

## FY18 Molecular Bioinformatics Core (mBIO) Services and Fees

*Prices will increase by 3% each fiscal year.*



The pricing structure listed below is effective as of July 1, 2017. For all custom projects, please consult the mBIO Core Director for a quote.

### Standard data analysis:

Workflow	Cost per sample
Whole Genome Sequencing	\$162
Whole Exome Sequencing	\$162
RNA Sequencing	\$162
Microarray Analysis	\$54
Differential Proteomics	\$75
ChIP-seq	\$100
ATAC-seq	\$100
Targeted Sequencing	\$75
<i>DeNovo</i> Assembly	\$1,000

### Individual Tasks:

Service	Cost
Quality Control Assurance	\$20 per sample
Alignment	\$100 per sample
Biostatistics	\$30 per sample
Variant Calls	\$75 per sample
I-Pathway Guide Analysis	\$75 per paired condition
String Analysis	\$50 per paired condition
Gene Set Enrichment Analysis	\$75 per paired condition
DAVID Pathway Analysis	\$50 per paired condition
Heatmap	\$10 per image
Principal Components Analysis (PCA)	\$20 per paired condition
Pearson's correlation coefficient	\$10 per image
Venn Diagram	\$10 per image
Volcano Plot	\$10 per image
Custom Scripting / Data Mining	\$70 per hour
Training	\$70 per hour
GEO Upload Support	\$200 per experiment
Experiment Design Assistance or Consultation	No charge for two hours; \$70/hr. afterward, or allocation of percent effort on grant



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**Long-term, secure storage of data:** \$0.13/GB

**GraphPad Prism 7.0 software site license:** \$125/copy/computer/year  
(Mac or PC)

### External customers:

External, academic: 50% surcharge

External, commercial: To be negotiated directly with the Office of Research