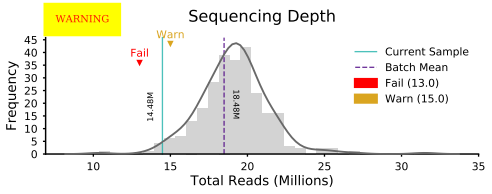
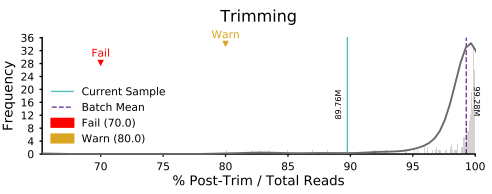
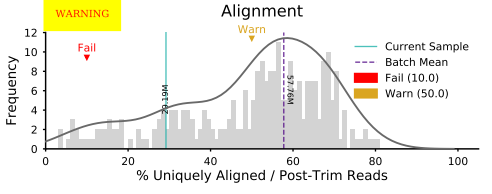
Sample : RNA_B11_01 Batch : batch_11

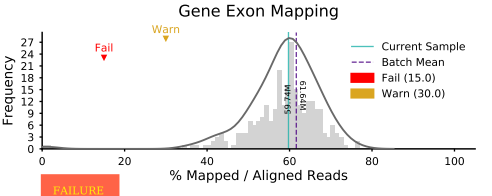
Warn cutoffs: Default alpha =0.8| Sequencing Depth = 15.0| Trimming = 80.0| Alignment = 50.0| Gene Exon Mapping = 30.0| Ribosomal RNA = 0.312| Sequence Contamination = 5.0| Gene Body Coverage = 0.8| Distribution of Gene Expression = 0.8

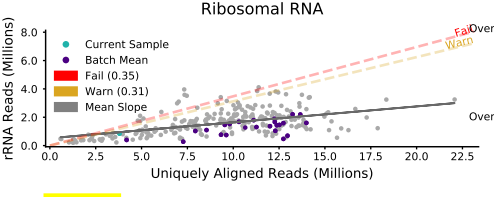
Fail cutoffs: Default alpha =0.9| Sequencing Depth = 13.0| Trimming = 70.0| Alignment = 10.0| Gene Exon Mapping = 15.0| Ribosomal RNA = 0.348| Sequence Contamination = 10.0| Gene Body Coverage = 0.9| Distribution of Gene Expression = 0.9

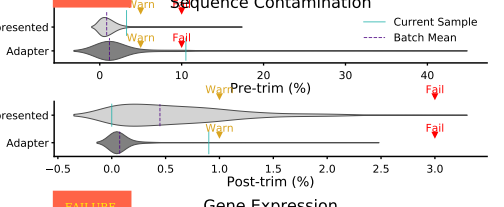


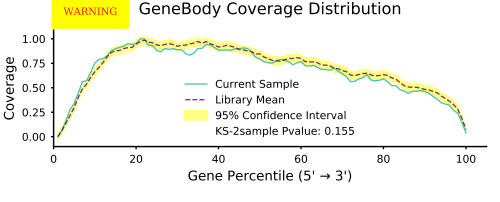


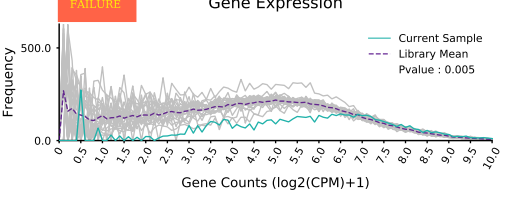










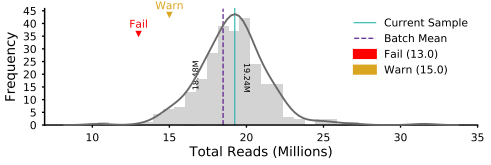


Sample : RNA_B11_02 Batch : batch_11

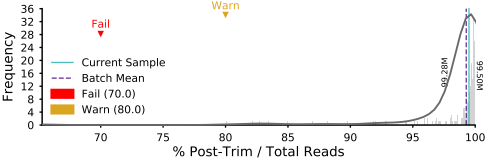
Warn cutoffs: Default alpha =0.8| Sequencing Depth = 15.0| Trimming = 80.0| Alignment = 50.0| Gene Exon Mapping = 30.0| Ribosomal RNA = 0.312| Sequence Contamination = 5.0| Gene Body Coverage = 0.8| Distribution of Gene Expression = 0.8

Fail cutoffs: Default alpha =0.9| Sequencing Depth = 13.0| Trimming = 70.0| Alignment = 10.0| Gene Exon Mapping = 15.0| Ribosomal RNA = 0.348| Sequence Contamination = 10.0| Gene Body Coverage = 0.9| Distribution of Gene Expression = 0.9

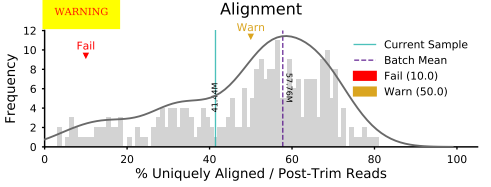
Sequencing Depth



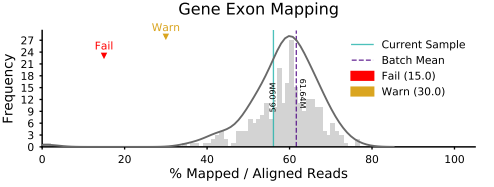
Trimming



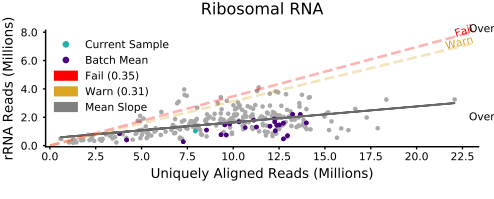
Alignment



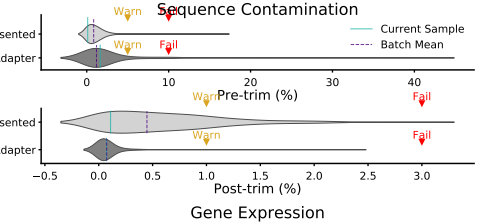
Gene Exon Mapping



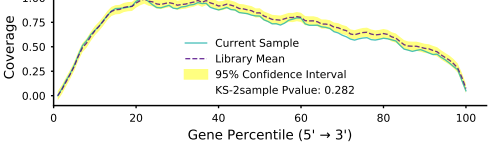
Ribosomal RNA



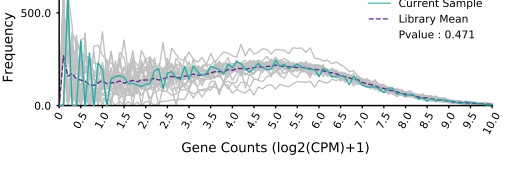
Sequence Contamination



GeneBody Coverage Distribution



Gene Expression

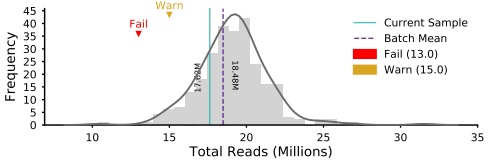


Sample : RNA_B11_03 Batch : batch_11

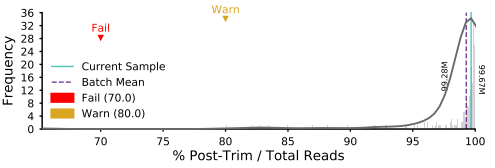
Warn cutoffs: Default alpha =0.8| Sequencing Depth = 15.0| Trimming = 80.0| Alignment = 50.0| Gene Exon Mapping = 30.0| Ribosomal RNA = 0.312| Sequence Contamination = 5.0| Gene Body Coverage = 0.8| Distribution of Gene Expression = 0.8

Fail cutoffs: Default alpha =0.9| Sequencing Depth = 13.0| Trimming = 70.0| Alignment = 10.0| Gene Exon Mapping = 15.0| Ribosomal RNA = 0.348| Sequence Contamination = 10.0| Gene Body Coverage = 0.9| Distribution of Gene Expression = 0.9

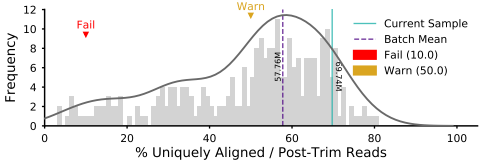
Sequencing Depth



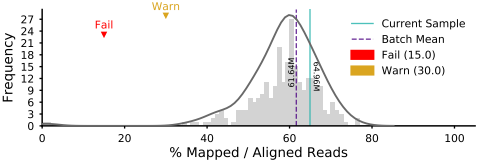
Trimming



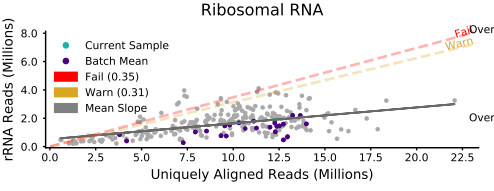
Alignment



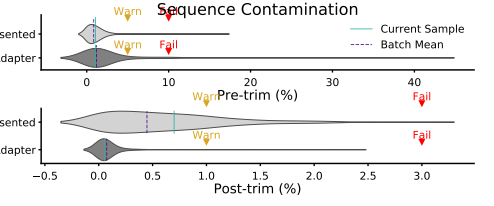
Gene Exon Mapping



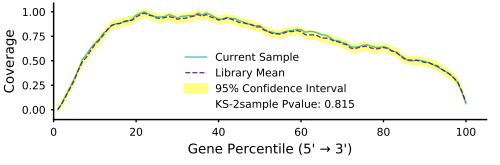
Ribosomal RNA



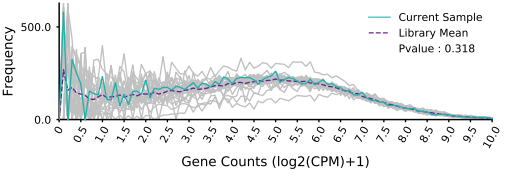
Sequence Contamination



GeneBody Coverage Distribution



Gene Expression

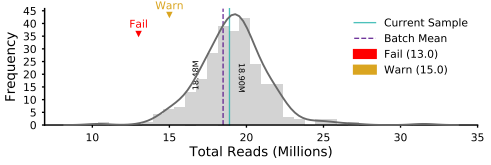


Sample : RNA_B11_04 Batch : batch_11

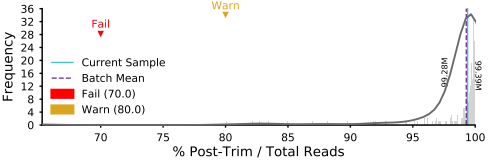
Warn cutoffs: Default alpha =0.8| Sequencing Depth = 15.0| Trimming = 80.0| Alignment = 50.0| Gene Exon Mapping = 30.0| Ribosomal RNA = 0.312| Sequence Contamination = 5.0| Gene Body Coverage = 0.8| Distribution of Gene Expression = 0.8

