

Requesting A Quote

Name:	Email:
PI Name:	PI Email:
Institution:	Department:
# of Samples:	Species:
HTSF Account Name (if Known):	

Submitting Material Type:

Raw Material
 Libraries
 Pools
 Other: _____

Source Material Type:

RNA
 Total RNA
 Low Concentration Total RNA
 Other: _____

DNA
 Whole Genomic DNA
 ChIP DNA
 Amplicons
 Other: _____

Library Type:

- KAPA Stranded mRNA
- KAPA Hyper DNA
- TruSeq Stranded RNA - Ribo-Zero Gold

- SMARTer/ Nextera low input RNA library
- Small RNA/ miRNA libraries
- 16s Metagenomic
- Other: _____

Pooling Plan:

No Pooling
 Pool in Groups of _____
 One Pool Including All Samples

Sequencing Platform:

Miseq Nano
Miseq
HiSeq 2500, Rapid Run
HiSeq 2500-v4 High Output
Hiseq 4000

Novaseq6000-SP
Novaseq6000-S1*
Novaseq6000-S2*
Novaseq 6000-S4*

*XP Options available. If single lane is requested. This may delay running of the flowcell.

TBD(If you are unsure of the depth of sequencing needed please)

Read Type Format:

- Single End
- Paried End

Preferred Cycle Length: _____

Number of lanes each pool should be run on: _____
(typically each pool is run on 1 lane. If unsure, put 9999)

Note: If you are unsure of the best platform for your needs, please list ether of below as preferred: We can Reply with the best platform for your need

- # reads/ sample: _____
- # clusters/ Sample: _____

Goal of Sequencing/ Hypothesis: