

ScHARe I Terra Accounts and Workspaces

Sign up for free temporary billing

If you have not filled out the 1-question form on the Think-a-Thon registration confirmation email already, please provide a Google email address in the chat

You will be:

- registered for ScHARe
- added to a free temporary billing project that will allow you to run all the Think-a-Thon materials with your instructors

You will be active on this billing project for about 1 day after the Think-a-Thon

If you want to access work-in-progress from the Think-a-Thon after this time, you will need to set up your own billing and copy any of your workspaces to your own billing



Science collaborative for Health disparities and Artificial intelligence bias Reduction

ScHARe



National Institute on Minority Health and Health Disparities

Office of Data Science Strategy

NIH



National Institute of Nursing Research

Thank you



Outline

- **5'** Introduction and setup
 - Experience poll
- **15'** ScHARe and Terra overview
 - Interest poll
- **20'** Registration and Account
- **30'** Billing and Costs
- **20'** Workspaces and Permissions
- **30'** Notebooks and Environment

Experience poll

Please check your level of experience with the following:

	None	Some	Proficient	Expert
Python				
R				
Cloud computing				
Terra				
Health disparities research				
Health outcomes research				
Algorithmic bias mitigation				

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Part I ScHARe and Terra overview



ScHARe is a cloud-based population science data platform designed to accelerate research in health disparities, health care outcomes, and artificial intelligence (AI) bias mitigation strategies

The platform offers researchers at all career levels and disciplines:

- Access to centralized social determinants of health and other social science datasets
- The ability to collaborate as they apply AI, machine learning, and other advanced analytical techniques to these datasets in a secure setting





ScHARe aims to fill three critical gaps:

- Foster research collaborations and increase participation of women and underrepresented populations with health disparities in data science
- Leverage social science (SDoH) research opportunities afforded by Big Data and cloud computing
- Advance Al bias mitigation strategies and ethical inquiry by increasing the use of diverse eyes and skills



ScHARe Components

ScHARe co-localizes within the cloud:

- Datasets (including social determinants of health and social science data) relevant to minority health, health disparities, and health care outcomes research
- Data repository to comply with the required hosting, managing, and sharing of data from NIMHD- and NINRfunded research programs
- Computational capabilities and secure, collaborative workspaces for students and all career level researchers
- Tools for collaboratively evaluating and mitigating biases associated with datasets and algorithms utilized to inform healthcare and policy decisions

Frameworks: Google Platform, Terra Interface, GitHub, NIMHD Web ScHARe Portal

Intramural & Extramural Resource





Researchers can access, link, analyze, and export **a wealth of datasets** within and across platforms relevant to research about health disparities, health care outcomes and bias mitigation, including:

- Google Cloud Public Datasets: publicly accessible, federated, de-identified datasets hosted by Google through the Google Cloud Public Dataset Program
 Example: American Community Survey (ACS)
- ScHARe Hosted Public Datasets: publicly accessible, de-identified datasets hosted by ScHARe Example: Behavioral Risk Factor Surveillance System (BRFSS)
- Funded Datasets on ScHARe: publicly accessible and controlled-access, funded program/project datasets using <u>Core Common Data Elements</u> shared by NIH grantees and intramural investigators to comply with the NIH Data Sharing Policy
 Examples: Jackson Heart Study (JHS); Extramural Grant Data; Intramural Project Data

Access to social and behavioral datasets

- The cloud offers access to vast repositories of data, and enables mapping and linking across sources
- Extremely large datasets are statistically analyzed to gain detailed insights, often using AI and substantial computerprocessing power
- Datasets can be linked together to see how patterns in one domain affect other areas
- Data can be structured into fixed fields or unstructured as free-flowing information
- Increase use of Dark Data



ScHARe Data Ecosystem will offer access to **300+ datasets**, including:

- Google Cloud Public Datasets
- ScHARe Hosted Public Datasets
 - American Community Survey
 - U.S. Census

- Social Vulnerability Index
- Food Access Research Atlas
- Medical Expenditure Panel Survey
- National Environmental Public Health Tracking Network
- Behavioral Risk Factor Surveillance System
- Repository for Funded Datasets on ScHARe
 Compliance with NIH Data Sharing Policy

Data organization

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On ScHARe, datasets are categorized by content based on the CDC **Social Determinants of Health categories**:

- 1. Economic Stability
- 2. Education Access and Quality
- 3. Health Care Access and Quality
- 4. Neighborhood and Built Environment
- 5. Social and Community Context

with the addition of:

- Health Behaviors
- Diseases and Conditions

Users will be able to **map and link** across datasets

Cloud computing strategies

- AI/ML can analyze massive amounts of data properly without human intervention
- Terra, standalone or in conjunction with Google Cloud Platform's Vertex AI, can support AL & MLbased analyses
- Workflows (pipelines) are a series of steps performed by a compute engine for bulk analysis
- Computing environments can be customized or standardized (using a custom Docker Image or a startup script)



 Uses workflows in Workflow Description Language (WDL), a language easy for humans to read, for batch processing data.

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- Python and R, including libraries most commonly used
- Enables customization of computing environments to ensure everyone in your group is using the same software
- Big query and Tensorflow access for advanced machine learning
- Enables researchers to create interactive Jupyter notebooks (documents that contain live code) and share data, analyses and results with their collaborators in real time
- For novice users, integration with SAS is planned

Cloud security

- The Terra system has been granted Authority to Operate as a FISMA Moderate impact system and is FedRAMP authorized
- NIST framework NIST-800-53 Rev 4 Moderate, to achieve compliance with industry-accepted security standards



- Secure Workspaces
- Duos used for organization and security
- **RAS** single-sign on in future

Al bias mitigation strategies

- Widespread use of AI raises a number of ethical, moral, and legal issues – likely not to go away
- Algorithms often are "black boxes"
- Biases can result from:
 - social/cultural context not considered
 - design limitations
 - data missingness and quality problems
 - algorithm development and model training
 - implementation
- If not rectified, biases may result in decisions that lead to discrimination, unequitable healthcare, and/or health disparities
- Lack of diverse perspectives Populations with health disparities are underrepresented in data science
- Guidelines and recommendations emerging from HHS, NIST, White House, etc.



Critical thinking can rectify AI biases.

