

SEQUENCING SUBMISSION FORM
Please ensure both pages are entirely filled out
ALL CUSTOMERS

Sample submitted by: _____

Email address: _____

PI: _____

PI email address: _____

Signature: _____

EXTERNAL CUSTOMERS ONLY

Institution/company name: _____

Bill to/accounts payable:

Name: _____

Address: _____

Telephone: _____

Email: _____

P.O. no.: _____

Name of organism _____

 Input nucleic acid: gDNA cDNA CHIP (shearing size _____)

 total RNA mRNA ribo-reduced RNA smallRNA

 Other (provide explanation of content: _____)

 Library prepared by lab (QC only). Kit used: _____

 FFPE extracted samples? Yes No

 Sample quantification method: Qubit/fluorometer Nanodrop/Spec. No initial quantification

 Library type: gDNA CHIP WGBS Exome 10x long read

 strand mRNA strand total RNA smallRNA 10x scRNA

 Other (discuss with Core before submitting): _____

Sample buffer: _____

 Indexing: Single Dual

No. samples per flowcell: _____

Sample information

Please fill out the following table with the appropriate information.

- Submit samples in 1.5 mL single tubes or a 96-well plate.
- All sample tubes/plates must be labeled with the sample name, PI last name and date.
- Sample names must be fewer than eight characters; only alphanumeric characters can be used with no spaces.
- If submitting > 8 samples, please attach an Excel list of all samples to this form and email all sample information to genomics@vai.org.

	Sample name	Conc. (ng/μL)	A ₂₆₀ /A ₂₈₀	A ₂₆₀ /A ₂₃₀	Volume (μL)	Total (μg)	Index sequence
Sample 1							
Sample 2							
Sample 3							
Sample 4							
Sample 5							
Sample 6							
Sample 7							
Sample 8							

For genomic DNA and exome projects only: Attach gel image here or send via email to genomics@vai.org.

Sequencing options:

NextSeq 500 Mid Output (200M reads)

2x75 bp

2x150 bp

NextSeq 500 High Output (400M reads)

1x75 bp

2x75 bp

2x150 bp

Custom primer (name): _____

Bioinformatics options:

VARI Bioinformatics and Biostatistics Core

External collaborator. Name: _____ Email: _____

Submitting lab/VARI collaborator. Name: _____ HPC account ID: _____

I have read and understand the Illumina Sequencing Policy found on the Genomics Core website (genomicscore.vai.org/resources).

Please initial: _____