

Accessing Sequencing Data

All of the information regarding your sequencing run will be on iLab.

Once your samples are submitted and everyone has approved the request charges, you can monitor the progress of your project on iLab.

Log into iLab and on the Next-Generation Sequencing Facility page expand your project by clicking the down arrow.

Expand the Forms and Request Details to monitor the progress of your project. Green indicates the step is started or complete.

▼ Forms and Request Details (see bottom of list to add items to this request)

Step	Description	Note	Status
Receive library samples	click to edit	click to edit	Finished
Library QC	click to edit	click to edit	Finished
Library Pooling	click to edit	click to edit	Not Started
Sequencing run started	click to edit	click to edit	Not Started
Transfer sequencing data	click to edit	click to edit	Not Started

- Library pooling: once complete you can find the final pool concentration
- Sequencing run started: here you can find the Run ID given by the sequencer, the concentration that was loaded on the sequencer and the % of PhiX added
- Transfer sequencing data: this indicates to the sequencing core how data was transferred

We will be communicating to you in the Comments section of iLab throughout the sequencing project, and we would greatly appreciate you communicating to us in this way as well. This ensures everyone in the sequencing core has the most up-to-date information and we can better serve you. When we do add a comment, iLab will send an email as well.

Once your sequencing project is complete, we will let you know through the Comments section. Additionally, we will upload any pertinent information related your sequencing run in the Attachments & URLs section. This may include:

- Bioanalyzer traces
- MiSeq Denature Worksheet (indicating how we loaded the sequencer)
- Sample Tracking Form containing QC data from the Qubit, Bioanalyzer and qPCR
- Links to the run and project folder on BaseSpace

Comments add comment

Oct 10 '16 10:47 AM Meghan Sloan said
Hi everyone,
I apologize, the last Sample tracking form I uploaded still had the same mistake.
The Sample tracking form that is on here now has the correct data.
Thanks,
Meghan

Attachments & URLs add attachment add url

- N-GSF-AG-7_Muse-196AB_High_Sensitivity_DNA... Meghan Sloan 2016 Oct 10 @09:39
- N-GSF-AG-7_Muse-196_High_Sensitivity_DNA_A... Meghan Sloan 2016 Oct 10 @09:40
- N-GSF-AG-7_Muse-196_MiSeq_Denature_workshe... Meghan Sloan 2016 Oct 10 @09:40
- N-GSF-AG-7_Muse-196_Sample_tracking_form.xlsx Meghan Sloan 2016 Oct 10 @10:39