ScHARe was created to:

- foster participation of populations with health disparities in data science
- promote the collaborative identification of bias mitigation strategies across the continuum
- create a culture of ethical inquiry and critical thinking whenever AI is utilized
- build community confidence in implementation approaches
- focus on implementation of Al bias guidelines and recommendations



CORE COMMON DATA ELEMENTS

NOVEL CDE FOCUSED REPOSITORY TO FOSTER INTEROPERABILITY

COMPLY WITH DATA SHARING POLICY - HOST PROJECT DATA

DATA ECOSYSTEM

- Map across datasets
- Map across platforms



Schare Common Data Elements and Harmonization

- Through significant efforts, data can be harmonized to a particular data dictionary
- To incorporate external data, users must harmonize in an adhoc manner that is specific to a particular use case, and requires significant manual effort
- It is difficult to share these mappings with others or re-use the ad-hoc harmonization in other projects



SchARe Harmonization through mapping

- Harmonization should be shared: data standards cannot be enforced on everyone
- We are building a mapping system to relate datasets to data dictionaries and transform them on the fly (we call the result of a transform a dataview)
- Mappings can easily be shared amongst the community: a user who mapped CDC data to ScHARe's DD can then share it with the community



SchARe Mapping extends between platforms

- By providing mappings and simple connector plugins asneeded, we can accommodate new data sources and map between data dictionaries
- This provides the flexibility to expand the ecosystem as new standards are developed and adopted
- Current common data elements can be incorporated into mappings immediately





This creates an extraordinary opportunity for high-impact collaborations across platforms

Learning how to use Terra on ScHARe will open up a world of possibilities, giving you access to an interdisciplinary wealth of datasets and resources

Interest poll

I am interested in (check all that apply):

□ Learning about Health Disparities and Health Outcomes research to apply my data science skills

□ Conducting my own research using AI/cloud computing and publishing papers

□ Connecting with new collaborators to conduct research using Al/cloud computing and publish papers

□ Learning to use AI tools and cloud computing to gain new skills for research using Big Data

Learning cloud computing resources to implement my own cloud

Developing bias mitigation and ethical AI strategies

 \Box Other

Two Ways to Sign up for ScHARe News



Scannable from your screen!





- Monthly sessions (2 hours)
- Instructional/interactive
- Designed for new and experienced users
- Research & analytic teams to:
 - Conduct health disparities, health outcomes, bias mitigation research
 - Analyze/create tools for bias mitigation
- Publications from team collaboration
- Networking
- Mentoring and coaching

Instructional

Research teams

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3	April 18, 2023 2.00-4.30.pm	•	Vite: Schöller 2 - Accounts, Worksports, and Analyses Descriptions: An inside load at UnAMP's Terra industries general introduction and features; take to create and configure an eccentral and set up talking. New to reade a worksport and all the appropriate generations, have to close to create and non-a notebook; data and workflows <u>participant</u> .	3	August 14, 2023 2:30-4:30 pm	7	Title: Data Science Projects 5 - Health Outcomes Description: Investigating the influence of non-clinical factors on dispertite in health care delivery a handwom assoint for researchers and students at all levels interstead in collaborating on SciAllie to develop innovative research questions and projects leading to publications.

Register:



bit.ly/think-a-thons

Scerare

Part II Registration and Account

Registering for ScHARe

Complete the following steps to register for ScHARe:

- 1. Visit the ScHARe portal on the NIMHD website: nimhd.nih.gov/schare
- 2. Click on the "Register for ScHARe" button

Funding Opportunities About Programs News and Events Resources Science Collaborative for Health disparities SchARe and Artificial intelligence bias REduction ScHARe is a cloud-based social science data platform designed to accelerate research in minority health and health disparities, health care outcomes, and artificial intelligence bias mitigation strategies. The platform offers researchers at all career levels: · Access to social determinants of health and other social science datasets. · The ability to collaborate as they apply artificial intelligence, machine learning, and other advanced analytical techniques to these datasets in a secure setting. ScHARe aims to foster research collaborations and increase participation of underrepresented populations in data science so that everyone can benefit from the research opportunities

Register for ScHARe

afforded by Big Data and cloud computing.





On the registration page:

Click on the "Register for ScHARe on Terra" button



Registering for SchARe

Complete the registration form

The ScHARe team will:

- review and approve your application
- send you an email with additional instructions within 3 business days
- Note: you will need a Gmail account or another email account (an institutional email, for example) associated with a Google identity. If you do not have it, you can create one here:

bit.ly/3QeUngh



* Required

Welcome to ScHARe

ScHARe (Science Collaborative for Health disparities and Artificial Intelligence bias REduction) is a National Institutes of Health (NIH) project powered by Terra (a scalable and open-source Google Cloud based platform) for biomedical researchers and their collaborators to access the same data and run analyses together in dedicated, secure online spaces.

ScHARe is a collaboration between the National Institute on Minority Health and Health Disparities (NIMHD) and the National Institute of Nursing Research (NINR) at the National Institutes of Health (NIH).

To obtain access to the ScHARe platform on Terra, please fill in the form below. The ScHARe team will review and approve your application and send you an email with additional instructions within 3 business days.

The questions marked with an asterisk (*) are required.

The email you will receive after ScHARe registration approval will ask you to **complete the following steps:**

1. Access the ScHARe Terra workspace at:

bit.ly/access-schare

2. Click on the blue "Log in" button

Welcome to Terra Community Workbench

Terra is a cloud-native platform for biomedical researchers to access data, run analysis tools, and collaborate. Learn more about Terra.

If you are a new user or returning user, click log in to continue.

LOG IN

-



- 4. Select "Sign in with Google"
- 5. Sign into Terra. Your username is the Google email address you provided to request access to ScHARe
- 6. Click "Next" and enter your Google account password to login

Terra	Sign in to continue to Terra
	Email or phone
G Sign in with Google	Forgot email?
Sign in with Microsoft	Create account Next

7. You will see a New User Registration page. Insert your name and contact email, then click on "Register"

	ERRA
New User Regi	stration
First Name *	Last Name *
Contact Email for Notifica	ations *
REGISTER CANCEL	

8. Review and accept the Terra Terms of Service



Access to social and behavioral datasets

- The cloud offers access to vast repositories of data, and enables mapping and linking across sources
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- Social Vulnerability Index
- Food Access Research Atlas
- Medical Expenditure Panel Survey
- National Environmental Public Health Tracking Network
- Behavioral Risk Factor Surveillance System
- Repository for Funded Datasets on ScHARe
 Compliance with NIH Data Sharing Policy

Scerare

Part III Workspaces and Permissions

What is a workspace

Workspaces are the building blocks of Terra - a dedicated space where you and your collaborators can access and organize the same data and tools and run analyses together

They are like **computational sandboxes** with everything you need to complete your project: data, analysis tools, documentation

You can use workspaces to:

- Link to data in the cloud for analysis, instead of downloading and storing it yourself
- Combine data from different sources in a single table for analysis
- Keep data organized with integrated spreadsheet-like tables - no matter where in the cloud the data are stored
- Visualize and analyze data in real time using Python and R (and soon, SAS)
- Find and run bulk analysis tools (workflows) even if you're not a programming expert
- Share reproducible analysis results
- Collaborate while maintaining control of your resources
Dashboard

The landing page (i.e., Dashboard) is your project overview - the questions you're trying to answer, the data and analysis tools you'll use, etc.

