

Managing disk quotas of Galaxy account

a) After logging in, click ‘User’ on the control panel and find ‘Histories’.

The screenshot shows the Deng Lab Galaxy interface. The 'User' menu is open, showing options like 'Logout', 'Datasets', 'Histories', 'Histories shared with me', 'Pages', 'Workflow Invocations', and 'Visualizations'. The 'Histories' page is displayed, showing a list of datasets with columns for Name, Items, Datasets, Tags, Sharing, Size on Disk, Created, Last Updated, and Status. The 'build' dataset is highlighted as the 'current history'.

b) On ‘Saved Histories’ page, click ‘Advanced Search’.

The screenshot shows the 'Saved Histories' page. The 'Advanced Search' button is highlighted with a red arrow. The table below shows the list of saved histories.

Name	Items	Datasets	Tags	Sharing	Size on Disk	Created	Last Updated	Status	
mem2021_Test8	223	189	32	2	2.3 GB	Apr 12, 2021	Nov 10, 2023		
build	488	286	134	68	1	23.6 GB	Oct 09, 2020	Oct 27, 2023	current history
Unnamed history	27	16	9	2	3.6 MB	Jan 06, 2022	Apr 20, 2023		
Paper_ENG_1	294	200	48	48	17.7 GB	Mar 11, 2020	Dec 17, 2021		
upgrade_test	14	10	4	4	912.1 MB	Dec 02, 2021	Dec 02, 2021		
CX 18s third	176	135	3	38	15.5 GB	Nov 14, 2018	Sep 23, 2021		
new	30	26	4	4	1.5 GB	May 09, 2021	May 26, 2021		
old_test_data	22	13	9	9	2.1 GB	Apr 12, 2021	Apr 14, 2021		
Near-Far18	14	14			1.7 MB	Mar 29, 2019	Mar 31, 2019		

c) See all histories of all statuses. Only those histories with the status ‘deleted permanently’ have been removed from disk storage and excluded from your quotas. Those only marked as ‘deleted’ will not display in the history set. Tick the box and click ‘Delete Permanently’ will remove the corresponding history from disk storage for good.

!!! Make sure you have copies of your important files. ‘Delete Permanently’ is non-reversible.

Tools

search tools

Upload Data

Upload

Amplicon sequence analysis tools

Functional genes sequences analysis tools

Statistics analysis tools

Ecological process analysis

16S GCN Prediction

Tax4Fun2

PICRUSt2

FunGuild

FAPROTAX

BugBase

LEfSe

Source Tracker

Translation and Clustering

Niche width analysis

Miscellaneous

WORKFLOWS

All workflows

Protein Cluster - Translation and

Saved Histories

Close Advanced Search

name:

tags:

sharing: accessible | all | private | public | shared

status: active | all | deleted

Name	Items	Datasets	Tags	Sharing	Size on Disk	Created	Last Updated	Status	
mem2021_Test8	223	189	32	2	2.3 GB	Apr 12, 2021	Nov 10, 2023		
training data	37	37			0 bytes	Nov 07, 2023	Nov 10, 2023	deleted permanently	
Unnamed history	0				0 bytes	Apr 04, 2023	Oct 31, 2023	deleted	
build	488	286	134	68	1	23.6 GB	Oct 09, 2020	Oct 27, 2023	current history
Unnamed history	27	16	9	2	3.6 MB	Jan 06, 2022	Apr 20, 2023		
Paper_ENG_1	294	200	46	48	17.7 GB	Mar 11, 2020	Dec 17, 2021		
Unnamed history	0				0 bytes	Dec 09, 2021	Dec 09, 2021	deleted permanently	
upgrade_test	14	10	4		912.1 MB	Dec 02, 2021	Dec 02, 2021		
Unnamed history	1	1			0 bytes	Nov 11, 2021	Nov 11, 2021	deleted permanently	
CX 18s third	176	135	3	38	15.5 GB	Nov 14, 2018	Sep 23, 2021		
new	30	26	6		1.5 GB	May 09, 2021	May 26, 2021		
old_test_data	22	13	9		2.1 GB	Apr 12, 2021	Apr 14, 2021		
fanhaonan	5	5			0 bytes	Jan 13, 2021	Jan 16, 2021	deleted permanently	

History

search datasets

build

420 shown, 68 deleted, 1 hidden

23.64 GB

488: ANOVA test for null model

487: Null model result

486: otu_forall1.txt

485: group_forall.txt

484: dissimilarity result of bray

483: Galaxy11-[sample_list.txt].tabular

482: Galaxy4-[Unoise8_otu_table.txt] (1).tabular

480: Response ratio result at 90 confidence

479: Response ratio plot at 90 confidence

478: Response ratio result at 90 confidence

477: Response ratio plot at 90 confidence

Using 69.4 GB