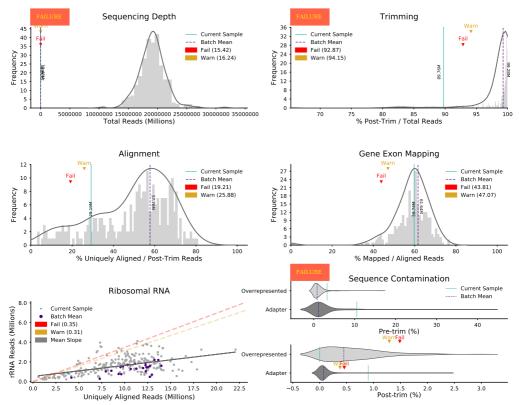
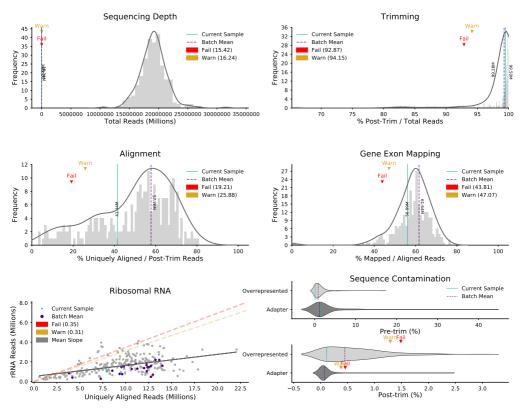
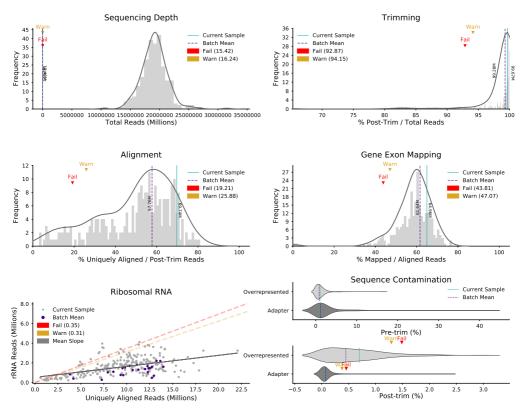
Warn cutoffs: Default alpha = 0.8] Sequencing Depth = 16.241476381729186| Trimming = 94.1462290640725| Algoment = 25.88342206791934| Gene Evon Mapping = 47.068386642306564| Ribosomal RNA = 0.312| Sequence Contamination = 0.37816525422038655| Gene Body Coverage = 0.8| Fail cutoffs: Default alpha = 0.9| Sequencing Depth = 15.422129911096862| Trimming = 92.86918155320676| Alignment = 19.211411756396192| Gene Exon Mapping = 43.813950731314556| Ribosomal RNA = 0.348| Sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 10.960615219281971| Gene Body Coverage = 10.96061

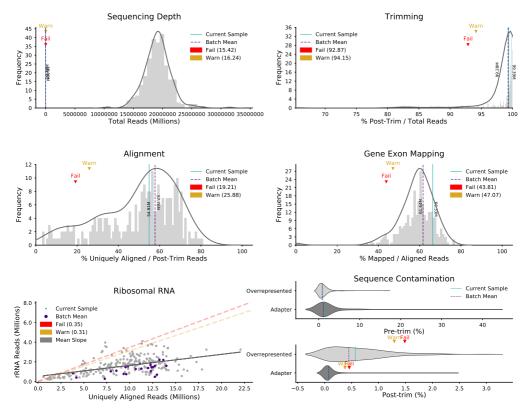




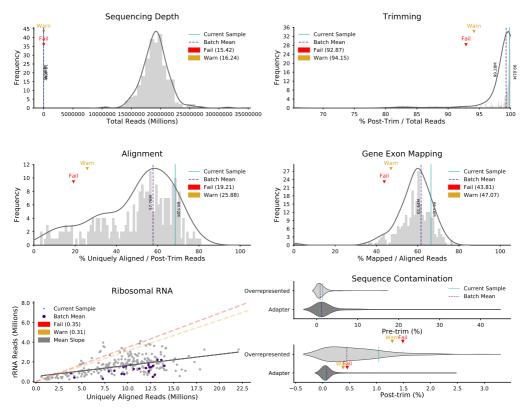
War durith: Default alpha = 0.81, Sequencing Depth = 16.2414783812198687, 1971 (April 2014) (Apr