Fail cutoffs: Default alpha =0.9| Sequencing Depth = 13.0| Trimming = 70.0| Alignment = 10.0| Gene Exon Mapping = 15.0| Ribosomal RNA = 0.348| Sequence Contamination = 10.0| Gene Body Coverage = 0.9| Distribution of Gene Expression = 0.9

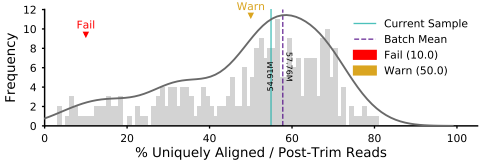
Sequencing Depth



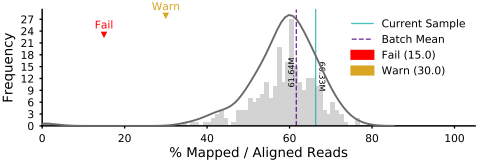
Trimming



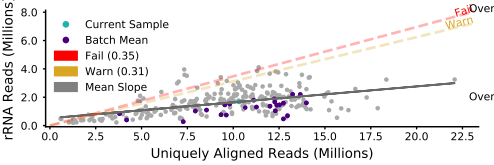
Alignment



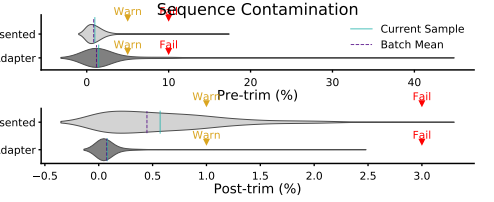
Gene Exon Mapping



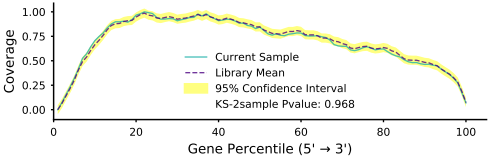
Ribosomal RNA



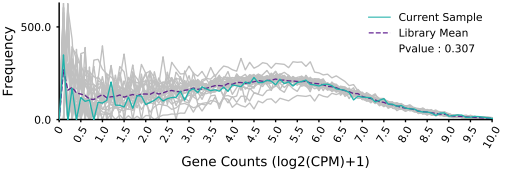
Sequence Contamination



GeneBody Coverage Distribution



Gene Expression

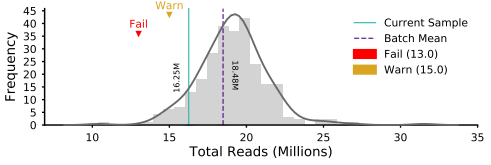


Sample : RNA_B11_05 Batch : batch_11

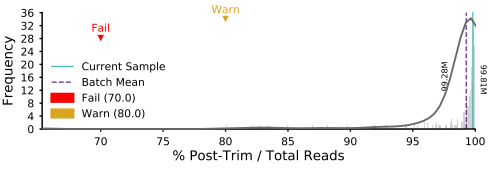
Warn cutoffs: Default alpha =0.8| Sequencing Depth = 15.0| Trimming = 80.0| Alignment = 50.0| Gene Exon Mapping = 30.0| Ribosomal RNA = 0.312| Sequence Contamination = 5.0| Gene Body Coverage = 0.8| Distribution of Gene Expression = 0.8

Fail cutoffs: Default alpha =0.9| Sequencing Depth = 13.0| Trimming = 70.0| Alignment = 10.0| Gene Exon Mapping = 15.0| Ribosomal RNA = 0.348| Sequence Contamination = 10.0| Gene Body Coverage = 0.9| Distribution of Gene Expression = 0.9

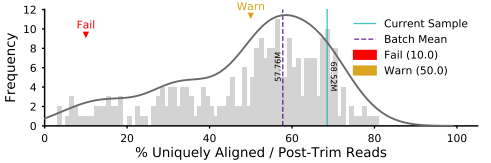
Sequencing Depth



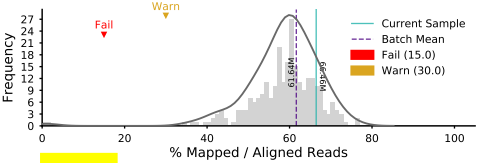
Trimming



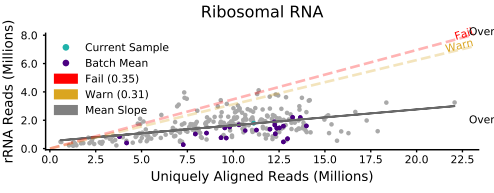
Alignment



Gene Exon Mapping

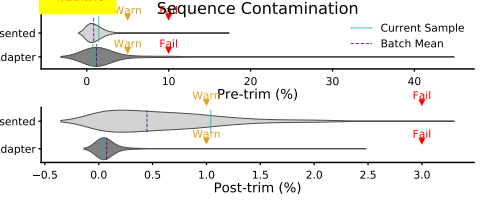


Ribosomal RNA

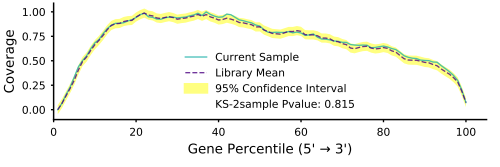


Sequence Contamination

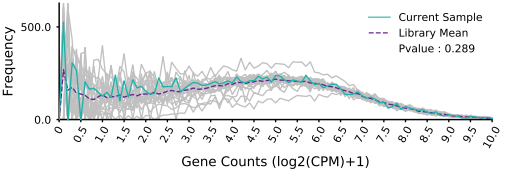
WARNING



GeneBody Coverage Distribution



Gene Expression

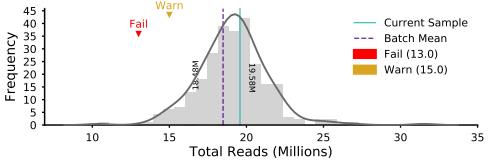


Sample : RNA_B11_06 Batch : batch_11

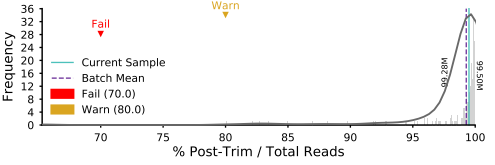
Warn cutoffs: Default alpha =0.8| Sequencing Depth = 15.0| Trimming = 80.0| Alignment = 50.0| Gene Exon Mapping = 30.0| Ribosomal RNA = 0.312| Sequence Contamination = 5.0| Gene Body Coverage = 0.8| Distribution of Gene Expression = 0.8

Fail cutoffs: Default alpha =0.9| Sequencing Depth = 13.0| Trimming = 70.0| Alignment = 10.0| Gene Exon Mapping = 15.0| Ribosomal RNA = 0.348| Sequence Contamination = 10.0| Gene Body Coverage = 0.9| Distribution of Gene Expression = 0.9

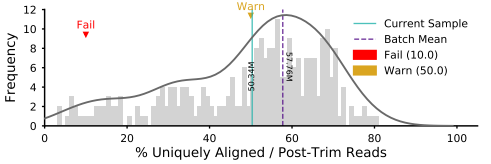
Sequencing Depth



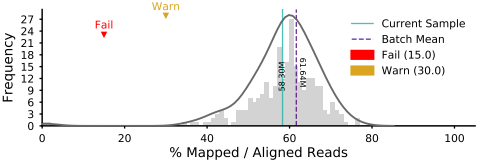
Trimming



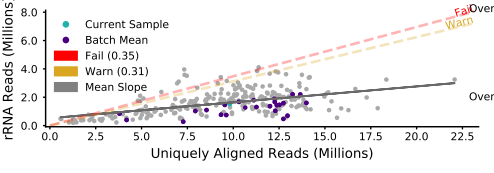
Alignment



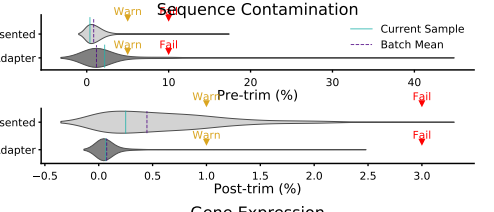
Gene Exon Mapping



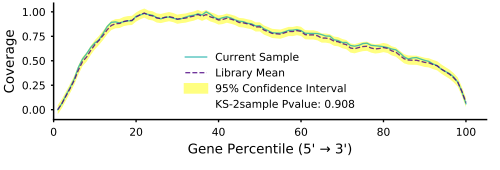
Ribosomal RNA



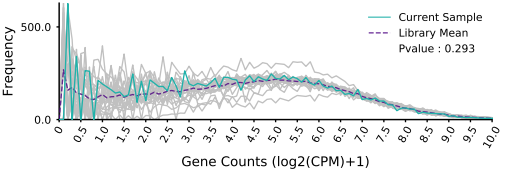
Sequence Contamination



GeneBody Coverage Distribution



Gene Expression

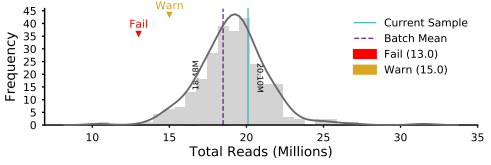


Sample : RNA_B11_07 Batch : batch_11

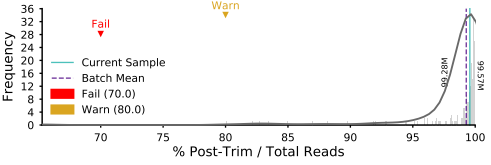
Warn cutoffs: Default alpha =0.8| Sequencing Depth = 15.0| Trimming = 80.0| Alignment = 50.0| Gene Exon Mapping = 30.0| Ribosomal RNA = 0.312| Sequence Contamination = 5.0| Gene Body Coverage = 0.8| Distribution of Gene Expression = 0.8

