

ILLUMINA SUBMISSION FORM

Date: _____

PLEASE ENSURE BOTH SIDES OF FORM ARE FILLED OUT ENTIRELY

PI: _____

Institution/Company Name: _____

Submitted By: _____

Financial Contact and Billing Address: _____

Telephone: _____

PI Email Address: _____

Submitter Email Address: _____

Account/P.O.No.: _____

Signature: _____

UW Affiliation: UWCCC Waisman Ctr.
 UW System ExternalFunding Source: NIH NSF GLBRC
 USDA OtherUW Institution: SMPH CALS L&S SOP
 COE Other _____**LIBRARY DETAILS**

TO BE PREPARED BY UWBC NEXT GEN SEQUENCING FACILITY (please choose one):

DNA: GBS: Amplicon: Other (discuss with facility before submitting):**- OR -**

LIBRARY ALREADY PREPARED:

Library Type: _____ Send QC Results? (Y/N): _____ Sample Buffer: _____

NON-SEQUENCING SERVICES (e.g. Pippin HT Size Selection, QC Only, Shearing, Transfer):

- Submission guidelines and sample sheet can be found here.
- Email the sample sheet to nextgen-seq@biotech.wisc.edu.
- All submissions must be accompanied by a hard copy of this submission form.

FOR OFFICE USE ONLY:

Submission # _____ Run # _____ Run Date _____ Billing Date _____ Order # _____

SEQUENCING OPTIONS

MiSeq

of samples _____

of flowcells _____

NovaSeq6000*

of samples _____

of lanes _____

*MiSeq nano required for all full flow cells and lanes.

NovaSeq6000 (2x150 shared ONLY)

of samples _____

of reads per sample (in millions)** _____** $\pm 20\%$ of stated value above

Use sample list to denote variable read requests.

Custom Sequencing Run

(scRNA, asymmetric read lengths, etc)

Must be approved by DNA Sequencing Facility.

Must be purchased as a full flow cell.

Sequencer: _____

Parameters: _____

Custom Sequencing Primers (Please provide at 100uM in no less than 50ul.)

Name: _____ Read (read 1, read 2, index 1): _____

BIOINFORMATICS OPTIONS (OPTIONAL)

Analysis to be done by (please choose one):

 UWBC Bioinformatics Resource Center (Please email brc@biotech.wisc.edu to inform them of your project.)

 Collaborator - Name _____ Email _____ UWNetID _____

SAMPLE DISPOSAL

We store dilutions of all libraries (both customer prepared and facility prepared) for 10 years. We will only store input DNA or customer prepared stock library for 6 months from submission date. If you would like your excess DNA/stock library back, please arrange a time with nextgen-seq@biotech.wisc.edu to pick up DNA. Please note that we will ship samples to customers at their expense. For shipping outside the US, customer must contract a courier.

Choose one of the following options:

Please dispose of input DNA and/or stock library 6 months from submission date.

I am responsible for making arrangements to pick up my excess DNA and/or libraries. I acknowledge that DNA and/or libraries will be disposed 6 months from submission. (Submitter and PI will be notified/reminded within the text of the email notifying them of "Illumina data ready for download.")

I have read and understand the Illumina Sequencing Policy found on the [UWBC website](#) (please initial): _____

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