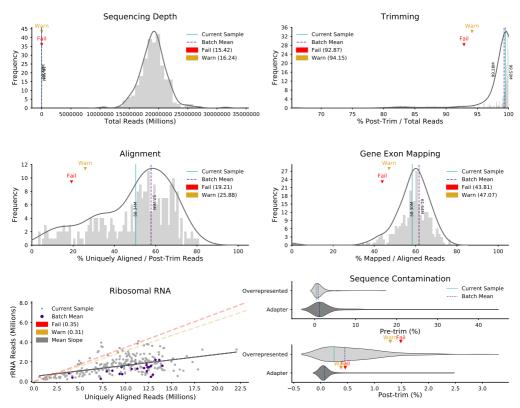


Warn cutoffs: Default alpha = 0.8] Sequencing Depth = 16.241476381729186| Trimming = 94.1462290640725| Algoment = 25.88342206791934| Gene Evon Mapping = 47.068386642306564| Ribosomal RNA = 0.312| Sequence Contamination = 0.37816525422038655| Gene Body Coverage = 0.8| Fail cutoffs: Default alpha = 0.9| Sequencing Depth = 15.422129911096862| Trimming = 92.86918155320676| Alignment = 19.211411756396192| Gene Exon Mapping = 43.813950731314556| Ribosomal RNA = 0.348| Sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 10.960615219281971| Gene Body Coverage = 10.96061

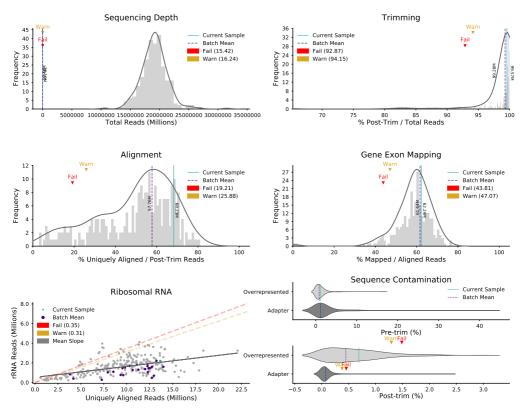


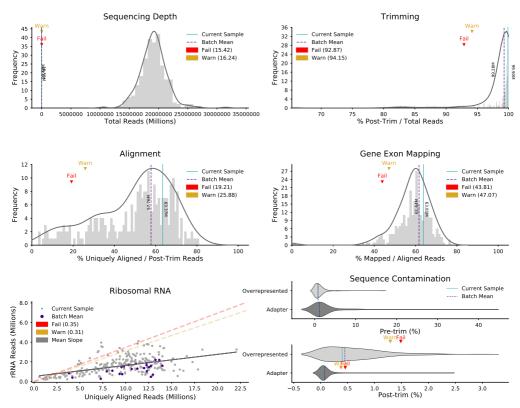
Warn cutoffs: Default alpha = 0.8] Sequencing Depth = 16.241476381729186| Trimming = 94.1462290640725| Algoment = 25.88342206791934| Gene Evon Mapping = 47.068386642306564| Ribosomal RNA = 0.312| Sequence Contamination = 0.37816525422038655| Gene Body Coverage = 0.8| Fail cutoffs: Default alpha = 0.9| Sequencing Depth = 15.422129911096862| Trimming = 92.86918155320676| Alignment = 19.211411756396192| Gene Exon Mapping = 43.813950731314556| Ribosomal RNA = 0.348| Sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 10.960615219281971| Gene Body Coverage = 10.96061



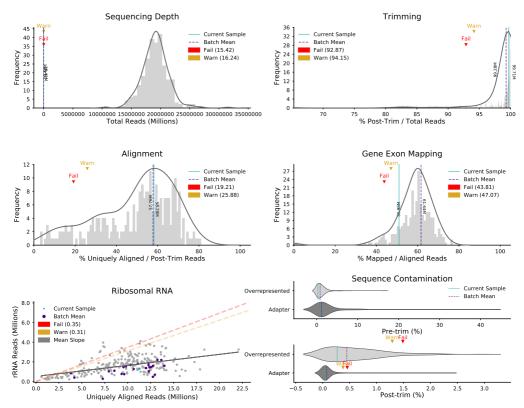


War durith: Default alpha = 0.81, Sequencing Depth = 16.2414783812198687, 1971 (April 2014) (Apr

