

Sample Submission Guidelines

AmpX – Express Amplicon Sequencing

| Sample Type | Purified PCR Product |
|----------------------------|--|
| Sample Size | 150-600 bp |
| Recommended Volume | 20 μL (15 μL minimum) |
| Recommended Concentration | 5-25 ng/μL (2 ng/μL minimum) |
| Illumina Adapter Sequences | For the forward sequencing read: 5'-TCCCTACACGACGCTCTTCCGATCTnnnnnn-3' For the reverse sequencing read: 5'-GTTCAGACGTGTGCTCTTCCGATCTnnnnnn-3' |

| RNA-Seq | mRNA-Seq | Total RNA-Seq |
|---------------------------|-----------------------------|------------------------------|
| Recommended Quantity | 1 μg (200 ng minimum) | 1 μg (200 ng minimum) |
| Recommended Concentration | >10 ng/µL (5 ng/µL minimum) | >50 ng/μL (10 ng/μL minimum) |
| Recommended Volume | 20 μL (15 μL minimum) | 20 μL (15 μL minimum) |
| Quality | RIN ≥ 7 | RIN ≥ 7 |
| Buffer | RNase-free water | RNase-free water |
| Shipment | Ship frozen on dry ice | Ship frozen on dry ice |

* RNA extraction service available

Whole Genome Sequencing

| Recommended Quantity | 1 μg (1 ng minimum) |
|---------------------------|--------------------------------------|
| Recommended Concentration | >10 ng/µL (5 ng/µL minimum) |
| Recommended Volume | 20 μL (15 μL minimum) |
| Quality | A260/280 1.8-2.0 |
| Buffer | Water, EB, or low TE (< 0.1 mM EDTA) |
| Shipment | Ship with ice packs |

* DNA extraction service available

Sequencing Only (for pre-made libraries)

| Recommended Concentration | >10nM (5nM minimum) |
|---------------------------|-----------------------|
| Recommended Volume | 20 μL (15 μL minimum) |
| Sample Type | Final libraries |
| Shipment | Ship with ice packs |

* Pooling and additional QC services available

We recommend using Qubit for concentration measurements. If your samples do not meet our minimum requirements, or if you do not see your sequencing type, please contact us at info@ampseq.com or (240) 654-6236. Our NGS team will evaluate your case and provide guidance on the best approach.