder or by clicking on the folder name breadcrumbs at the top of the window. The forward and backward buttons in your web browser can also be used to navigate between folder levels.

+ New ▶

Home

Shared with me

Community

Analysis History

© UNC Charlotte, all rights reserved

Visualization Tools	Name ^	Size ^	Last Modified ^
	analyses	-	Jun 17, 2020
	LocalQuickload	-	Mar 12, 2021
	my_quickload	-	Mar 2, 2021
	release_test	-	Nov 17, 2022
	example.txt	4 Bytes	Nov 10, 2022

Page 1

Right Click options:

Copy Name

- Save the name of the file to your clipboard.

Rename...

- Edit the name of your file.

Cut

- Hold your file or folder in the clipboard

Paste

- Move your file to this location

Delete

- Delete the file

Download

- Download the file

Mange Link

- Opens a panel allowing you to make your file private or public.
- If public, a link will become visible.

Metadata

Copy Name

Rename...

Cut

Paste

Delete

Download

Manage Link

Metadata

Analyse

- Opens a panel allowing you to set:
 - IGB related species and genome version of your file
 - Track foreground and background color
 - Track name shown within IGB
 - Additional comments

Analyse

- Opens a panel showing any available analyses. Three types are currently available:
 - Bam files
 - coverage graph
 - scaled coverage graph
 - Fasta files
 - Find enriched sequence motifs