Fail cutoffs: Default alpha =0.9| Sequencing Depth = 13.0| Trimming = 70.0| Alignment = 10.0| Gene Exon Mapping = 15.0| Ribosomal RNA = 0.348| Sequence Contamination = 10.0| Gene Body Coverage = 0.9| Distribution of Gene Expression = 0.9

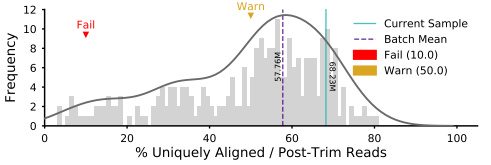
Sequencing Depth



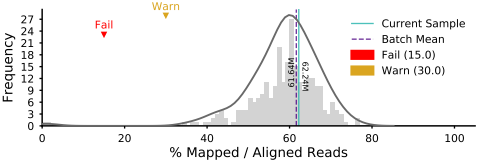
Trimming



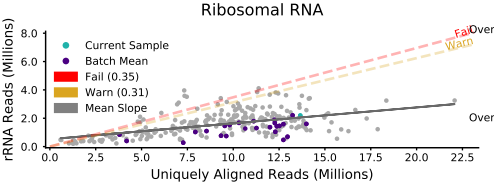
Alignment



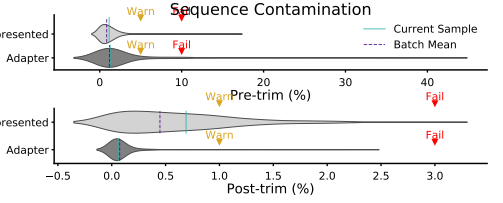
Gene Exon Mapping



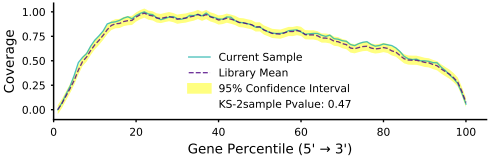
Ribosomal RNA



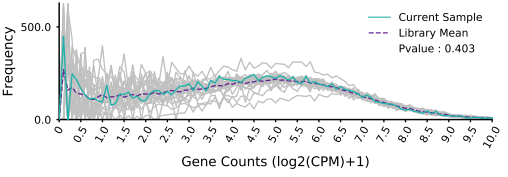
Sequence Contamination



GeneBody Coverage Distribution



Gene Expression

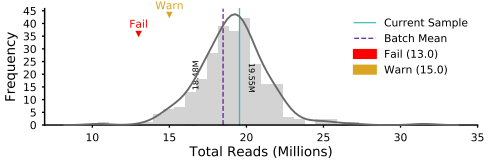


Sample : RNA_B11_08 Batch : batch_11

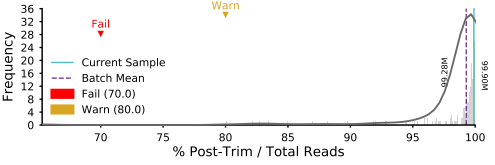
Warn cutoffs: Default alpha =0.8| Sequencing Depth = 15.0| Trimming = 80.0| Alignment = 50.0| Gene Exon Mapping = 30.0| Ribosomal RNA = 0.312| Sequence Contamination = 5.0| Gene Body Coverage = 0.8| Distribution of Gene Expression = 0.8

Fail cutoffs: Default alpha =0.9| Sequencing Depth = 13.0| Trimming = 70.0| Alignment = 10.0| Gene Exon Mapping = 15.0| Ribosomal RNA = 0.348| Sequence Contamination = 10.0| Gene Body Coverage = 0.9| Distribution of Gene Expression = 0.9

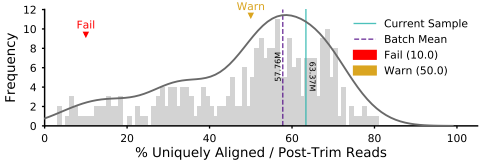
Sequencing Depth



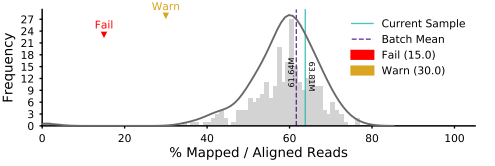
Trimming



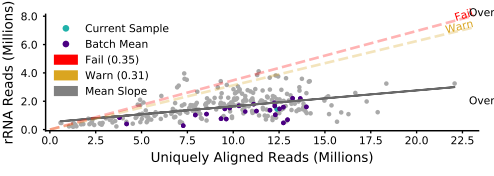
Alignment



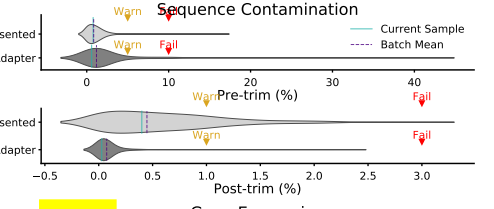
Gene Exon Mapping



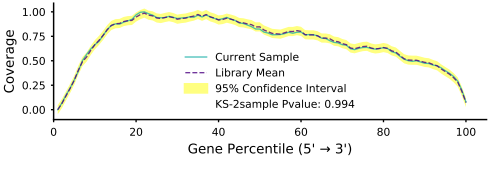
Ribosomal RNA



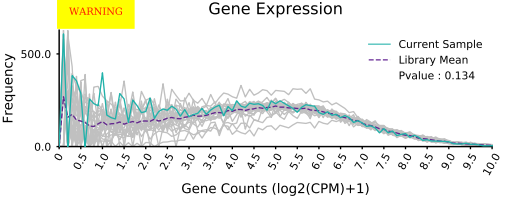
Sequence Contamination



GeneBody Coverage Distribution



Gene Expression

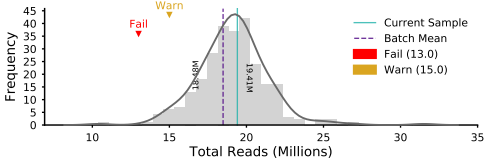


Sample : RNA_B11_09 Batch : batch_11

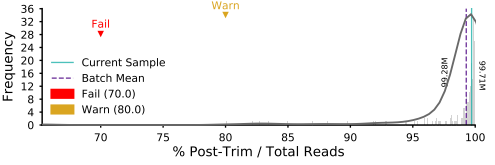
Warn cutoffs: Default alpha =0.8| Sequencing Depth = 15.0| Trimming = 80.0| Alignment = 50.0| Gene Exon Mapping = 30.0| Ribosomal RNA = 0.312| Sequence Contamination = 5.0| Gene Body Coverage = 0.8| Distribution of Gene Expression = 0.8

Fail cutoffs: Default alpha =0.9| Sequencing Depth = 13.0| Trimming = 70.0| Alignment = 10.0| Gene Exon Mapping = 15.0| Ribosomal RNA = 0.348| Sequence Contamination = 10.0| Gene Body Coverage = 0.9| Distribution of Gene Expression = 0.9

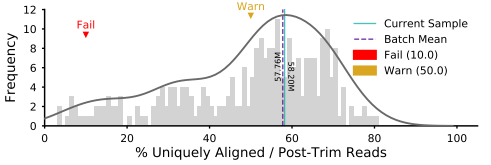
Sequencing Depth



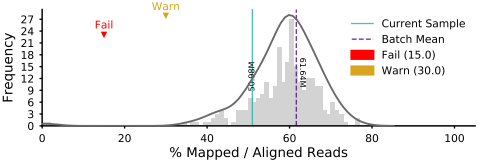
Trimming



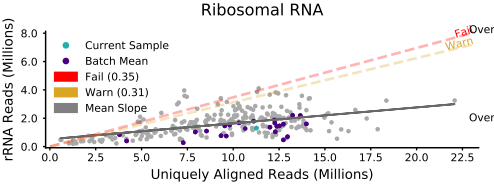
Alignment



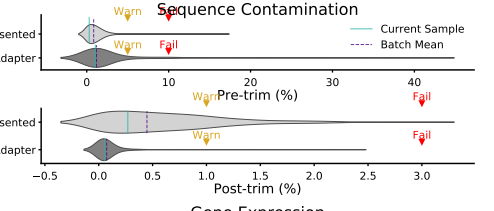
Gene Exon Mapping



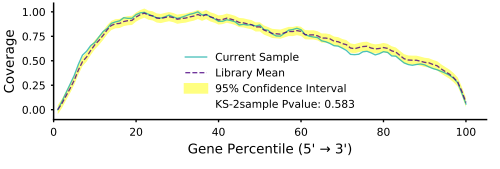
Ribosomal RNA



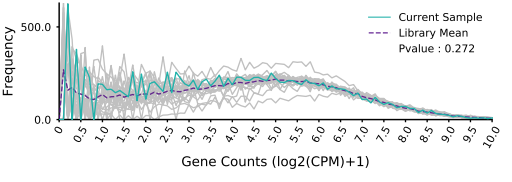
Sequence Contamination



GeneBody Coverage Distribution



Gene Expression

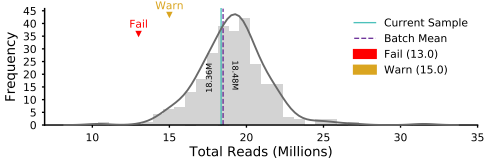


Sample : RNA_B11_10 Batch : batch_11

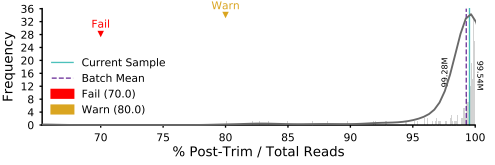
Warn cutoffs: Default alpha =0.8| Sequencing Depth = 15.0| Trimming = 80.0| Alignment = 50.0| Gene Exon Mapping = 30.0| Ribosomal RNA = 0.312| Sequence Contamination = 5.0| Gene Body Coverage = 0.8| Distribution of Gene Expression = 0.8

Fail cutoffs: Default alpha =0.9| Sequencing Depth = 13.0| Trimming = 70.0| Alignment = 10.0| Gene Exon Mapping = 15.0| Ribosomal RNA = 0.348| Sequence Contamination = 10.0| Gene Body Coverage = 0.9| Distribution of Gene Expression = 0.9

