

<b>To be used for estimates only: Please fill out inquiry form at <a href="http://www.bumc.bu.edu/bnorc/core-services-overview/core-user-request-form/">http://www.bumc.bu.edu/bnorc/core-services-overview/core-user-request-form/</a> or contact: <a href="mailto:funge.core@gmail.com">funge.core@gmail.com</a> for quotes</b>		
<b>Project consultation</b>	<b>BNORC</b>	<b>Non-BNORC</b>
Initial project set up	\$100	\$150
Project consultation (hourly)	\$150	\$200
<b>Sequencing Per Flowcell</b>	<b>BNORC</b>	<b>Non-BNORC</b>
NextSeq High 75 cycle	\$1,903	\$2,508
NextSeq High 150 cycle	\$3,287	\$4,329
NextSeq High 300 cycle	\$5,106	\$6,721
	<b>BNORC</b>	<b>Non-BNORC</b>
<b>Full Service Library Preparation - includes QC</b>	per sample	per sample
Crosslinking	\$15	\$20
Shear Check	\$150	\$200
Shearing	\$52	\$70
ChIP setup	\$37	\$50
ChIP antibody	\$7	\$10
DNA, cDNA or ChIP libraries	\$236	\$317
RNA preparation	\$25	\$33
mRNA libraries	\$247	\$332
Poly-A selection	\$40	\$53
Ribo-reduction	\$70	\$93
ATAC-seq	please inquire	please inquire
If submitting tissue requiring core to setup and perform FACS, charge for FACS will be \$150/hr		

	<b>BNORC</b>	<b>Non-BNORC</b>
<b>Miscellaneous</b>	per sample	per sample
BioAnalyzer Analysis (standalone)	\$10	\$13
Qubit quantification (Initial set up fee)	\$10	\$13.50
+ per sample	\$1	\$1.35
	<b>BNORC</b>	<b>Non-BNORC</b>
<b>Bioinformatic Consultation (per hour)</b>	\$100	\$135
<b>Per flowcell (Estimates)</b>		
Standard ChIPseq/ATACseq output (QC/alignment/browser track generation)	\$300	\$400
+ Peak calling, library normalization, counts	\$500	\$665
+ Differential enrichment/GSEA	\$800	\$1,064
Motif Enrichment	\$100	\$133
Standard RNAseq output (QC/alignment/browser track generation)	\$300	\$399
+ Transcript determination, normalization, counts	\$500	\$665
+ Differential expression/GSEA	\$800	\$1,064
Longterm Storage/TB/month	\$20	\$27