

## UC San Diego Center for Computational Biology & Bioinformatics

### Scope of Work 2020 RNA-seq Analysis

*UCSD Internal rate: \$180/hour*

*Non-profit rate: \$261/hour*

*For-profit rate: \$362/hour*

#### **Bulk RNA-seq**

##### **Primary Analysis**

1. Raw data quality control (method: FASTQC)
2. Alignment of raw data to genome (method: STAR)
3. Gene or transcript level quantification (method: RSEM or Kallisto)

*Cost: 0.5 hours/sample for  $\leq 20$  samples ; 0.25 hours/sample for any additional samples above 20 (bulk rate)*

##### **Secondary Analysis**

1. Assessment of experimental design and analysis design
2. Quality control of read quantification & outlier detection (method: MDS plot)
3. Normalization (method: TMM normalization)
4. Differential expression analysis (method: limma-voom)
5. Functional enrichment analysis & pathway analysis (method: Hypergeometric test and GSEA)
6. Standard figures: MDS plots, MD plots and heatmaps of top 25 DE genes

*Cost: 15-25 hours. Additional time in 5 hour increments for additional design complexity and custom downstream requests or visualizations.*

##### **Data Overlay**

1. Assessment of second data set and methods for data overlay (~2-5 hours)
2. Correlate RNA-seq results and other dataset (~10-25 hours)
3. Overlay RNA-seq results and other dataset on significantly dysregulated pathways (~5-10 hours)

*Cost: Hourly*

##### **Deliverables:**

1. Consultation meeting to discuss experimental design, biological questions and analysis
2. Results files from all analyses outlined
3. Open source code from all analyses
4. Summary powerpoint presentation of all results
5. Post-analysis meeting to discuss results

## **Single cell RNA-seq**

### **Primary Analysis**

- CellRanger

*Cost: 1 hour/sample*

### **Secondary Analysis**

- Seurat

*Cost: 20-40 hours depending on scope. Additional time in 5 hour increments for additional design complexity and custom downstream requests or visualizations.*

### **Trajectory Analysis**

*Cost: 20-60 hours depending on scope.*

## **Spatial Transcriptomics**

### **Visium**

Primary analysis cost: 2 hours/sample

Secondary analysis cost: 20-40 hours depending on scope.

### **Nanostring GeoMX**

Cost: 20-40 hours depending on scope.