Sequencing Depth

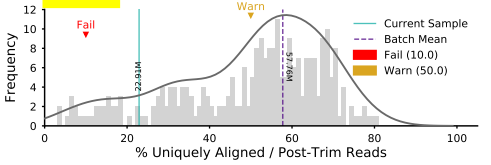


Trimming

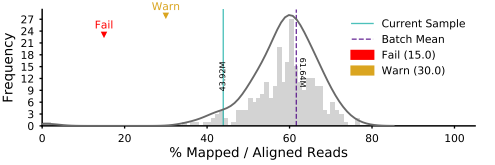


WARNING

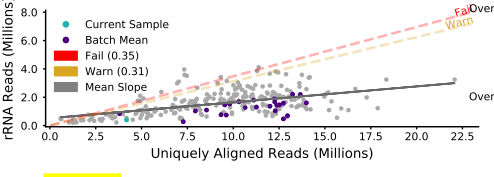
Alignment



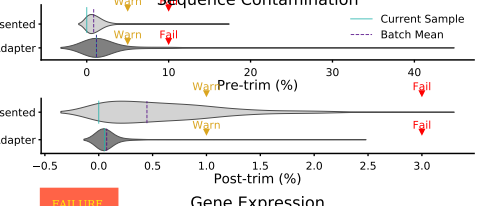
Gene Exon Mapping



Ribosomal RNA

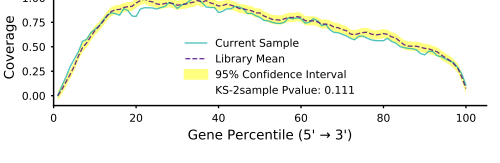


Sequence Contamination



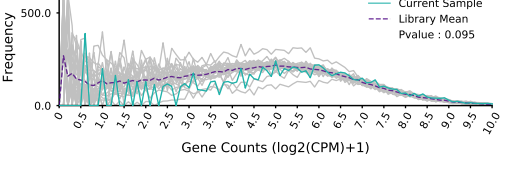
WARNING

GeneBody Coverage Distribution



FAILURE

Gene Expression

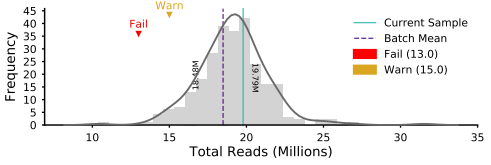


Sample : RNA_B11_11 Batch : batch_11

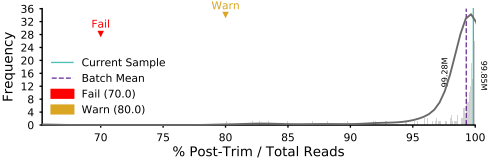
Warn cutoffs: Default alpha =0.8| Sequencing Depth = 15.0| Trimming = 80.0| Alignment = 50.0| Gene Exon Mapping = 30.0| Ribosomal RNA = 0.312| Sequence Contamination = 5.0| Gene Body Coverage = 0.8| Distribution of Gene Expression = 0.8

Fail cutoffs: Default alpha =0.9| Sequencing Depth = 13.0| Trimming = 70.0| Alignment = 10.0| Gene Exon Mapping = 15.0| Ribosomal RNA = 0.348| Sequence Contamination = 10.0| Gene Body Coverage = 0.9| Distribution of Gene Expression = 0.9

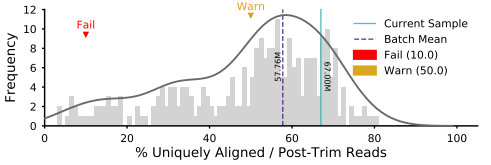
Sequencing Depth



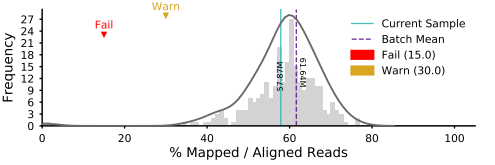
Trimming



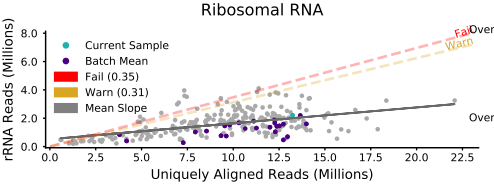
Alignment



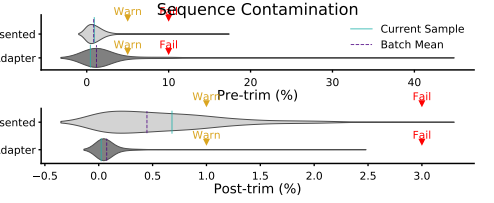
Gene Exon Mapping



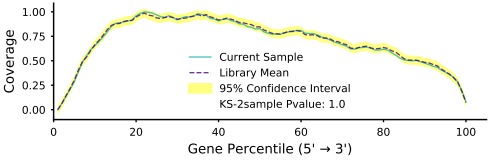
Ribosomal RNA



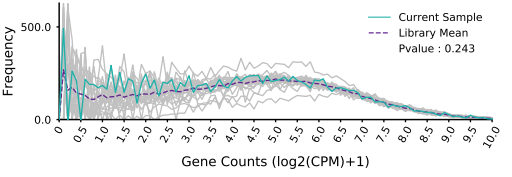
Sequence Contamination



GeneBody Coverage Distribution



Gene Expression

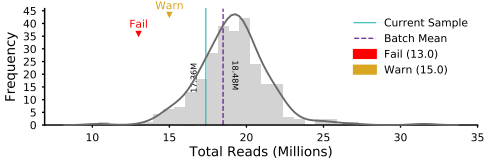


Sample : RNA_B11_12 Batch : batch_11

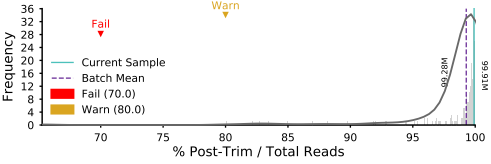
Warn cutoffs: Default alpha =0.8| Sequencing Depth = 15.0| Trimming = 80.0| Alignment = 50.0| Gene Exon Mapping = 30.0| Ribosomal RNA = 0.312| Sequence Contamination = 5.0| Gene Body Coverage = 0.8| Distribution of Gene Expression = 0.8

Fail cutoffs: Default alpha =0.9| Sequencing Depth = 13.0| Trimming = 70.0| Alignment = 10.0| Gene Exon Mapping = 15.0| Ribosomal RNA = 0.348| Sequence Contamination = 10.0| Gene Body Coverage = 0.9| Distribution of Gene Expression = 0.9

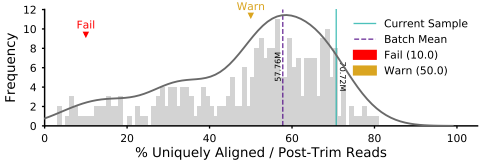
Sequencing Depth



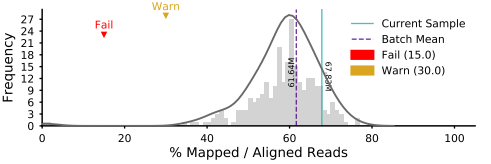
Trimming



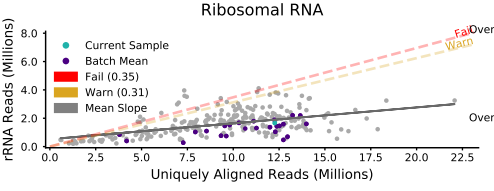
Alignment



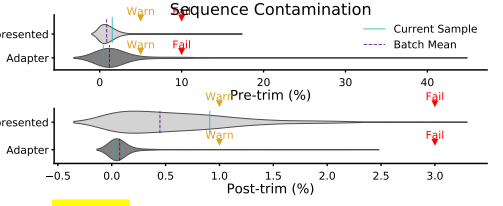
Gene Exon Mapping



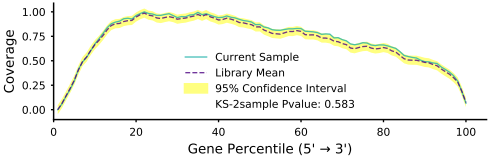
Ribosomal RNA



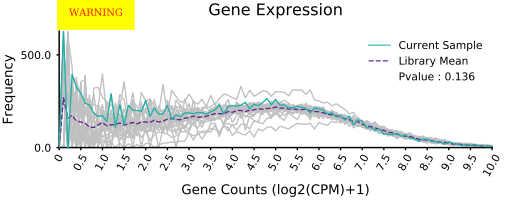
Sequence Contamination



GeneBody Coverage Distribution



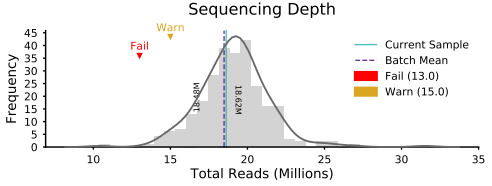
Gene Expression

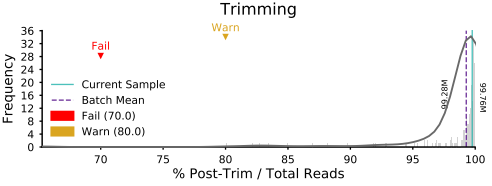


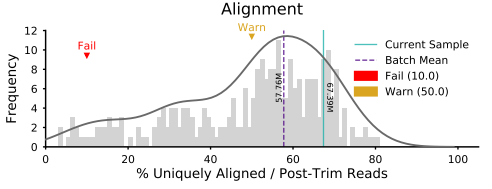
Sample : RNA_B11_13 Batch : batch_11

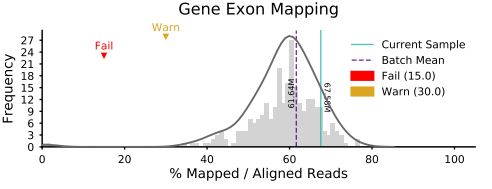
Warn cutoffs: Default alpha =0.8| Sequencing Depth = 15.0| Trimming = 80.0| Alignment = 50.0| Gene Exon Mapping = 30.0| Ribosomal RNA = 0.312| Sequence Contamination = 5.0| Gene Body Coverage = 0.8| Distribution of Gene Expression = 0.8