The dashboard uses the **Markdown language**, which lets you organize text with headers and include links and additional references



Science Collaborative for Health disparities and Al bias REduction

ScHARe (Science Collaborative for Health Disparities and Artificial Intelligence bias Reduction) is a cloud-based research collaboration platform developed by the National Institute on Minority Health and Health Disparities (NIMHD) and the National Institute of Nursing Research (NINR).

The aim of the ScHARe program is to increase participation of underrepresented populations in data science and cloud computing so that everyone can benefit from the research opportunities afforded by Big Data.

An integral part of the program is the <u>ScHARe Think-a-Thons</u> webinar series, conceived to help prepare underrepresented researchers, students, and their collaborators to use the ScHARe platform and learn how to leverage Big Data and cloud computing. Participants will share knowledge and skills, and form cross-disciplinary, multi-level collaborations around innovative research projects that can lead to breakthrough publications.

Dashboard

Workspace details are populated automatically in the right column of the Dashboard

Expandable sections include:

- Workspace information: creation date, date last updated, your access level
- Cloud information: location of workspace storage, estimated storage cost and size, etc.
- Workspace owners
- Workspace tags: only visible to owners, tags are useful for searching and indexing

WORKSPACE INFORMATION	~
Last Updated	3/6/2023
Creation Date	1/10/2023
Workflow Submissions	0
Access Level	Project Owner
CLOUD INFORMATION	~
Cloud Name	
Location	us us-central1 (lowa)
Google Project ID	terra-d3cb8107 📋
Bucket Name	fc-secure-d6e25d73 📋
Estimated Storage Cost Updated on 3/11/2023	\$0.43
Bucket Size Updated on 3/11/2023	21.4 GiB
Open bucket in browser @	
Open project in Google Cloud	Console 2*
OWNERS ()	~

Data

In the **Data** tab, like spreadsheets built right into the workspace, **data tables help keep track of all project data**, no matter where files are stored in the cloud

- In the ScHARe workspace, click on the Data tab
- Under Tables, you will see a list of dataset categories
- If you click on a category, you will see a list of relevant datasets
- Scroll to the right to learn more about each dataset

More information will be provided in the next Think-a-Thons

Workspaces ScHARe/ScHARe > Data						
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EducationAccessAndQuality (47)	0	0	AdjustedGraduationRate_2015-2016	Education Access and Quality		
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HealthCareAccessAndQuality (10)	0	0	AdjustedGraduationRate_2017-2018	Education Access and Quality		
MultipleCategories (15)	0	0	AdjustedGraduationPlate_2018-2019	Education Access and Quality		
III NeighborhoodAndBuiltErwironment (10)	0	0	ECPP_EarlyChildhoodProgramParticip	Education Access and Quality		
SocialAndCommunityContext (4)	0	0	ECPP_EarlyChildhoodProgramParticip	Education Access and Quality		
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Analyses

In the **Analyses** tab, you can interact with the data and perform/share analyses using Python or R

- In the ScHARe workspace, click on the **Analyses** tab
- This tab contains instructional notebooks and analysis tutorials on how to use the ScHARe resources in the Terra environment

More information on our instructional notebooks will be provided later

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Your Analyses	+ START
Application	Name 1
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Jupyter	01_Introduction to Terra Cloud Environment.ipynb
Jupyter	02_Introduction to Terra Jupyter Notebooks.ipynb
Jupyter	03_R Environment setup.ipynb
Jupyter	04_Python 3 Environment setup.ipynb

Workflows

In the **Workflows** tab, you will find workflows for bulk analyses

- These are the sorts of repetitive analyses that can be automated
- Workflows in Terra are written in the human-readable Workflow Description Language (WDL)

More information will be provided in the next Think-a-Thons

DASHBOARD DATA ANALYSES WORKFLOWS WORKFLOWS Find a Workflow To get started, click Find a Workflow		HARe >	spaces > ScHARe/S rkflows	CES Wo	WORKSPA	= 💿
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Creating your first workspace

Let's create your first Terra workspace!

For the purpose of this tutorial, let's assume that you intend to create a workspace that will allow you to work with two groups of collaborators:

- Group 1 Internal collaborators: researchers in your lab, who must be able to access your data, perform computations, and work with you to write the collaborative notebooks you plan to use to share your results with the scientific community
- Group 2 External collaborators: researchers at another institution, who you want to be able to see your data, notebooks and analyses, but without the possibility of modifying them

Click on the menu in the top left corner of the page, then on "Groups"



On the Groups page, select "Create a New Group" and proceed to **create two different groups**, one for each of the two groups of collaborators previously identified



On the Groups page, select "Create a New Group" and proceed to **create two different groups**, one for each of the two groups of collaborators previously identified



For each group, click on the name of the group and, in the following screen, on "Add User"



Add the Google email address of at least one researcher to each group. If you want one or more of your collaborators to be able to manage users and groups, check the "Can manage users (admin)" box

	User email *
ROUP MANAGEMENT: GROUP-1-INTE	RNAL researcher1@gmail.com
+ Add User	Role
Email 👃	CANCEL ADD USER

You now have two lists of collaborators with whom you can share your workspace, assigning different roles. It's time to create an **authorization domain** for your workspace!

What is an authorization domain (AD)?

Data in the cloud is more secure than data stored locally. Terra has several layers of protection.

ADs are data protection mechanisms - **managed groups with strictly enforced workspace permissions** that, like a badge, follow the original workspace and all copies of it.

