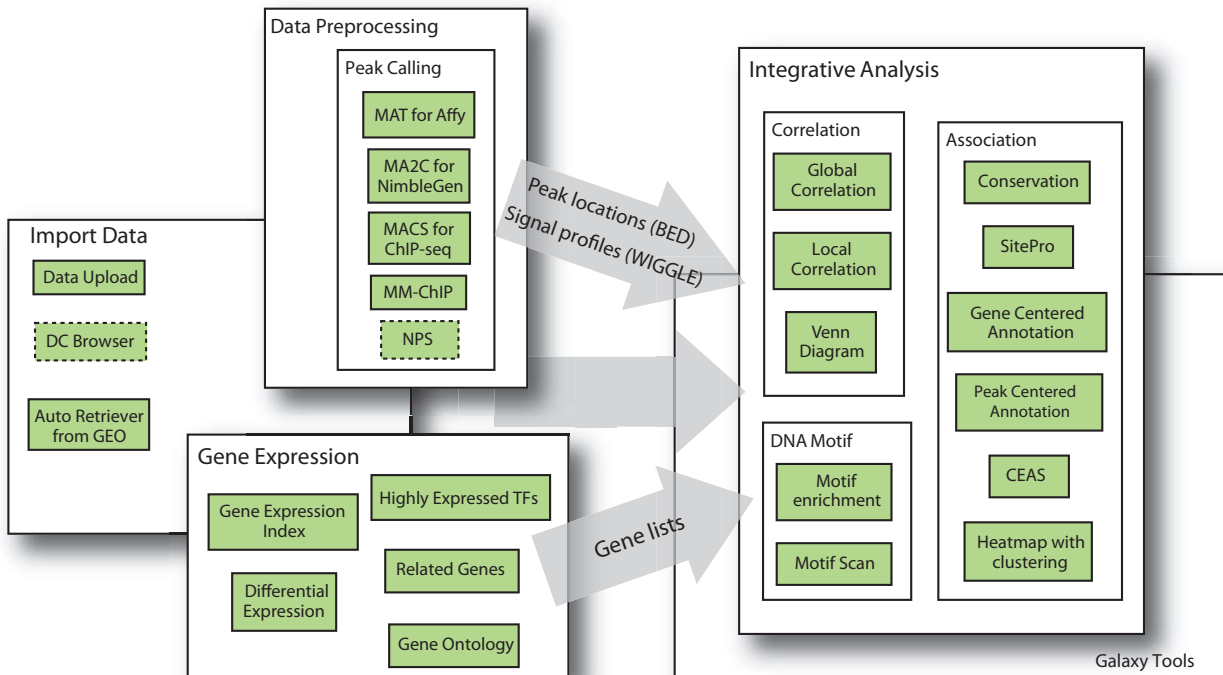




Galaxy / Cistrome Analyze Data Workflow Shared Data Lab Admin Help User

Tools Options

CISTROME TOOLBOX

- Import Data
- Data Preprocessing
- Gene Expression
- Integrative Analysis
- Liftover/Other

GALAXY TOOLBOX

- Get Data
- Text Manipulation
- Filter and Sort
- Join, Subtract and Group
- Convert Formats
- Extract Features
- Fetch Sequences
- Operate on Genomic Intervals
- Statistics
- Wavelet Analysis
- Graph/Display Data
- Regional Variation
- Multiple regression
- Multivariate Analysis

GALAXY NGS TOOLBOX BETA

- NGS: SAM Tools
- NGS: RNA Analysis
- Workflows

Welcome to Galaxy/Cistrome!

Thanks to your support and test for our Cistrome AP site, which is based on Galaxy, a Metaserver for integrative analysis of genomic data. Please check our project trac site [here](#) for more information.

Server Maintenance Policy

In order to provide a stable system, we schedule periodical server maintenance on every Saturday from 6:00pm to 8:00pm EST. During maintenance, we will:

1. Update Cistrome server codes if necessary.
2. Clean unused and unlinked data files older than 4 weeks.
3. Delete inactive user accounts older than 8 weeks.
4. Clean ASPERA directory for file upload, and reset ASPERA password. So please contact us again if necessary.

Due to the limit on the server, jobs running on Cistrome server have the following restrictions:

1. Only 15 jobs can be run simultaneously. Other jobs will be put into queue.
2. Any job generating over 10 Giga Bytes output will be stopped by Galaxy.
3. Cistrome will check every hour for jobs running over 24hrs (after the job actually runs on the server), then terminate them. If your job is terminated in this way, the output will appear green and empty. We will improve this feature later and provide more information in the job info.

Cistrome Analysis Pipeline (AP) Module

The Cistrome AP module is to organize the analysis tools in a pipeline for

History Options

summits by peak height

- 108: Sort on data 102
- 107: Convert on data
- 106: ceas job log
- 105: CEAS report
- 104: MACS diagnosis
- 103: MACS job log on
- 102: MACS wiggle on
- 101: MACS xls on
- 100: MACS summits on
- 99: MACS peaks on
- 69: Downregulated genes in group1 vs group0