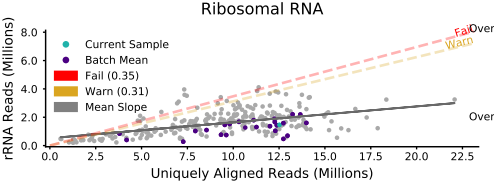
Fail cutoffs: Default alpha =0.9| Sequencing Depth = 13.0| Trimming = 70.0| Alignment = 10.0| Gene Exon Mapping = 15.0| Ribosomal RNA = 0.348| Sequence Contamination = 10.0| Gene Body Coverage = 0.9| Distribution of Gene Expression = 0.9

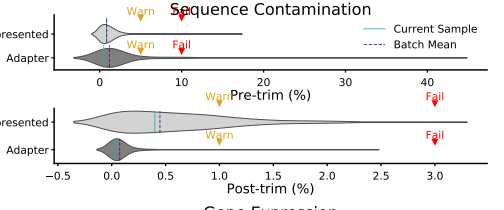


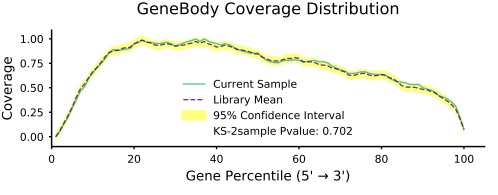


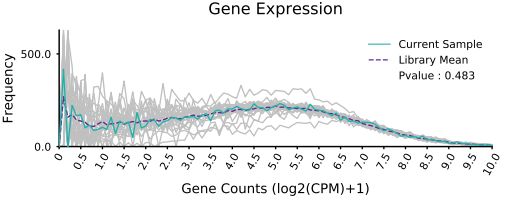










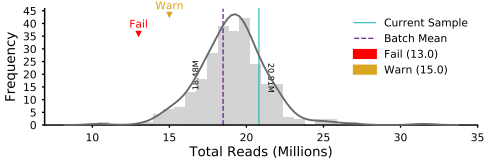


Sample : RNA_B11_14 Batch : batch_11

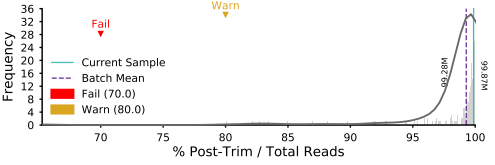
Warn cutoffs: Default alpha =0.8| Sequencing Depth = 15.0| Trimming = 80.0| Alignment = 50.0| Gene Exon Mapping = 30.0| Ribosomal RNA = 0.312| Sequence Contamination = 5.0| Gene Body Coverage = 0.8| Distribution of Gene Expression = 0.8

Fail cutoffs: Default alpha =0.9| Sequencing Depth = 13.0| Trimming = 70.0| Alignment = 10.0| Gene Exon Mapping = 15.0| Ribosomal RNA = 0.348| Sequence Contamination = 10.0| Gene Body Coverage = 0.9| Distribution of Gene Expression = 0.9

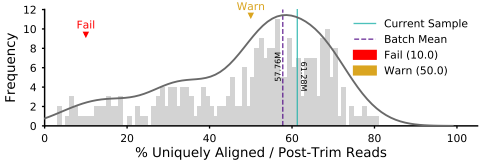
Sequencing Depth



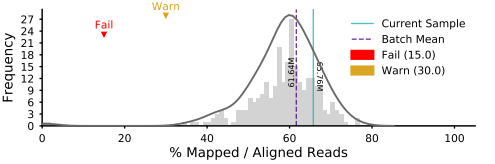
Trimming



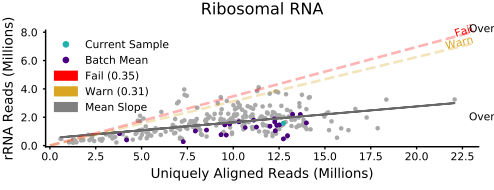
Alignment



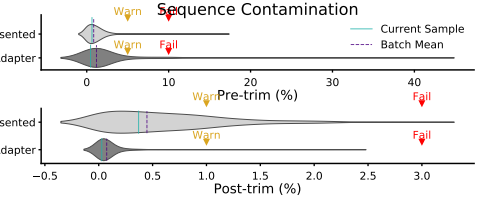
Gene Exon Mapping



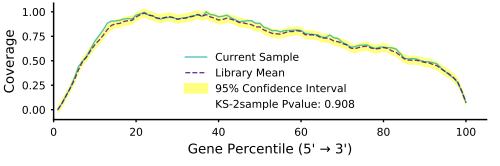
Ribosomal RNA



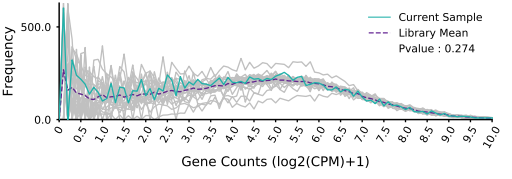
Sequence Contamination



GeneBody Coverage Distribution



Gene Expression

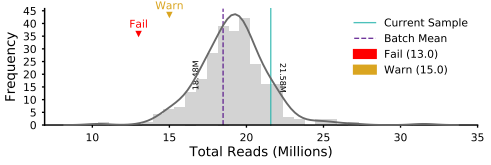


Sample : RNA_B11_15 Batch : batch_11

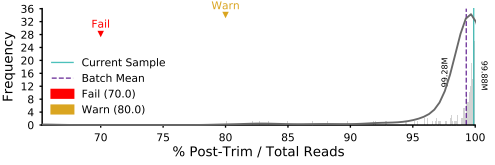
Warn cutoffs: Default alpha =0.8| Sequencing Depth = 15.0| Trimming = 80.0| Alignment = 50.0| Gene Exon Mapping = 30.0| Ribosomal RNA = 0.312| Sequence Contamination = 5.0| Gene Body Coverage = 0.8| Distribution of Gene Expression = 0.8

Fail cutoffs: Default alpha =0.9| Sequencing Depth = 13.0| Trimming = 70.0| Alignment = 10.0| Gene Exon Mapping = 15.0| Ribosomal RNA = 0.348| Sequence Contamination = 10.0| Gene Body Coverage = 0.9| Distribution of Gene Expression = 0.9

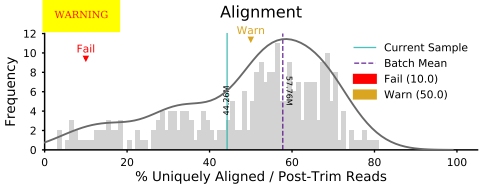
Sequencing Depth



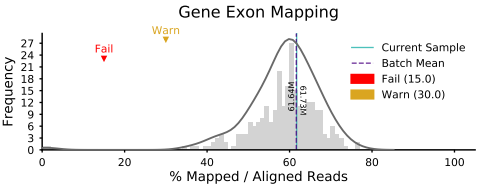
Trimming



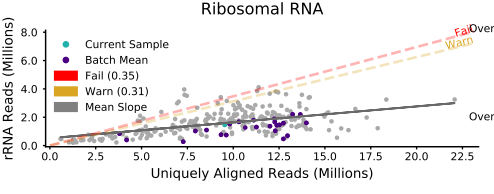
Alignment



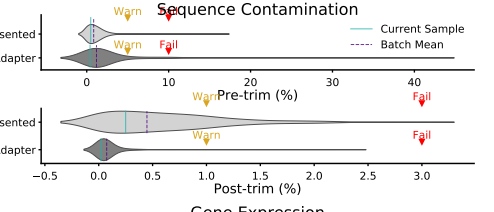
Gene Exon Mapping



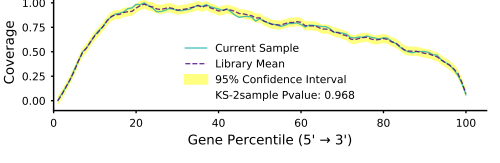
Ribosomal RNA



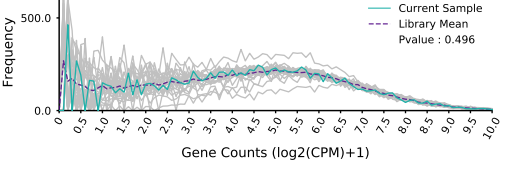
Sequence Contamination



GeneBody Coverage Distribution



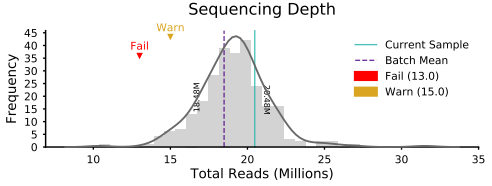
Gene Expression

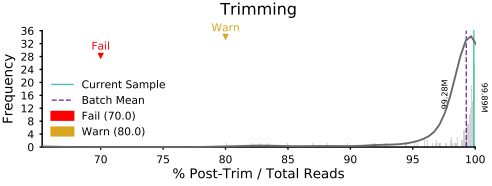


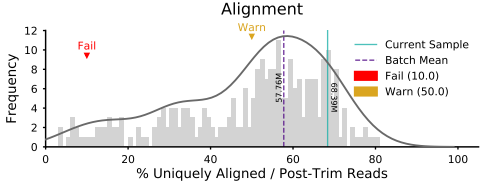
Sample : RNA_B11_16 Batch : batch_11

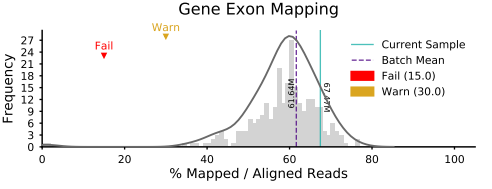
Warn cutoffs: Default alpha =0.8| Sequencing Depth = 15.0| Trimming = 80.0| Alignment = 50.0| Gene Exon Mapping = 30.0| Ribosomal RNA = 0.312| Sequence Contamination = 5.0| Gene Body Coverage = 0.8| Distribution of Gene Expression = 0.8