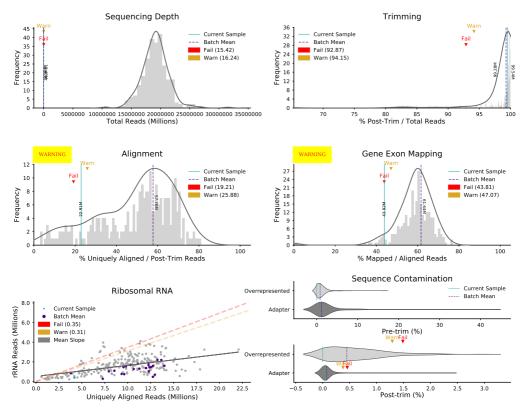


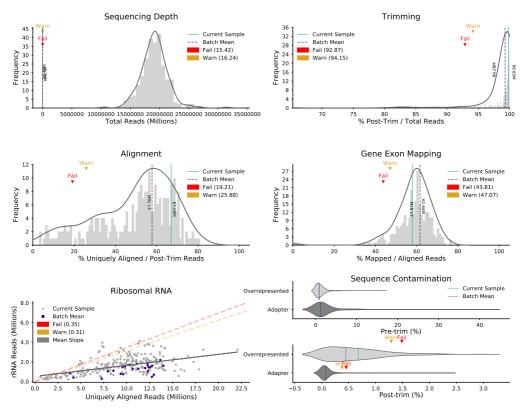


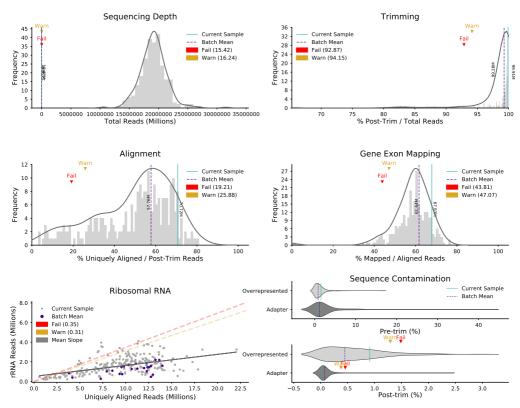
Warn cutoffs: Default alpha = 0.8] Sequencing Depth = 16.241476381729186| Trimming = 94.1462290640725| Algoment = 25.88342206791934| Gene Evon Mapping = 47.068386642306564| Ribosomal RNA = 0.312| Sequence Contamination = 0.37816525422038655| Gene Body Coverage = 0.8| Fail cutoffs: Default alpha = 0.9| Sequencing Depth = 15.422129911096862| Trimming = 92.86918155320676| Alignment = 19.211411756396192| Gene Exon Mapping = 43.813950731314556| Ribosomal RNA = 0.348| Sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 0.9| Direction of the contamination = 10.860615219281971| Gene Body Coverage = 10.960615219281971| Gene Body Coverage = 10.96061



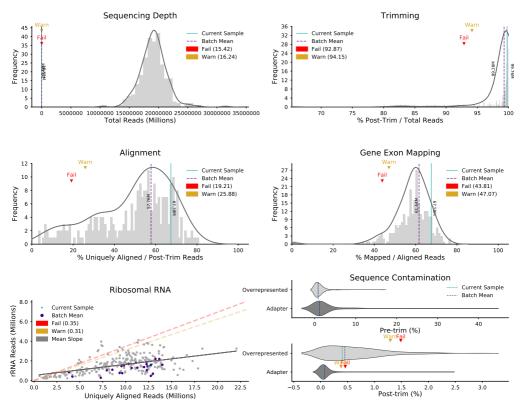
Warm cutoffs: Default alpha = 0.8| Sequencing Depth = 16.241476381729186| Trimming = 94.1462290640725| Alignment = 25.883432206791934| Gene Exon Mapping = 47.068386642306564| Ribosomal RNA = 0.312| Sequence Contamination = 0.37816525422038655| Gene Body Coverage = 0.8| In Fail Cutoffs: Default alpha = 0.9| Sequencing Depth = 15.42212991109862| Trimming = 92.86918155320676| Alignment = 19.211411756396192| Gene Exon Mapping = 43.813950731314556| Ribosomal RNA = 0.348| Sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence of the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence of the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to

