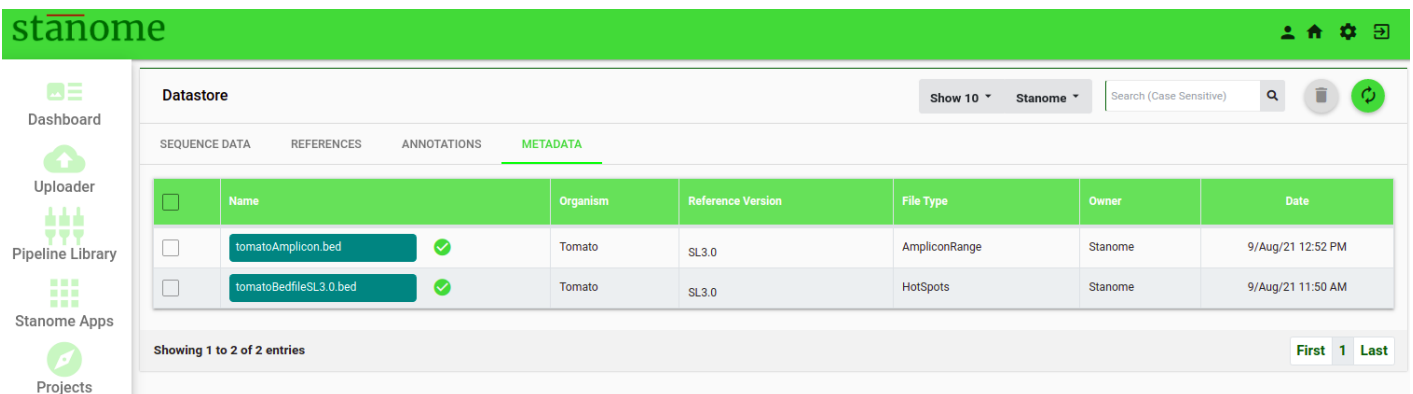# Metadata

Click the **Metadata** tab after clicking the icon to access the **Metadata** window (Fig.). The **Metadata** window shows the list view of Gene lists, Hotspots and Amplicon ranges uploaded by stanome or the user. These files are used as metadata during analysis. The window is restricted to display 10 (or more) Metadata files per page. The search box (Fig.) on the top right corner of the page allows users to filter files with name(s), type (i.e. gene list or hotspots), platform, or organism name. Metadata files are also checked for format and file integrity and their correspondence to a reference genome version. Only the validated files are available for consumption on the platform, and the remaining will be greyed out with the "Validation Failed" message.

The list view shows following six fields :

- **Name:** Filename
- **Organism:** Name of the organism to which metadata belongs to
- **Reference Version:** Reference version to which metadata file correspond to
- **FileType:** Type of the file e.g. Gene list, Hotspots, or Amplicon range
- **Owner:** Owner of the metadata file
- **Date:** Date of upload