The AD guarantees that only members of the AD can clone (copy) the original workspace. If anyone tries to share a cloned (copied) workspace with a colleague who doesn't have the right badge (doesn't belong to the AD), the recipient won't be able to view the workspace. **Your data and analyses are safe!**

Create an **authorization domain** for your workspace, by adding a new group called "Collaborators" in the "Groups" section.

	oup-	GROUP MANAGEMENT
GROUP MANAGEMENT	Enter a unique name *	+ Create a New Group
+ Create a New Group	Only letters, numbers, underscores, and dashes allowed	Group Name
Group Name	CANCEL CREATE CROUP	Collaborators
Group-1-Internal-Collaborators	C.	
Group-2-External-Collaborators	Group-2-External-Cr	Group-1-Internal-Collaborators
ScHARe	ScHAPeghfrecloud:	Group-2-External-Collaborators
ScHARe-Think-a-Thons	ScHARe-Think a-Thi	

Both your groups of collaborators will need to **belong to your "Collaborators" authorization domain** to be able to access your workspace. Simply add the group email associated with each group (circled in blue below) to the "Collaborators" group list: every email address in your groups will now belong to the "Collaborators" authorization domain as well. You can add additional users to each group at any time



	Add user to Terra Group User email *	
GROUP MANAGEMENT: COLLABORATORS	Group-1-Internal-Collaborators@firecloud.org	
+ Add User	Role	
Email 1	CANCEL ADD USER	
	Admin	
	GROUP MANAGEMENT: COLLABORATORS	
	Allow anyone to request access 🕕	
	+ Add User	
	Email 1	Roles
	Group-1-Internal-Collaborators@firecloud.org	Member
	Group-2-External-Collaborators@firecloud.org	Member

It's now time to create your workspace!

In the "Workspaces" section, click on the "+" sign next to "Workspaces"



Assign a **name** to your workspace, select our free temporary **billing project "ScHARe-Temp"** (or create one following the instructions provided during this Think-a-Thon), and add a **description**

Create a New Workspace Workspace name *	Create a New Workspace Workspace name *
My-Workspace	My-Workspace
Billing project *	Billing project *
Select a billing project 🗸	Select a billing project 🗸 🗸
Description	ScHARe-Temp
My first workspace	
CANCEL GREATE WORKSPACE	CANCEL CREATE WORKSPACE

Assign the "Collaborators" **authorization domain** to the workspace

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workspace name	
My-Workspace	
Billing project *	
ScHARe-Temp	~
Bucket location ()	
us-central1 (lowa) (default)	~
Description	
My first workspace	
Authorization demain 0	
Authorization domain 0	
Select circuius	~

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ScHARe-Temp	~
Bucket location ()	
us-central1 (lowa) (default)	~
Description	
My first workspace	
Authorization domain ()	
Select groups	~
and the second	
Collaborators	
Group-1-Internal-Collaborators	
Group-1-Internal-Collaborators Group-2-External-Collaborators	POV
Collaborators Group-1-Internal-Collaborators Group-2-External-Collaborators ScHARe	av

Workspace name *	
My-Workspace	
Billing project *	
ScHARe-Temp	~
Bucket location ()	
us-central1 (lowa) (default)	~
Description	
My first workspace	
Authorization domain 🕕	
Collaborators ×	~

Congratulations, your workspace was successfully created!

Ξ 🍺		CES Da	kspaces > ScHARe-T shboard	emp/My-Workspace >	103		COVID-19 Data & Tools
DASHBOARD	DATA	ANALYSES	WORKFLOWS	JOB HISTORY			
ABOUT THE W					WORKSPACE INFORMATIO	N	×
My Thist Worksp	ace				Last Updated	3/11/2023	
					Creation Date	3/11/2023	
					Workflow Submissions	0	
					Access Level	Owner	
					CLOUD INFORMATION		>
					OWNERS		>
					AUTHORIZATION DOMAIN		>

You are now ready to **share it with the two groups of collaborators** you created:

- 1. Click on the menu in the top left corner of the page, then on "Workspaces"
- Identify your workspace in the list of workspaces provided on screen and click on the corresponding vertical three-dot menu (A), then on "Share" (B)



MY	WORKSPACES (2)	NEW AND INTERESTING (6)	FEATURED (61)	PUBLIC (318)			P	Clone	
	Name			Last Modified	t	Created By	Access Level	Share	Α
	My-Workspace My first workspace			Mar 11, 2023		@gmail.com	Owner	Lock	
$^{\pm}$	ScHARe Think-a	Thons		Mar 6, 2023		calzonil2@nih.gov	Writer	💼 Delete	0

In the drop-down menu, select the group email corresponding to your **first group of internal** collaborators

Share Workspace	Share Workspace
User email	User email
Add people or groups	Add people or groups
Current Collaborators	Group-1-Internal-Collaborators@firecloud.org
@gmail.com Owner v Can share Can compute	Collaborators@firecloud.org Group-2-External-Collaborators@firecloud.org ScHARe@firecloud.org
Share with Support No CANCEL SAVE	ScHARe-Think-a-Thons@firecloud.org Share with Support No CANCEL SAVE



Assigning permissions – Group 1 (Writer)

Since you want this group to be able to:

- access your data
- perform computations
- help you to write the notebooks you will use to share results with the scientific community select the "Writer" role for this group in the dropdown menu and check the "Can compute" box

Any user belonging to the group will be able to **modify** your workspace and analysis notebooks, **and perform computations** for which you will be billed through your Terra Billing Project

Notice how you are the owner of the workspace, and can add other co-owners if desired

ser email		
Add people or groups		ADD
urrent Collaborators		
@gmail.com		
Owner 🗸 🗸	 ✓ Can share ✓ Can compute 	
Group-1-Internal-Collaborate	ors@firecloud.org	
Writer 🗸	Can share Can compute	×
Group-2-External-Collaborat	ors@firecloud.org	
Reader v	Can share Can compute	×

Assigning permissions – Group 2 (Reader)

Since you want this group to be able to see your data, notebooks and analyses, but without the possibility of modifying them, select the "**Reader**" role for this group in the drop-down menu and do not check the "**Can compute**" box. If you also want the group to be able to share your work, check on the "**Can share**" box

Any user belonging to the group will be able to **see**, **but not modify** your workspace and analysis notebooks, and they will **not be able to perform computations** – unless they copy your notebooks and create their own Terra Billing Project, sustaining the cost of the computations

To allow collaborators from Group 1 to perform **computations** for which you will sustain the cost, you have to give them **permission to use your Terra Billing Project**

