

Downloading your data RUN from BaseSpace

Open a Basespace account (free) here-

<https://login.illumina.com/platform-services-manager/?rURL=https://basespace.illumina.com&clientId=basespace&clientVars=aHR0cHM6Ly9iYXNlc3BhY2UuaWxsdW1pbmEuY29tL2Rhc2hib2FyZA&redirectMethod=GET#/register>

Please fill out your email address associated with your BS account in the appropriate place in the Sample Submission Form

Once your sequencing run is complete you will receive an invitation for ownership transfer of your run and project to your account. After accepting the transfer all the run's data will be available only to you and will be removed from the TGC account. Please make sure to download and backup your data!

To better understand the technical aspects of your run, please visit the following link from *Illumina Software Resources/BaseSpace™ Sequence Hub: does my sequencing run look good?*-

<https://developer.illumina.com/news-updates/basespace-sequence-hub-does-my-sequencing-run-look-good>

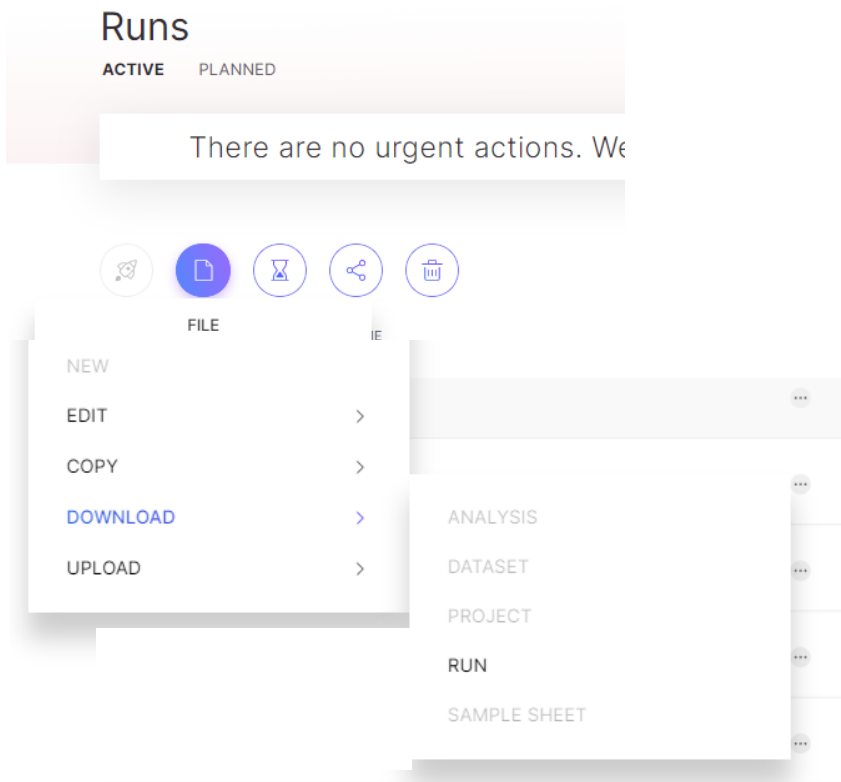
For general help with Illumina Basespace please visit- <https://help.basespace.illumina.com/>

To download data from a run as a package of FASTQ files or SAV files (technical data of your run), use the following steps:

(1) Go to RUNS and open/choose the desired run.



(2) Select File, point to Download, and then select Run.




(3) Select the file type you want to download.

- FASTQ—FASTQ files.
- Sequencing Analysis Viewer (SAV)—InterOp and other files required to run SAV.

Data is downloaded in a compressed (*.zip) file format.

Download Run ×

Run Name	Size
	697.13 MB


BaseSpace Sequence Hub Downloader must be installed. It's a one-time installation and provides fast and secure downloads via SSL.

[INSTALL DOWNLOADER](#)

Select the file types to be downloaded:

FASTQ

SAV

 No analysis files are available for this run

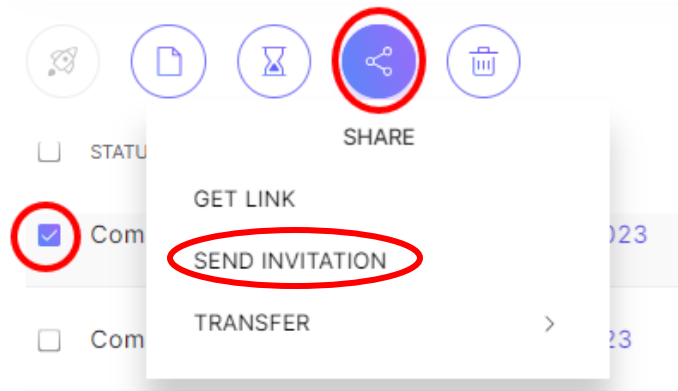
[CLOSE](#) [DOWNLOAD](#)

If you do not have the FASTQ option available, go to **PROJECTS** and open/choose the desired run. Repeat the same process (File- Download- Project).



