



How do you make sense of all those proposals? Use our [RNA-seq Deliverables Checklist](#) to see what you're getting, and compare this to other providers.

RNA-seq Deliverables Checklist			
<u>Deliverable</u>	<i>Cofactor Genomics</i>	<i>Provider 2:</i>	<i>Provider 3:</i>
Ribosomal Treatment	Ribo-depleted/whole transcriptome		
Molecular Spike-ins	Included		
Library QC prior to sequencing	Included		
FASTQC Quality Plots	Included		
Guaranteed delivery of Q25 data or better (mean)	Included		
Reads/Sample	30 million		
Guaranteed delivery of reads	Included		
Sequence Alignments to Genome	Included		
Sequence Alignments to Transcriptome	Included		
Read Clusters and Average Coverage Calculated	Included		
Data Normalization Across Samples	Included		
Differential Expression Comparison including Fold-Change	Included		
Statistical Analysis including p-values and coefficient of variance (CV) between replicates	Included		
Transcriptome Saturation Analysis for each sample	Included		
Analysis Consultation with Bioinformatics Team	Included		