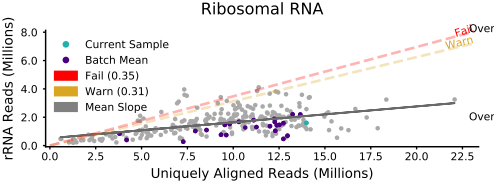
Fail cutoffs: Default alpha =0.9| Sequencing Depth = 13.0| Trimming = 70.0| Alignment = 10.0| Gene Exon Mapping = 15.0| Ribosomal RNA = 0.348| Sequence Contamination = 10.0| Gene Body Coverage = 0.9| Distribution of Gene Expression = 0.9

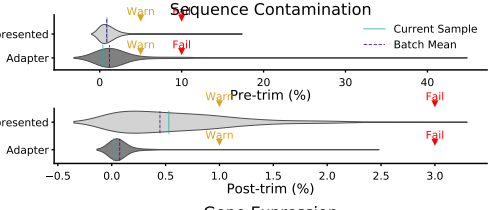


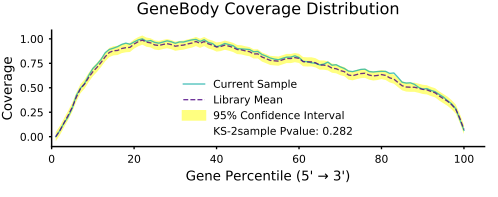


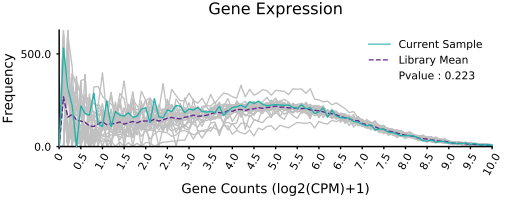










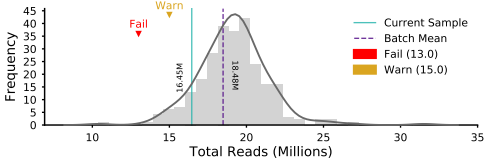


Sample : RNA_B11_17 Batch : batch_11

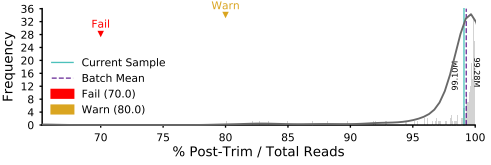
Warn cutoffs: Default alpha =0.8| Sequencing Depth = 15.0| Trimming = 80.0| Alignment = 50.0| Gene Exon Mapping = 30.0| Ribosomal RNA = 0.312| Sequence Contamination = 5.0| Gene Body Coverage = 0.8| Distribution of Gene Expression = 0.8

Fail cutoffs: Default alpha =0.9| Sequencing Depth = 13.0| Trimming = 70.0| Alignment = 10.0| Gene Exon Mapping = 15.0| Ribosomal RNA = 0.348| Sequence Contamination = 10.0| Gene Body Coverage = 0.9| Distribution of Gene Expression = 0.9

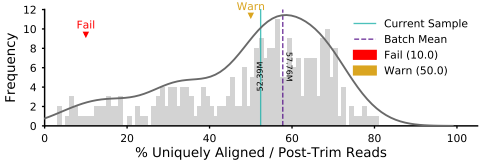
Sequencing Depth



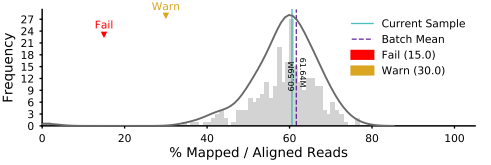
Trimming



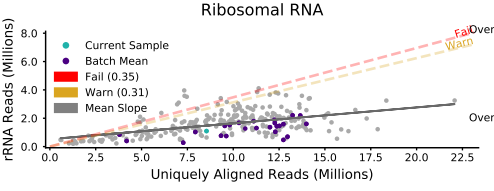
Alignment



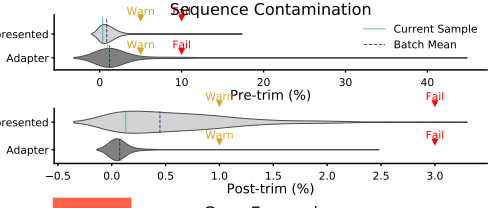
Gene Exon Mapping



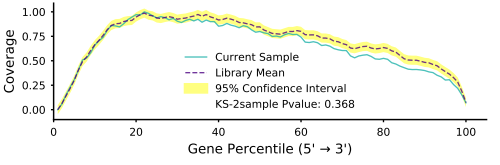
Ribosomal RNA



Sequence Contamination

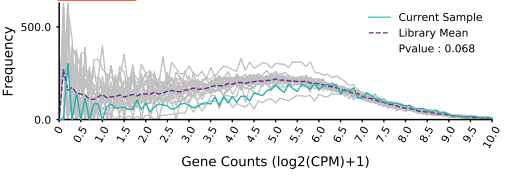


GeneBody Coverage Distribution



Gene Expression

FAILURE

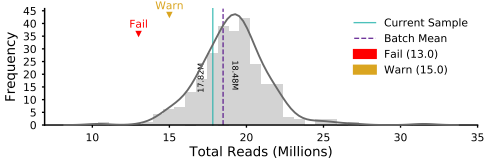


Sample : RNA_B11_18 Batch : batch_11

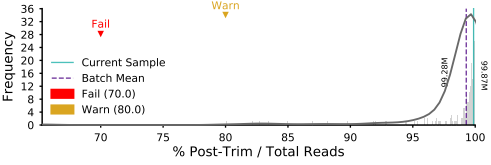
Warn cutoffs: Default alpha =0.8| Sequencing Depth = 15.0| Trimming = 80.0| Alignment = 50.0| Gene Exon Mapping = 30.0| Ribosomal RNA = 0.312| Sequence Contamination = 5.0| Gene Body Coverage = 0.8| Distribution of Gene Expression = 0.8

Fail cutoffs: Default alpha =0.9| Sequencing Depth = 13.0| Trimming = 70.0| Alignment = 10.0| Gene Exon Mapping = 15.0| Ribosomal RNA = 0.348| Sequence Contamination = 10.0| Gene Body Coverage = 0.9| Distribution of Gene Expression = 0.9

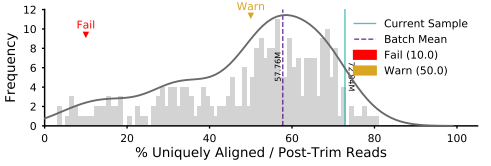
Sequencing Depth



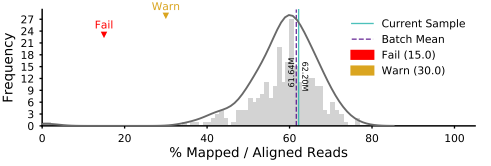
Trimming



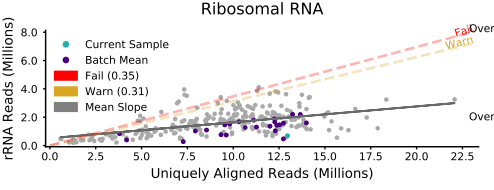
Alignment



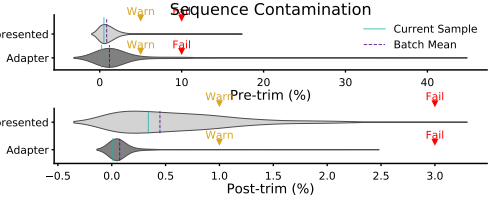
Gene Exon Mapping



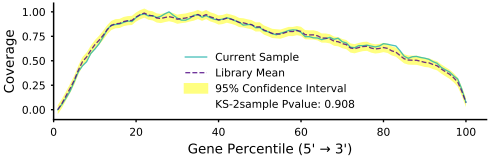
Ribosomal RNA



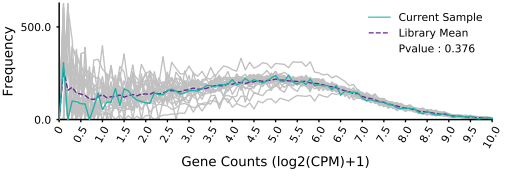
Sequence Contamination



GeneBody Coverage Distribution



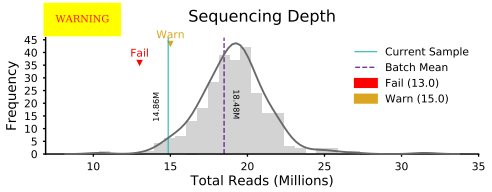
Gene Expression

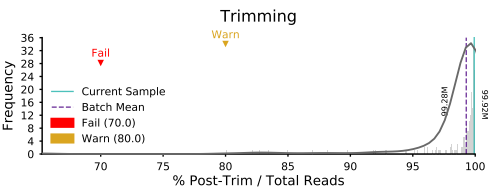


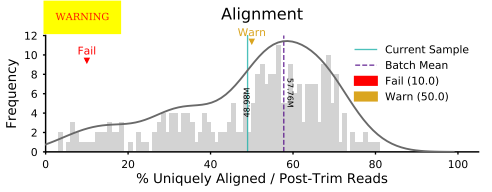
Sample : RNA_B11_19 Batch : batch_11

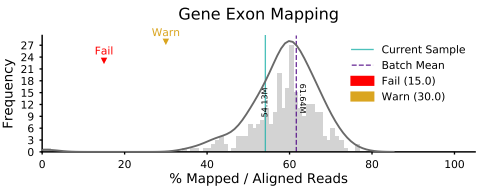
Warn cutoffs: Default alpha =0.8| Sequencing Depth = 15.0| Trimming = 80.0| Alignment = 50.0| Gene Exon Mapping = 30.0| Ribosomal RNA = 0.312| Sequence Contamination = 5.0| Gene Body Coverage = 0.8| Distribution of Gene Expression = 0.8

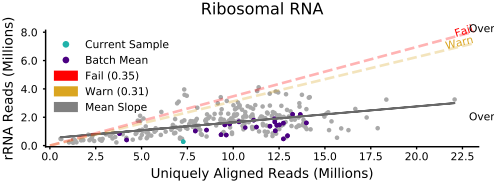
Fail cutoffs: Default alpha =0.9| Sequencing Depth = 13.0| Trimming = 70.0| Alignment = 10.0| Gene Exon Mapping = 15.0| Ribosomal RNA = 0.348| Sequence Contamination = 10.0| Gene Body Coverage = 0.9| Distribution of Gene Expression = 0.9