We will demonstrate how you can do this using the free temporary billing project associated to your workspace

Note - This part of the tutorial is **for demonstration purposes only**. As owners of the free billing project you used to create your demo workspace, we are the only ones able to share it with others. Revisit these slides **after creating your own billing project: you will be able to replicate these steps** using the billing project you own

Instructions:

1. Click on the menu in the top left corner of the page, then on "Billing"



- 2. Identify the **billing project associated with your workspace** in the list of Billing Projects owned by you, and **click on it**
- 3. Click on the "Members" tab

	lling > Billing Project cHARe-Temp
BILLING PROJECTS O CREATE	ScHARe-Temp View billing account
Owned by You 🗸 🗸 🗸	You are the only owner of this shared billing pro account. More information about managing sh
ScHARe 2	Workspaces Members Spend report
ScHARe-Temp	
nimhd-ai-bias-mitigation	Name 1
Shared with You 🗸	My-Workspace
T101-Jan2021	ScHARe Think-a-Thons
	ThinkAThonTest
	Wiz test

4. Click on "Add User", add the **group email** associated with collaborators from Group 1 to the Billing Project, and confirm. If you prefer, you can add specific **individual collaborators** instead of the entire group

	tilling > Billing Project SCHARe-Temp	Add user to Billing Project
BILLING PROJECTS O CREATE	ScHARe-Temp View billing account	User email *
Owned by You 🗸	You are the only owner of this shared billing proj account. More information about managing sha	Group-1-Internal-Collaborators@firecloud.org
ScHARe		Role
ScHARe-Temp 4	Workspaces Members Spend report	Can manage users (Owner)
nimhd-al-blas-mitigation	+ Add User	Warning: Adding any user to this project will mean they can incur costs to the billing associated with this project.
Shared with You 🗸 🗸	ScHARe-Think-a-Thons/Efirecloud.org	CANCEL ADD USER
T101-Jan2021	Construction of Construction of Construction	
	calzonil2@nih.gov	

Success! Your collaborators can now use your Billing Project to perform computations in the cloud

	Billing > Billing Project ScHARe-Temp	
BILLING PROJECTS O CREAT	ScHARe-Temp View billing account	
Owned by You	 You are the only owner of this shared billing project. Consider account. More information about managing shared billing 	er adding another owner to ensure someone is able projects.
ScHARe	Workspaces Members Spend report	
ScHARe-Temp	+ Add User	
nimhd-ai-blas-mitigation	Email 1	Roles
Shared with You	Group-1-Internal-Collaborators@firecloud.org	User
T101-Jan2021	ScHARe-Think-a-Thons@firecloud.org	User
	calzonil2@nih.gov	Owner

One last thing. If you are interested in using the data resources of a workspace or replicating the analyses showcased in its notebooks, and have the appropriate permissions to do so, you can create a copy of such workspace for your personal use. This operation is called **"cloning" the workspace**

You are encouraged to clone the ScHARe workspace and use its resources

Here is how you can do it. In this example, we will clone the workspace "ScHARe Think-a-Thons", a copy of the ScHARe workspace we created for this Think-a-Thon

1. Click on the menu in the top left corner of the page, then on "Workspaces"



2. Identify the workspace you want to clone in the list of workspaces provided on screen and click on the corresponding vertical three-dot menu (A), then on "Clone" (B)

		- N99992	А ₀
Workspaces 😌 Dedicated spaces for you and your collaborators to access and analyze data tog V Recently Viewed	ether. Learn more about workspaces.		
My-Workspace ScHARe Think-a-Thons Viewed Mar 11, 2023, 4:29 PM	AM 🛆		
Search by keyword. Tags V Ac	eess levels v Billing project	✓ Submission status ✓	Cloud platform V
Name	Last Modified † Created By	Access Level	В
My-Workspace My first workspace	Mar 11, 2023 ny152us@gm	nail.com Owner	Clone () A
* ScHARe Think-a-Thons	Mar 6, 2023 calzonil2@nił	ngov Writer	Lock (3) → Leave

- 3. Input a **name** for the workspace copy
- 4. Select the **Billing Project** you want to associate with the workspace. For this example, you can select our free temporary Billing Project "ScHARe-Temp"
- 5. Select the **bucket location**. A bucket location can only be set when creating a workspace. For this example, you can leave the default unmodified
- 6. Change the **Description** if desired
- A cloned workspace will inherit the Authorization Domain (AD) groups of the original workspace. Add an additional AD if needed

ScHARe Think-a-Thons copy	
Billing project *	
ScHARe-Temp	~
Bucket location ()	
us-central1 (lowa) (default)	Y
Description	
	^
### Science Collaborative for Health disparities	~
Authorization domain () Inherited groups: ScHARe, ScHARe-Think-a-Thons	
Select groups	~

Success! The cloned workspace is now listed among your workspaces

You can freely access all of its **resources**

Workspaces 🔂						
Dedicated spaces for you and your colla	aborators to a	ccess and analyze data	together. Learn	more about	workspaces.	
 Recently Viewed 						
ScHARe Think-a-Thons copy	N	My-Workspace		ScH	ARe Think-a-Thons	
Viewed Mar 13, 2023, 2:46 AM	a v	fiewed Mar 11, 2023, 4	:29 PM 🤇	View	nd Mar 11, 2023, 6:02 Al	м
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Scerare

Part IV Notebooks and Environment

What is a notebook?

A Jupyter Notebook is an interactive analysis tool that includes:

- code cells for manipulating and visualizing data in real time (Terra notebooks support Python or R)
- documentation to make it easier to share and reproduce your analysis

To get the most out of this tutorial you should be familiar with **programming**. If you are not, the code in our notebooks is very easy to understand and reuse, and our tutorial will still help you understand how notebooks work

We will:

- cover the basics of creating your first notebook
- explore the instructional notebooks available in the ScHARe workspace and run one of them

Why use notebooks?