Share your project with another BS account:

- (1) Check the box of the run you want to share and then, using the 'Share' button, send a sharing invitation to your collaborator (Send Invitation).



- (2) Fill in your collaborator Basespace user account email and then press the 'Add Collaborator' button.

Share this runs ×

231026_VH00495_255_AACW3V2M5

Invite a collaborator
Email address

Optional Message
Add a personal message (optional)

Remaining characters : 140/140

Share the associated Project(s) as well (**Recommended**)
Unchecking this box will limit what the recipient will have access to.

ADD COLLABORATOR

CANCEL **SAVE SETTINGS**

Share this runs ×

231026_VH00495_255_AACW3V2M5

Invite a collaborator
Email address ←

Optional Message
Add a personal message (optional)

Remaining characters : 140/140

Share the associated Project(s) as well (**Recommended**)
Unchecking this box will limit what the recipient will have access to.

ADD COLLABORATOR

CANCEL **SAVE SETTINGS**

(3) Save the settings.

Share this runs ×

230711_VH00495_227_AAC37NGHV

Invite a collaborator
Email address

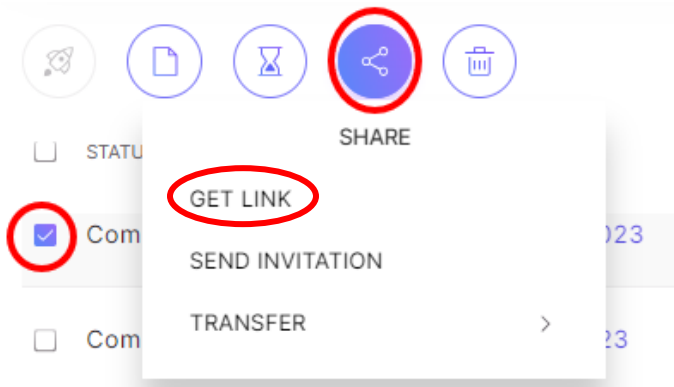
Optional Message

Remaining characters : 140/140

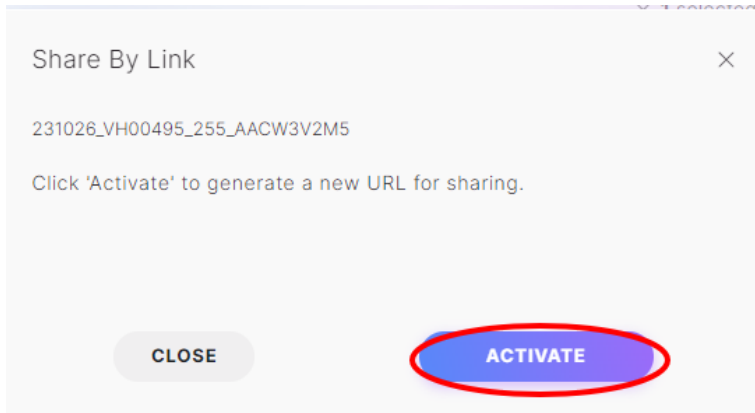
Share the associated Project(s) as well (**Recommended**)
Unchecking this box will limit what the recipient will have access to.

Share your run with a non-BS user:

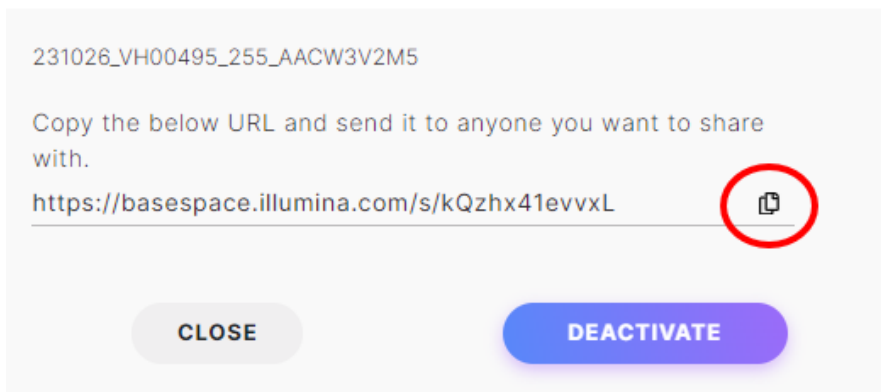
(4) Check the box of the run you want to share and then, using the 'Share' button, create a link (Get Link).



(5) Activate the link.



(6) Copy the link.



If you encounter problems, contact the TGC team.