## Sharing data using QuickLoad sites

## About IGB QuickLoad

Use IGB QuickLoad (QL) to share data files on-line using a Web site or cloud storage service like Dropbox. Using IGB QuickLoad, you can set up and configure data sets so that they'll be listed in the IGB **Data Access** panel, making it easy for lab members, collaborators, and members of the public to browse and load your data as new tracks in IGB.

QuickLoad sites can be local or remote. Typically, you'll create a QuickLoad site on your local computer, test it there, and once you're happy with how the data look when loaded into IGB, you'll transfer it to a Web site.

## How to set up an IGB QuickLoad site

First, find out if your genome is already supported by the main IGB QuickLoad site.

If your genome is already there, then you should make a satellite QuickLoad site. A satellite site should not contain sequence data or gene models annotations as these are already available from the main IGB QuickLoad site.

To find out if your genome is already supported, look for it in the the main IGB QuickLoad site's contents.txt file.

IGB genome version names 	contents.txt
H_sapiens_Dec <sup>2</sup> 2013 H_sapiens_Feb_2009 H_sapiens_Mar_2006 H_sapiens_May_2004 G_gallus_Nov_2011 G_max_Aug_2010 Glycine G_max_Jan_2014 Glycine These appear in the "genome version" mer in the IGB Current	max Gmax_275_v2.0
Sequence tab	

If your genome is listed, go to: Create a satellite IGB QuickLoad site for genomes already supported in the main IGB QuickLoad.

If your genome is not listed, go to: Sharing data for a new or custom genome not already part of IGB QuickLoad.