A notebook integrates code and its output into a single document where you can run code, display the output, and also add explanations, formulas, and charts

Using notebooks:

- is now a major part of the data science workflow at research institutions across the globe
- can make your work more transparent, understandable, repeatable, and shareable
- will speed up your workflow and make it easier to communicate and share your results

ScHARe notebooks

Let's take a first look at what a notebook can do by checking out the instructional notebooks that **ScHARe offers to help novice users** learn how to use the workspace and its resources

A list of the available notebooks is provided on the right. **We will access and run a copy of one of these notebooks**, as an example

List of ScHARe instructional notebooks

- **00_List of Datasets Available on ScHARe**: a list of the datasets available in the ScHARe Datasets collection.
- **01_Introduction to Terra Cloud Environment**: an introduction to the Terra platform and cloud environment.
- 02_Introduction to Terra Jupyter Notebooks: an introduction to Jupyter Notebooks on the Terra platform.
- **03_R Environment setup**: instructions on how to setup your cloud environment for R-based notebooks.
- 04_Python 3 Environment setup: instructions on how to setup your cloud environment for Python 3-based notebooks.
- 05_How to access plot and save data from public BigQuery datasets using R: instructions on how to access, plot, and save data from datasets available on the cloud through the Google Cloud Public Datasets Program, using R.
- 06_How to access plot and save data from public BigQuery datasets using Python 3: instructions on how to access, plot, and save data from datasets available on the cloud through the Google Cloud Public Datasets Program, using Python 3.
- 07_How to access plot and save data from ScHARe hosted datasets using Python 3: instructions on how to access, plot, and save data from datasets hosted by ScHARe in this workspace.
- 08_How to upload access plot and save data stored locally using Python 3: instructions on how to import to Terra, access, plot, and save data from datasets stored locally on your computer.

Creating a notebook

- Click on the menu in the top left corner of the page, then on "Workspaces"
- 2. Click on the new workspace you created earlier today

	KSPACES	, 📃 🧔
Luca Calzoni	Vlabo	Dedicated space Recently View My-Workspa Viewed Mar 1
Library	~	Search by key
Support	~	MY WORKSPA Name
COVID-19 Data & To		TI ScHAR

Wo	rkspaces 🔂
Dedic	ated spaces for you and your collabor
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My	-Workspace
Vie	wed Mar 13, 2023, 3:51 AM
Sea	rch by keyword
MY	WORKSPACES (2) NEW AND INTE
	Name
*	My-Workspace
	My first workspace

Creating a notebook

- Click on the "Analyses" tab
- 4. Click on the "Start" button


- 5. Select "Jupyter"
- 6. In the next window, assign a **name** to the notebook and choose a programming language ("**Python 3**")
- 7. Click "Create analysis"

Select an application	Create a new notebook Name of the notebook *		Create a new notebook Name of the notebook *
R Studio	Language *	v	Language *
= Galaxy	Python 3 R	~	Python 3
Or Click / Drag to upload an analysis file			
0			

- 8. You will now be asked to configure your **Cloud Environment.** In this window, you can **configure** the ondemand availability of **computer system resources**, especially data storage and computing power, needed to perform your computations:
 - Application configuration: the software application + programming languages + packages used to create your cloud environment. You can leave this unchanged
 - Cloud compute profile: the CPU and RAM available to run your application, which determines how much processing can be done at a time.

Notice how changing the parameters changes the estimated costs to run at the top of the window. **Find the balance that's right for you.** More compute power costs more, and you don't want to request (and pay for) significantly more than your computation needs

0.05 per hr	\$0.00 per hr	\$0.40 per month
pplication configuratio	n O	
Default: (GATK 4.2.4.0,	Python 3.7.12, R 4.2.2)	~
What's installed on this e	environment?	Updated: Dec 7, 2022 Version: 2.2.9
URI		
CPUs 1 v Memo	ry (GB) 3.75 🗸	
Enable GPUs man	Learn more about GPU cost	t and restrictions, 🖻
Enable GPUs man	Learn more about GPU cost	t and restrictions, 🖻

- CPUs, Memory: leave the default values unchanged
- Enable GPUs: checking the box allows you to add graphics processing units (GPUs) to your environment. GPUs are optimized for linear algebra computations, such as matrix multiplication, and for training artificial intelligence and deep learning models. Leave the box unchecked
- Enable autopause: recommended. Once your browser becomes inactive and 30 minutes pass, the cloud environment autopauses to save on costs
- Location: physical location where the data is going to reside. Leave the default value unchanged
- Persistent disk: Terra attaches a persistent disk to your cloud compute to provide an option to keep the data on the disk after you delete your compute. A minimal cost per hour is associated with maintaining the disk, even when the cloud compute is paused. Select "Standard" and "10 GB" and click on "Create"

CPU type	NVIDIA Tesla T4	~ GPUs	1 ~	
Compute t	ype			
Standard	VM			~
so cation us-centra	minutes of inactivity	v		
30 C Location US-centra Persistent of Where your	minutes of inactivity minutes of inactivity	Learn more a	bout persistent o	disks and
30 C Location us-centra Persistent of Persistent of where your Disk Type	minutes of inactivity	Learn more a Disk Size (G	ibout persistent o	disks and

• Success! Your environment has been configured, and your notebook created. Click on its name to open it

Ξ 💿	NORKSPA	Work CES Ana	spaces → ScHARe-Te Iyses	emp/My-Workspace >
DASHBOARD	DATA	ANALYSES	WORKFLOWS	JOB HISTORY
Your Anal	yses +	START		
Application	n	Name 🌡		
Jupyter Jup	yter	Test.ipynb		

Creating the cloud environment will take 3-5 minutes

= 💿	WORKS	SPACES	Worl Tes	kspaces > ScHARe t .ipynb	Ten	np/My-Workspace > analyses >
DASHBOARD	DATA	AN	ALYSES	WORKFLOWS		JOB HISTORY
PREVIEW (READ	ONLY)	OPEN		ROUND MODE	1	Creating cloud environment. You can navigate away and return in 3-5 minutes.
in []:						

Since this newly-created notebook is empty, we will open and run one of the **ScHARe instructional notebooks** instead, to give you a closer look at **how notebooks work**.

