



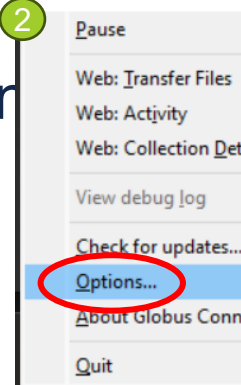
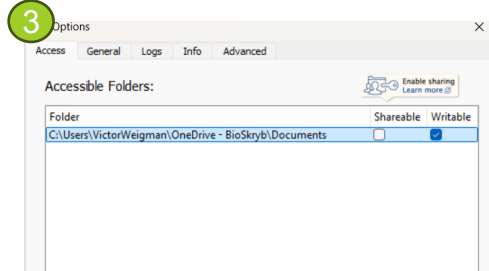


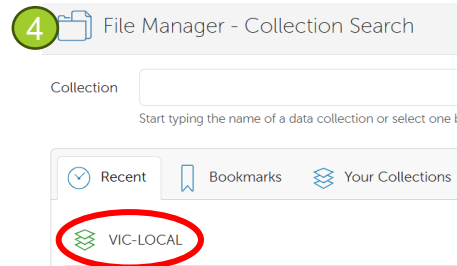
# Globus Data Transfer Guide

July 2025



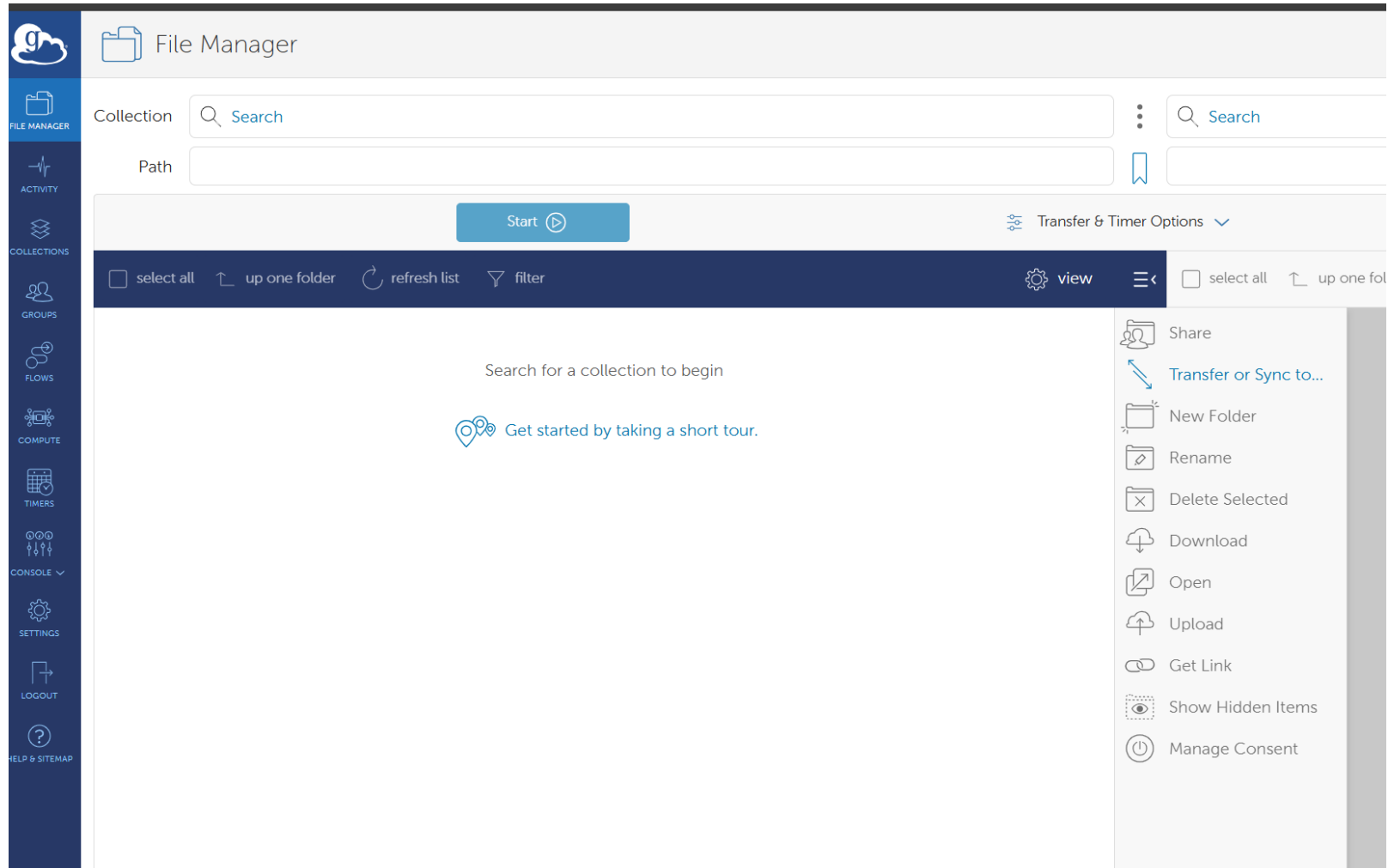
# Ensure you have a destination

- If your institution already uses globus, your account should have access to their collections and your space in them
- If you do not, you should get Globus Connect Personal ([here](#))
- [1] You will be able to see network shares your device can see, you just have to mount it to GCP (icon in task tray)  
- [2] Will need to right click on the icon to pull up menu, select 'Options' 
- [3] This will bring up a set of directories. Add new ones with "+" 
- Go to path as you normally would in File Explorer
- [4] That Collection will show up in the Globus File Manger (with whatever you name it)




