

Molecular Bioinformatics Core  
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Number of Samples: \_\_\_\_\_ Order ID: \_\_\_\_\_  
Software Platform Used: \_\_\_\_\_  
Completed File Location: \_\_\_\_\_  
Quote Amount \_\_\_\_\_ Completion Date: \_\_/\_\_/\_\_

### Order Request Form

Date Samples Submitted:	Principal Investigator:
Contact Person:	Department:
Contact Phone:	Account No.:
Contact Email:	Species:
Sequencing Platform:	Authorized Signature:

- **Fastq Location:** \_\_\_\_\_
- **Results will be stored on server for 8 weeks ONLY**

Number of Samples: \_\_\_\_\_

Sequencing Platform: \_\_\_\_\_

Control / Experiment Setup

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### Service Request

- |   |   |
|---|---|
| <input type="checkbox"/> Eukaryote Genome Assembly                  | <input type="checkbox"/> BAM files                        |
| <input type="checkbox"/> Prokaryote Genome Assembly                 | <input type="checkbox"/> Differential Expression Analysis |
| <input type="checkbox"/> Transcriptome Assembly                     | <input type="checkbox"/> Epigenetics                      |
| <input type="checkbox"/> Sequence Alignment<br>(Model Organism)     | <input type="checkbox"/> Variant Analysis                 |
| <input type="checkbox"/> Sequence Alignment<br>(Non-model Organism) | <input type="checkbox"/> Figures for publication          |