- Click on the menu in the top left corner of the page, then on "Workspaces"
- 2. Click on the ScHARe Think-a-Thons workspace that was shared with you by the instructors

	=
L Luca Calzoni V	Works Dedicated
Workspaces	My-Wo Viewed
🚺 Library 🗸	Search b
? Support ~	
COVID-19 Data & Tools	My Sc

Wo	rkspaces 🔂	
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Sei	rch by keyword	16
MY	WORKSPACES (2) NEW AND IN	TER
	Name	
31	My-Workspace	
	My first workspace	

 Click on the
 "02_Introduction to Terra Jupyter Notebooks" notebook

	Workspaces - ScHARe-Temp/ScHARe Think-a-Thons - CES Analyses
DASHBOARD DATA	ANALYSES WORKFLOWS JOB HISTORY
Your Analyses +	START
Application	Name ↓
Jupyter	00_List of Datasets Available on ScHARe.ipynb
Jupyter Jupyter	01_Introduction to Terra Cloud Environment.ipynb
Jupyter Jupyter	02_Introduction to Terra Jupyter Notebooks.ipynb
Jupyter	03_R Environment setup.ipynb

4. Click on "**Open**"

Ξ 🙋	WORKSPA	CES 02	kspaces > ScHARe-To _Introduction	emp/ScHARe Think-a-Thons > analyses > to Terra Jupyter Notebooks.ipynb
DASHBOARD	DATA	ANALYSES	WORKFLOWS	JOB HISTORY
PREVIEW (REA	D-ONLY)	OPEN :		
Int: Maybe	roduction 1	Dipyter notebooks and y	Votebooks you're interested in a cra	ash course in what they are and how to start using them
Not	ebook overv	iew		
This no	otebook will help you	understand:		
• W	hy notebooks are use	d in biomedical resear	ch	
• Tł	ne relationship betwee	n the notebook and th	ne workspace	
• Ju	pyter Notebook basic	s: how to use a notebo	ook, install packages, and	d import modules
• 0	ommon libraries in da	ta analysis and popula	r tutorial notebooks	

- Configure the environment as shown before, then click on "Create"
- 6. Follow the step-by-step instructions in the notebook, which will help you understand:
 - why notebooks are used in research
 - the relationship between the notebook and the workspace
 - how to use a notebook, install packages, and import modules
 - common libraries in data analysis



SCHARE

Part V Billing and Costs

What are the cloud costs of working on Terra?

The Terra platform is free to use

However, the following operations in Terra **may incur** charges:

1. Virtual Machine compute costs

In cloud computing, a **virtual machine** is an emulation of a computer system that provides the functionality of a physical computer

Terra allows you to **customize** the characteristics of your virtual machine based on your computation needs (more on this later)

- A high-performance machine costs more
- You will be charged for the time you use the machine

Cloud compute profile		
CPUs 1 v Memo	ory (GB) 3.75 🗸	
Enable GPUs	Learn more about GPU cost and re	strictions. 🗗
Standard VM		~
Location and a (lowa) (defa	ault)	
Persistent disk Persistent disks store and your disk is mounted.	alysis data. Learn more about persis	stent disks and who
Disk Type	Disk Size (GB)	
Standard	× 10	

What are the cloud costs of working on Terra?

2. Data storage

• You will be charged for any data stored in the storage spaces ("**buckets**") associated with your account

3. Data egress (i.e. moving data) costs

- As we will see later, when creating a bucket to store data you are asked to set its location ("region"), i.e. the physical place where the data is going to reside (e.g.: "US-CENTRAL1 Iowa", or "US-EAST4 Northern Virginia"). More info on regions <u>here</u>. You will pay to move stored data to a bucket in a different region
- You will pay to download data from a bucket that is configured as a Requester Pays bucket. With such buckets, the data requester pays the cost of the data download instead of the bucket owner

How will I be charged for these costs?

Terra runs on Google Cloud Platform (GCP). All Terra costs are GCP fees that are ultimately paid for by a **Google Cloud Billing account** linked to Terra – specifically, to a **Terra Billing project**



How will I be charged for these costs?

Will I incur any costs today?

Today and for one day after the Think-a-Thon, **access to a free temporary billing project** will allow you to run all the materials with your instructors

What happens after tomorrow?

You will no longer have access to the free temporary billing project. If you want to access work-inprogress from the Think-a-Thon, you will need to **set up your own billing** and copy any of your workspaces to your own billing

Next, we will show you how to set up your own billing

Get \$300 in free Google Cloud credits

If you've never used Google Cloud before, **you are eligible for \$300 in free Google Cloud credits** you can use for working in Terra

Conditions for Google Cloud credits eligibility

- You haven't previously signed up for the Free Trial
- You've never been a paying customer of Google Cloud, Google Maps Platform, or Firebase
- If you're part of an organization that uses Google Cloud, your email will likely not be eligible



Google Cloud

What can I do with my credits in Terra?

The credits will cover anything that has a cost in Terra - such as storing data and running analyses. You can't use credits to add <u>GPUs</u> to your computing resources, and you are limited to 4 workspaces at a time

How long will my \$300 credits be available?

Your credits will be available for 3 months, or until you have used up all \$300. Once your credits run out or expire, you can upgrade to a paid account

Step 1. Set up a Google Cloud Billing account

- 1. Go to the **Google Cloud console** at <u>https://console.cloud.google.com/</u> and sign in with your Terra user ID. If you haven't already set up a billing account, you'll be invitated to activate your free trial
- 2. Click the activate button and follow the instructions



Step 1. Set up a Google Cloud Billing account

9	SWITCH ACCOUNT
Country	
United States	
What best describes your orga	nization or needs?
Please select	
Terms of Service	Google Cloud Platform Terms of
Service, Supplemental Free service of any applicable se	Trial Terms of Service, and the terms or ervices and APIs.

Access to all Cloud Platform Products

Get everything you need to build and run your apps, websites and services, including Firebase and the Google Maps API.

\$300 credit for free

Put Google Cloud to work with \$300 in credit to spend over the next 90 days.

No autocharge after free trial ends

We ask you for your credit card to make sure you are not a robot. You won't be charged unless you manually upgrade to a paid account.



Step 1. Set up a Google Cloud Billing account

- You'll need to verify your identity with a one-time verification sent to a cell phone, and give a credit card, PayPal account or bank account (you won't be billed until the credits expire)
- Verify the Google Cloud Billing account in the <u>Billing page</u>. You should see My Billing Account in the top left
- 5. Google will create a project, **My First Project**, funded by your free credits, in the **My Projects** tab



Step 2. Link the Cloud Billing account to Terra

The next step is to link the Google Cloud Billing Account to your Terra account, so that Terra and Google can communicate about cost and billing

You must use the same Google ID for both the Cloud Billing account and your Terra user name

- 1. When logged into Google with your Terra user ID, go to the Google Cloud Console Billing page.
- 2. Select the checkbox beside the Google Cloud billing account you will use for Terra.
- 3. On the right panel, below **Permissions**, select the **Add Principal** button.
- 4. Add "terra-billing@terra.bio" under New Principal in the form.
- 5. In the dropdown, select the role **Billing > Billing Account User**.
- 6. Click Add.
- 7. Click on the **Save** button

Note: "terra-billing@terra.bio" will appear in the list as "terra-billing@firecloud.org." This is expected.