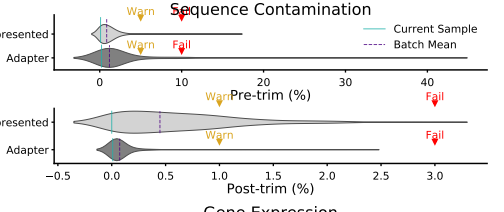


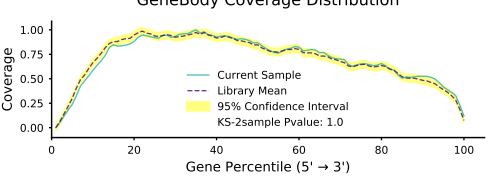


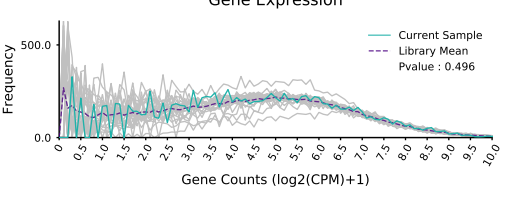










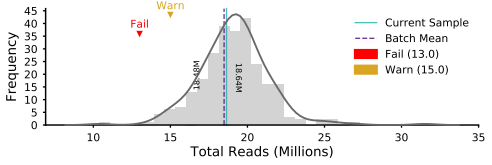


Sample : RNA_B11_20 Batch : batch_11

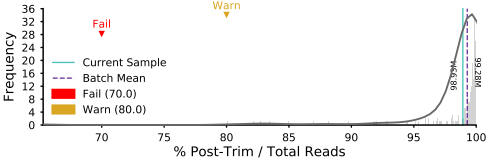
Warn cutoffs: Default alpha =0.8| Sequencing Depth = 15.0| Trimming = 80.0| Alignment = 50.0| Gene Exon Mapping = 30.0| Ribosomal RNA = 0.312| Sequence Contamination = 5.0| Gene Body Coverage = 0.8| Distribution of Gene Expression = 0.8

Fail cutoffs: Default alpha =0.9| Sequencing Depth = 13.0| Trimming = 70.0| Alignment = 10.0| Gene Exon Mapping = 15.0| Ribosomal RNA = 0.348| Sequence Contamination = 10.0| Gene Body Coverage = 0.9| Distribution of Gene Expression = 0.9

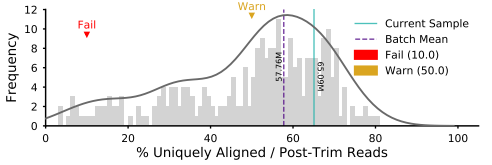
Sequencing Depth



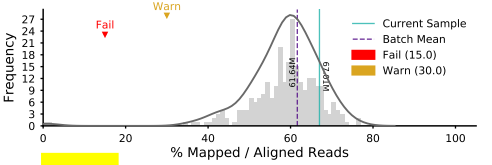
Trimming



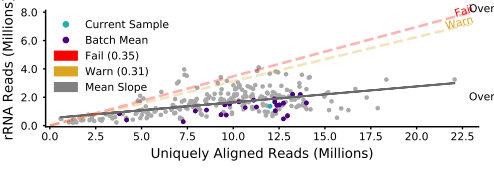
Alignment



Gene Exon Mapping

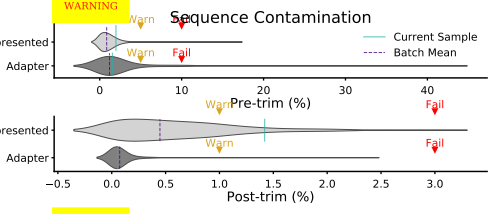


Ribosomal RNA

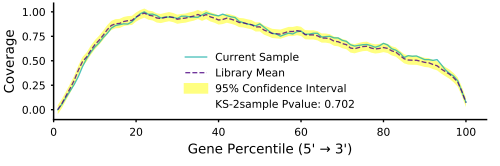


Sequence Contamination

WARNING

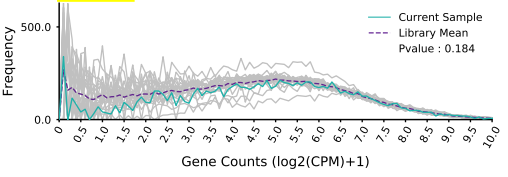


GeneBody Coverage Distribution



Gene Expression

WARNING

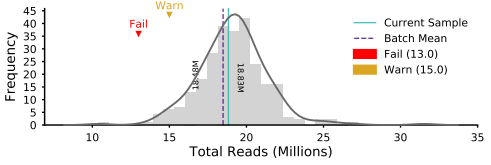


Sample : RNA_B11_21 Batch : batch_11

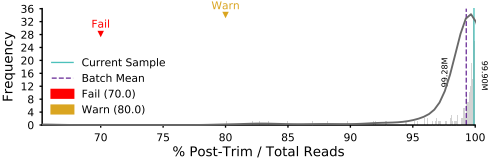
Warn cutoffs: Default alpha =0.8| Sequencing Depth = 15.0| Trimming = 80.0| Alignment = 50.0| Gene Exon Mapping = 30.0| Ribosomal RNA = 0.312| Sequence Contamination = 5.0| Gene Body Coverage = 0.8| Distribution of Gene Expression = 0.8

Fail cutoffs: Default alpha =0.9| Sequencing Depth = 13.0| Trimming = 70.0| Alignment = 10.0| Gene Exon Mapping = 15.0| Ribosomal RNA = 0.348| Sequence Contamination = 10.0| Gene Body Coverage = 0.9| Distribution of Gene Expression = 0.9

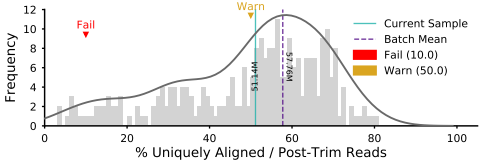
Sequencing Depth



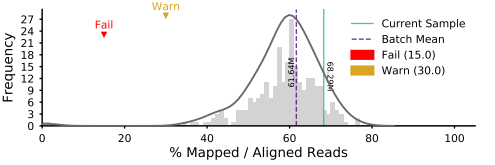
Trimming



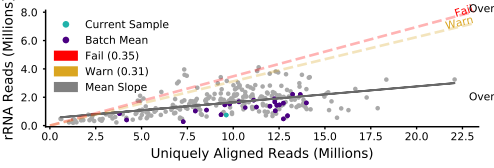
Alignment



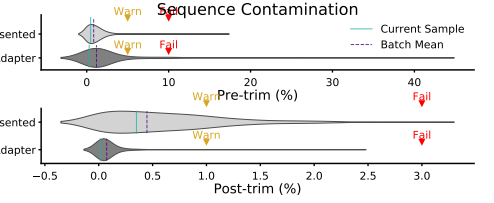
Gene Exon Mapping



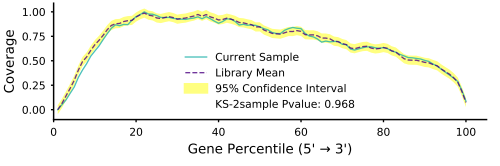
Ribosomal RNA



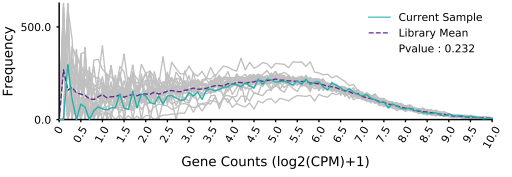
Sequence Contamination



GeneBody Coverage Distribution



Gene Expression

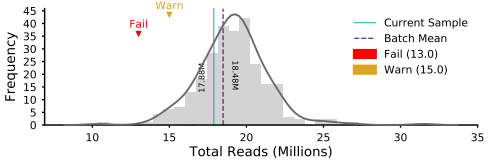


Sample : RNA_B11_22 Batch : batch_11

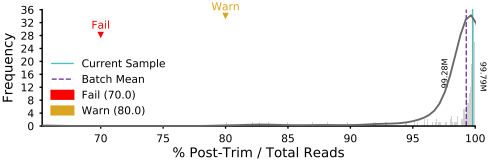
Warn cutoffs: Default alpha =0.8| Sequencing Depth = 15.0| Trimming = 80.0| Alignment = 50.0| Gene Exon Mapping = 30.0| Ribosomal RNA = 0.312| Sequence Contamination = 5.0| Gene Body Coverage = 0.8| Distribution of Gene Expression = 0.8

Fail cutoffs: Default alpha =0.9| Sequencing Depth = 13.0| Trimming = 70.0| Alignment = 10.0| Gene Exon Mapping = 15.0| Ribosomal RNA = 0.348| Sequence Contamination = 10.0| Gene Body Coverage = 0.9| Distribution of Gene Expression = 0.9

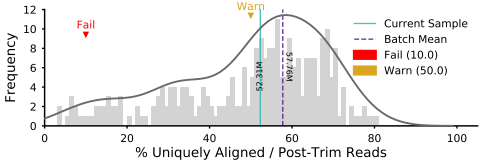
Sequencing Depth



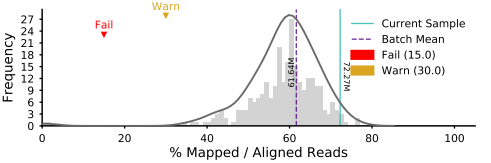
Trimming



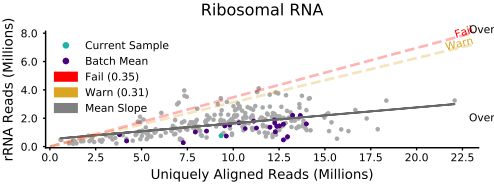
Alignment



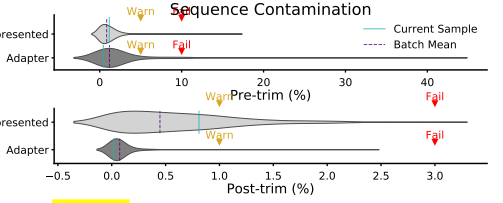
Gene Exon Mapping



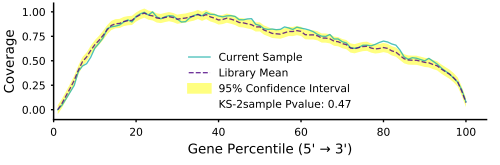
Ribosomal RNA



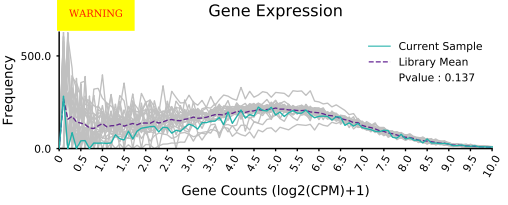
Sequence Contamination



GeneBody Coverage Distribution



Gene Expression

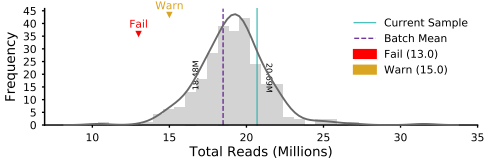


Sample : RNA_B11_23 Batch : batch_11

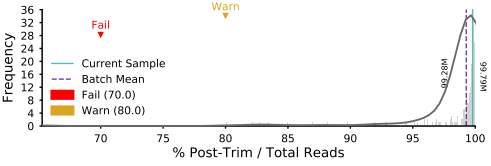
Warn cutoffs: Default alpha =0.8| Sequencing Depth = 15.0| Trimming = 80.0| Alignment = 50.0| Gene Exon Mapping = 30.0| Ribosomal RNA = 0.312| Sequence Contamination = 5.0| Gene Body Coverage = 0.8| Distribution of Gene Expression = 0.8

