**Technical Terms:**

1. Overview of Service

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| --- | --- |
| Species Name | *VAR\_SPECIES* |
| Sample Type | Enriched RNA Sample |
| Sample Number | SAM\_NUM\_1 |
| Library Type | RIP-Seq Library |
| Bioinformatics | VAR\_ALY |
| Turnaround Time | VAR\_TAT working days starting after we receive the library preparation confirmation from client |

1. Sample Requirements

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| Sample Type | Remarks | Amount | RIN | Peaks | Concentration | Purity |
| Enriched RNA Sample | Strongly Recommended | ≥150 ng | ≥8 (Animal)  ≥7 (Plant, Fungi) | For unfragmented sample, the main peak should be higher that 1000bp. | ≥5 ng/μL | OD260/280>2.0  No degradation or DNA contamination |
| Required | ≥100 ng | ≥8 (Animal)  ≥7 (Plant, Fungi) |