Step 3. Create a Terra Billing project

Once Terra is linked to a Cloud Billing account, you can create a Terra Billing project that will allow you to create a workspace to store and analyze data

- 1. Go to the **<u>Billing page</u>** from the main navigation (click on **your name** to expand the drop-down, and select **Billing**)
- 2. Click on the "+ Create" button at the top left
- 3. If prompted to **Enable Billing Permissions**, select the **Google identity** of the Google Cloud Billing account, and click **Allow**. This lets Terra access Cloud Billing accounts associated with your Terra user name (Google ID).
- 4. Enter a **unique name** for your Terra Billing project
- 5. Select the **Google Cloud Billing account** that will fund the Billing project

You may see multiple Cloud Billing accounts that you can select for this Terra Billing project. If you need to locate a Billing account ID, navigate to the **Google Developers Console** and click on **Billing**. Look for the number below **Billing account ID**

You can **ESTIMATE COSTS**:

- 1. analysis costs
- 2. cloud storage costs
- 3. egress (i.e., data moving) costs

You can **CHECK ACTUAL COSTS** in the Google Cloud Platform Console

You can **REDUCE COSTS** in several ways (for advanced users)

Provo per nr	< \$0.01 per hr	\$2.00 per month	-
Application configurat	ion O		
Default (GATH 4-2.0.0	Tymas 37.10, #4.1.1)		
Whet's installed on this	environment?	Upstated Sep 23, 2021	
UR			
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avandaria vw			
Persistent disk size (G Denistent disk size (G Denistent disk size a your disk is mounted.	N Nalyski diata. Luarri mioni abi	out perclaterit disks and where	

- Adjust settings to optimize cost (VM and disk)
- 2) Estimate costs using real-time cost/hour in Cloud Environment widget



- Updates based on the machine configuration you choose
- Total cost (estimate) = (cost/hour) x (hours the VM will be active) + cost of the Persistent Disk
- Autopause function saves money!

You can **ESTIMATE COSTS**:

- 1. analysis costs
- 2. cloud storage costs
- 3. egress (i.e., data moving) costs

You can **CHECK ACTUAL COSTS** in the Google Cloud Platform Console

You can **REDUCE COSTS** in several ways (for advanced users)

ASHBOARD DATA NOTEBOOKS WORKFLOWS JOB HISTORY			1				
ABOUT THE WORKSPACE	WORKSPACE	INFORMATION					
This workspace reproduces the fundamental steps in a genome wide association study (GWAS),	3/19/2020	2010/2020 0/27/2021					
using 1,000 Genomes Project (phase 3) genotypes and simulated phenotypes.		ACCOMPLAYER Written					
1 Explore phenotypes and population structure (Jupyter Notebook - Hail/Python)	\$0.07	amp-128-op					
 Test for genetic associations using mixed-models and generate summary visualizations (WDL workflow) 	OWNERS	X					
he output of the notebook (part 1) serves as the input to the workflow (part 2).	tmajaria@bros	dinstitute org					
nstructions for applying the analyses presented in this workspace on your own data are provided in he penultimate section of this documentation.	TAGS O		-				
Notes on data in this workspace	Add a taig 🗸 🗸			Estimated cloud			
To demonstrate an analysis that could be run on typical whole genome sequence data, this workspace provides mock phenotype data generated from publicly available 1000 Genomes phase 3 genotypes. Phenotypes have been simulated based on individual genotypes and known		1000 Genomes X GWAS X Dupyter Notabooks X WDLs X		storage costs for your workspace			
ssociated loci for multiple complex traits. The <u>GCTA activate</u> ⁶ was used with lists of causal variants and an estimate of narrow sense heritability ⁶ for each phenotype.	Google Bucket						
fraits and sources for causal variants	Location: #	ulti-region: US	-				

You can **ESTIMATE COSTS**:

- 1. analysis costs
- 2. cloud storage costs
- 3. egress (i.e., data moving) costs

You can **CHECK ACTUAL COSTS** in the Google Cloud Platform Console

You can **REDUCE COSTS** in several ways (for advanced users)

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B pheno-data (2504)	D	DOI 3 Mor	ore Information	
g program (1)		cios		DONE
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You can **ESTIMATE COSTS**:

- 1. analysis costs
- 2. cloud storage costs
- 3. egress (i.e., data moving) costs

You can **CHECK ACTUAL COSTS** in the Google Cloud Platform Console

You can **REDUCE COSTS** in several ways (for advanced users)

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You can **ESTIMATE COSTS**:

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You can **CHECK ACTUAL COSTS** in the Google Cloud Platform Console

You can **REDUCE COSTS** in several ways (guides are for advanced users) Terra allows you to find the right balance between cost and time

Saving on workflow costs

- ► Delete intermediate files: <u>guide</u>
- ► Call-caching: guide
- ► Checkpointing: guide
- ► Preemptible VMs: guide

Saving Cloud Environment costs

- ► Size application compute appropriately: <u>guide</u>
- Move generated data to regional or nearline storage: <u>guide</u>
- Autopause: <u>guide</u>

Saving on storage costs

- ► Ask how much are you storing, where are you storing it, and how frequently will you access it?
- ► Move data to regional or nearline storage: <u>guide</u>



Thank you

Next Think-a-Thons: Data Sets Review and Analyzing Data

Terra tutorials and resources

If you are new to Terra, we recommend exploring the following resources:

- <u>Overview Articles</u>: Review high-level docs that outline what you can do in Terra, how to set up an account and account billing, and how to access, manage, and analyze data in the cloud
- Video Guides: Watch live demos of the Terra platform's useful features
- <u>Terra Courses</u>: Learn about Terra with free modules on the Leanpub online learning platform
- <u>Data Tables QuickStart Tutorial</u>: Learn what data tables are and how to create, modify, and use them in analyses
- Notebooks QuickStart Tutorial: Learn how to access and visualize data using a notebook
- <u>Machine Learning Advanced Tutorial</u>: Learn how Terra can support machine learning-based analysis

Next Think-a-Thons:



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References and credits

• **Tutorials and notebooks:** The Broad Institute, Inc., Verily Life Sciences, LLC