# Log into Globus, go to File Manager

[File Manager | Globus](#)




# Demo Data '0\_PublicDatasets'




Pulls up all collections. Guide: [Demo Data - BaseJumper Docs](#)










 File Manager

Collection

Path

Start 

☐ select all  up one folder  refresh list  filter

NAME 	LAST MODIFIED
 0.Export	12/31/1969, 06:59 PM
 Embryo	12/31/1969, 06:59 PM
 Exome	12/31/1969, 06:59 PM
 ResolveDNA	12/31/1969, 06:59 PM
 ResolveOME	12/31/1969, 06:59 PM
 ResolveOME_Biorxiv	12/31/1969, 06:59 PM
 TargetEnrichment	12/31/1969, 06:59 PM
 Ultima	12/31/1969, 06:59 PM

If you are looking to pull data out of your organization workspace, click on Collections and you will be able to see it

# Transfer

Source

Destination

File Manager

Collection: 0\_Public-Datasets → VIC-LOCAL

Path: /ResolveOME/PBMCs/ → /C:/Users/VictorWeigman/Downloads/Data/

Click 'Start' to transfer

Start

Select Files / Folders

Collection

Path

Start

Transfer & Timer Options

select none up one folder refresh list filter

view

select all up one folder refresh list filter

NAME	LAST MODIFIED	SIZE
<input checked="" type="checkbox"/> PBMCs-frs-run1-a-RNA-S41_S41_L001_R1_001.fastq.gz	12/20/2023, 08:39 AM	18.68 MB
<input checked="" type="checkbox"/> PBMCs-frs-run1-a-RNA-S41_S41_L001_R2_001.fastq.gz	12/20/2023, 08:39 AM	18.83 MB
<input checked="" type="checkbox"/> PBMCs-frs-run1-a-RNA-S5_S5_L001_R1_001.fastq.gz	12/20/2023, 08:39 AM	14.35 MB
<input checked="" type="checkbox"/> PBMCs-frs-run1-a-RNA-S5_S5_L001_R2_001.fastq.gz	12/20/2023, 08:39 AM	14.41 MB
<input checked="" type="checkbox"/> PBMCs-frs-run1-a-RNA-S52_S52_L001_R1_001.fastq.gz	12/20/2023, 08:39 AM	13.85 MB
<input checked="" type="checkbox"/> PBMCs-frs-run1-a-RNA-S52_S52_L001_R2_001.fastq.gz	12/20/2023, 08:39 AM	13.76 MB
<input checked="" type="checkbox"/> PBMCs-frs-run1-a-RNA-S63_S63_L001_R1_001.fastq.gz	12/20/2023, 08:39 AM	14.88 MB
<input checked="" type="checkbox"/> PBMCs-frs-run1-a-RNA-S63_S63_L001_R2_001.fastq.gz	12/20/2023, 08:39 AM	14.89 MB
<input checked="" type="checkbox"/> PBMCs-frs-run1-a-RNA-S74_S74_L001_R1_001.fastq.gz	12/20/2023, 08:39 AM	13.91 MB
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<input checked="" type="checkbox"/> PBMCs-frs-run1-a-RNA-S84_S84_L001_R1_001.fastq.gz	12/20/2023, 08:39 AM	15.52 MB
<input checked="" type="checkbox"/> PBMCs-frs-run1-a-RNA-S84_S84_L001_R2_001.fastq.gz	12/20/2023, 08:39 AM	15.52 MB
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<input checked="" type="checkbox"/> PBMCs-frs-run1-b-RNA-S16_S16_L001_R1_001.fastq.gz	12/20/2023, 08:39 AM	18.25 MB
<input checked="" type="checkbox"/> PBMCs-frs-run1-b-RNA-S16_S16_L001_R2_001.fastq.gz	12/20/2023, 08:39 AM	18.44 MB
<input checked="" type="checkbox"/> PBMCs-frs-run1-b-RNA-S28_S28_L001_R1_001.fastq.gz	12/20/2023, 08:40 AM	16.74 MB
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<input checked="" type="checkbox"/> PBMCs-frs-run1-b-RNA-S29_S29_L001_R1_001.fastq.gz	12/20/2023, 08:39 AM	22.27 MB

180 items selected

Permissions

Transfer or Sync to...

New Folder

Rename

Delete Selected

Download

Open

Upload

Get Link

Show Hidden Items

Manage Consent

This folder is empty.

# Closing notables

- When you start, you can leave the window as it will start a job in the background
- You will get an email from transfer start and completion
- The 'Start' button works on both sides (so you can upload if you select start in the 'Destination' to move to source
  - This works for all non-Demo collections. You cannot upload to Demo Datasets
- If you can't see a path in your destination you think should be there try to put '/' (root path) in the Path field and press Enter
- If you are stuck, email [basejumper@bioskryb.com](mailto:basejumper@bioskryb.com)