Fail cutoffs: Default alpha =0.9| Sequencing Depth = 13.0| Trimming = 70.0| Alignment = 10.0| Gene Exon Mapping = 15.0| Ribosomal RNA = 0.348| Sequence Contamination = 10.0| Gene Body Coverage = 0.9| Distribution of Gene Expression = 0.9

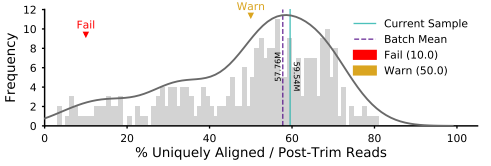
Sequencing Depth



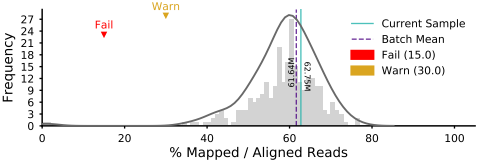
Trimming



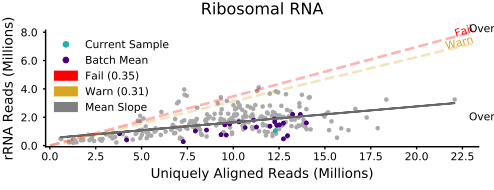
Alignment



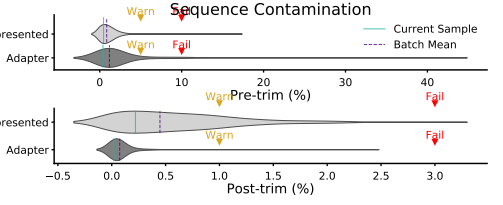
Gene Exon Mapping



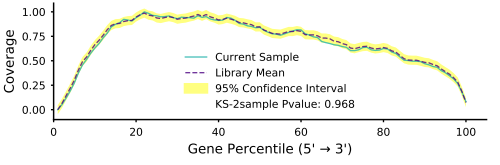
Ribosomal RNA



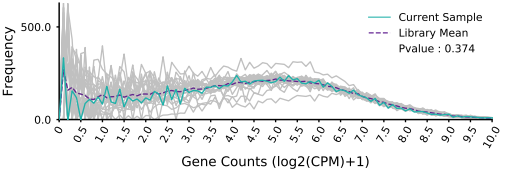
Sequence Contamination



GeneBody Coverage Distribution



Gene Expression

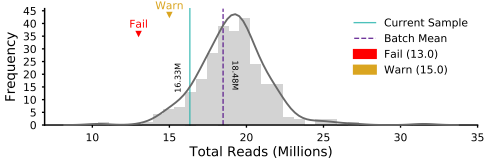


Sample : RNA_B11_24 Batch : batch_11

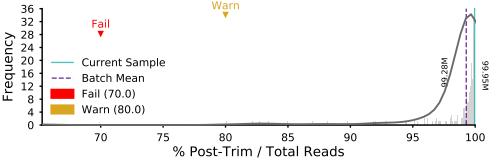
Warn cutoffs: Default alpha =0.8| Sequencing Depth = 15.0| Trimming = 80.0| Alignment = 50.0| Gene Exon Mapping = 30.0| Ribosomal RNA = 0.312| Sequence Contamination = 5.0| Gene Body Coverage = 0.8| Distribution of Gene Expression = 0.8

Fail cutoffs: Default alpha =0.9| Sequencing Depth = 13.0| Trimming = 70.0| Alignment = 10.0| Gene Exon Mapping = 15.0| Ribosomal RNA = 0.348| Sequence Contamination = 10.0| Gene Body Coverage = 0.9| Distribution of Gene Expression = 0.9

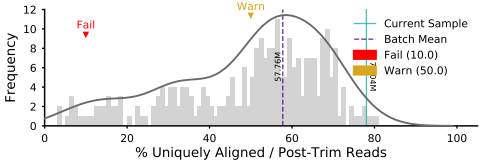
Sequencing Depth



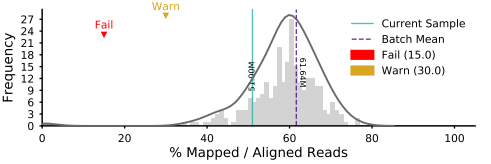
Trimming



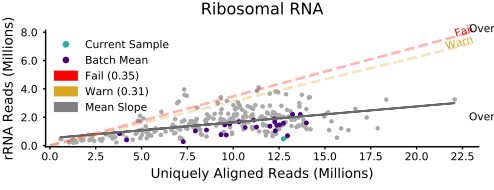
Alignment



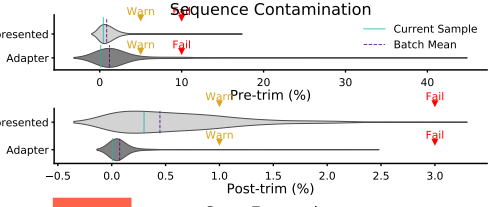
Gene Exon Mapping



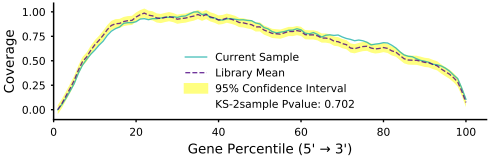
Ribosomal RNA



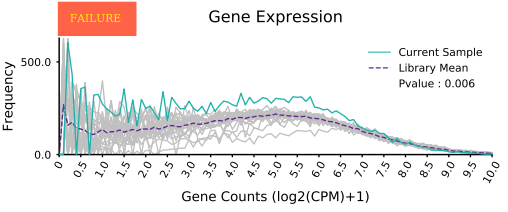
Sequence Contamination



GeneBody Coverage Distribution



Gene Expression

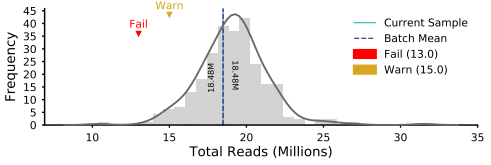


Sample : Batch_Mean Batch : batch_11

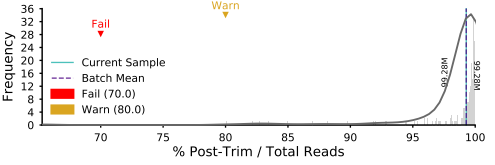
Warn cutoffs: Default alpha =0.8| Sequencing Depth = 15.0| Trimming = 80.0| Alignment = 50.0| Gene Exon Mapping = 30.0| Ribosomal RNA = 0.312| Sequence Contamination = 5.0| Gene Body Coverage = 0.8| Distribution of Gene Expression = 0.8

Fail cutoffs: Default alpha =0.9| Sequencing Depth = 13.0| Trimming = 70.0| Alignment = 10.0| Gene Exon Mapping = 15.0| Ribosomal RNA = 0.348| Sequence Contamination = 10.0| Gene Body Coverage = 0.9| Distribution of Gene Expression = 0.9

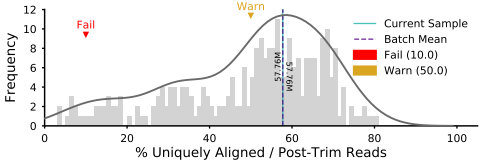
Sequencing Depth



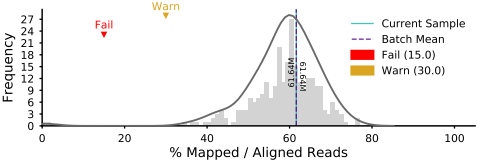
Trimming



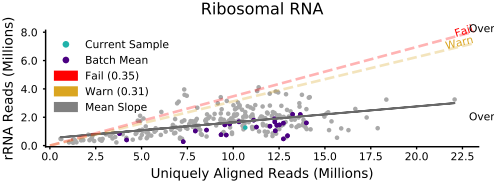
Alignment



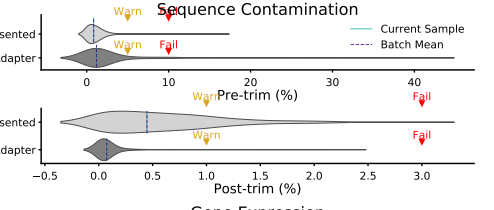
Gene Exon Mapping



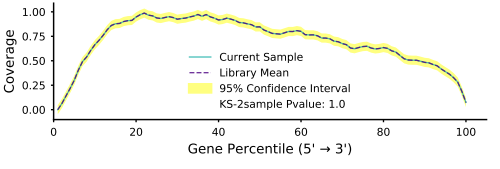
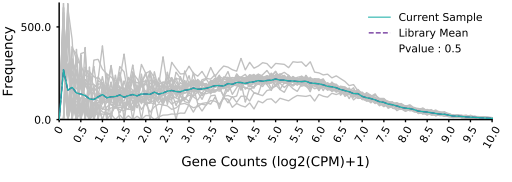
Ribosomal RNA



Sequence Contamination



GeneBody Coverage Distribution



Gene Expression