Local Quickload

General Function Checklist

This page tests local QuickLoads using relative file paths.

(To test absolute file paths see QuickLoad Saver release testing documentation)

Add files to your computer

- 1. Download the following file: relative space.zip
- 2. Extract it to where you would like your local QuickLoad to be stored. (Desktop, C drive, etc.)
- Files were extracted to local machine:
- mac
- linux
- windows

Add the QuickLoad to IGB

- 1. Open the Data Sources tab in Preferences.
- 2. Click Add... to open the Add Data Source window.
- 3. Click Choose local folder.
- 4. Choose the unzipped QuickLoad folder "relative space".
- 5. Click Submit.

Data Sources

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z	IGB Quickload	Quickload	http://igbquickload.org/quickload/								
2	Pollen	Quickload	http://www.igbquickload.org/pollen/								
Ż	Cytokinin	Quickload	http://www.igbquickload.org/cyto/								
Ż	Blueberry	Quickload	http://www.igbquickload.org/blueberry/								
Ż	UCSC	DAS	http://genome.cse.ucsc.edu/cgi-bin/das/								
ţ,	UofUtahBioinfoCore	DAS2	http://bioserver.hci.utah.edu:8080/DAS2DB/								
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Name:	Your server name	
Type:	Quickload	0
URL:		
		Choose local folder
	Cancel	Submit

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• The data on your QuickLoad is reachable by IGB and not highlighted in Red, Yellow, or Greyed out.

mac

linux

windows

Access local QuickLoad data

- 1. Close the **Preferences** window.
- 2. Open the $\ensuremath{\textbf{E_unicornis_Jul_2043}}$ genome.
- The correct **Species** (E_unicornis) and **Genome Version** (Jul_2043) loaded.
- mac
- linux
- windows

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• Data loads automatically and appears the same as the above image.

mac

linux

windows

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• Your QuickLoad appears in the Available Data section.

linux
windows

Zoom to the following coordinates, then click Load Sequence: chrXVI:387,629-387,744



- The sequence data loads along the coordinates track.
- mac
- linux
- windows