

FY17 Molecular Bioinformatics Core (mBIO) Services and Fees



The pricing structure listed below is effective as of **December 1, 2016**. For all custom projects, please consult the mBIO Core Director for a quote.

Data analysis, standard workflows: \$52.00/hour
DNA-seq (whole genome sequencing, WGS, or whole exome sequencing WES), RNA-seq, ChIP-seq and *de novo* genome assembly, pathway analysis, GEO data repository upload, microarray analysis, single nucleotide polymorphism (SNP) analysis, and data annotation. Development of custom analysis pipelines is also billed at the hourly rate.

Typical experiments:

RNA-seq, WGS, WES, ChIP-seq:	\$156/sample (3h)
Microarray analysis:	\$52/array (1h)
De novo assembly:	\$1,560 (30h)

Long-term, secure storage of data: \$123/TB

Ingenuity Pathway Analysis (IPA) software access:

Per sign-in/project: \$150/project

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