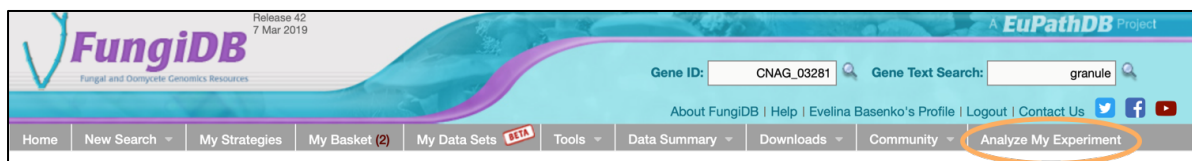


EuPathDB Galaxy workspaces - Introduction

EuPathDB Galaxy-based workspace offers pre-loaded genomes, private data analysis and display, and the ability to share and export analysis results. Developed in partnership with Globus Genomics and accessed through the *Analyze My Experiment* tab on the home page of FungiDB, the EuPathDB Galaxy instance houses a large variety of bioinformatics tools. Users can create their own private Galaxy workspace by creating a Globus account through FungiDB or any other EuPathDB database. After an account is created, sample workflows can be imported for editing, new custom workflows can be created which can be stored privately or shared with the wider community or between EuPathDB Galaxy users. Results can be exported or shared with collaborators, but the instance is not meant as a repository for long term data storage.



To use EuPathDB Galaxy services click on the *Analyze my Experiment* tab and follow prompts to set up a free account.

The workspace has four major components:

- the left panel has a list of available tools;
- the main welcome page is the interactive interface that houses pre-configured workflows, workflows editor, *etc.*;
- the top menu controls the main interface;
- the right panel provides access to histories, deleted datasets, and other useful functions.