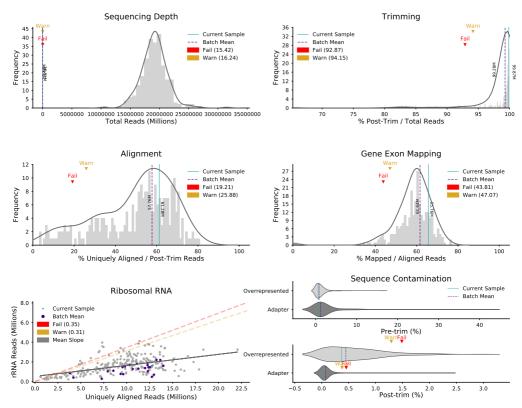


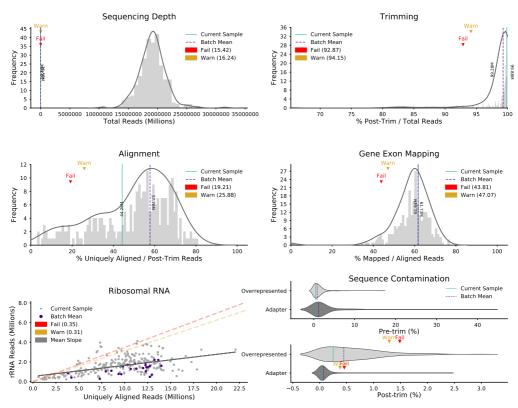


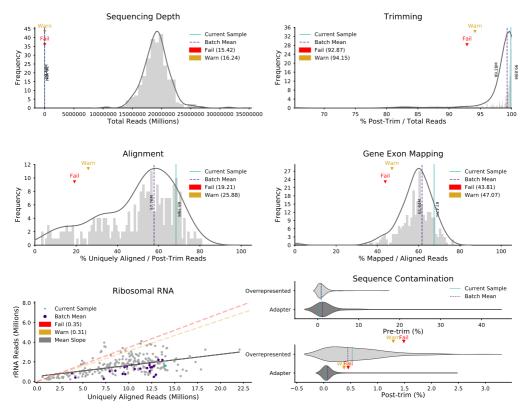


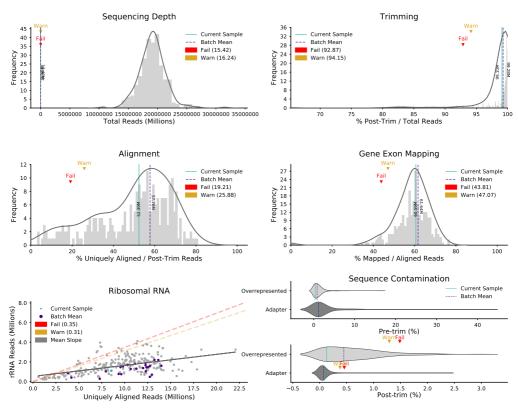
Warm cutoffs: Default alpha = 0.8| Sequencing Depth = 16.241476381729186| Trimming = 94.1462290640725| Alignment = 25.883432206791934| Gene Exon Mapping = 47.068386642306564| Ribosomal RNA = 0.312| Sequence Contamination = 0.37816525422038655| Gene Body Coverage = 0.8| Linear Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Display to the sequence Contamination = 10.860615219281971| Gene Body Cov

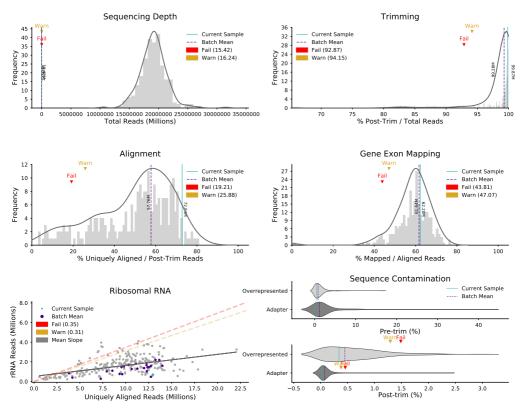




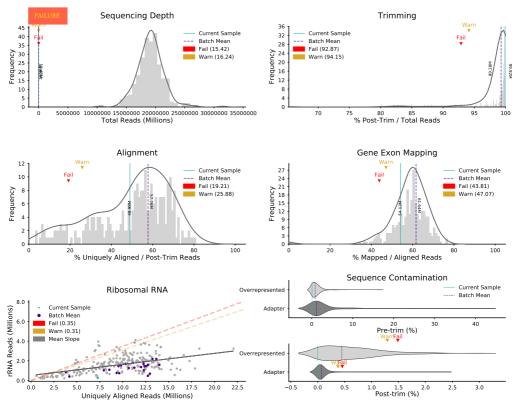




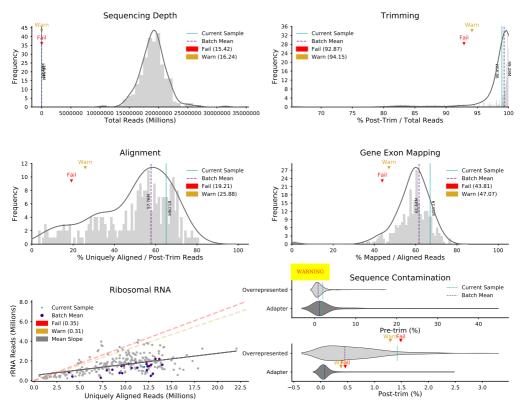


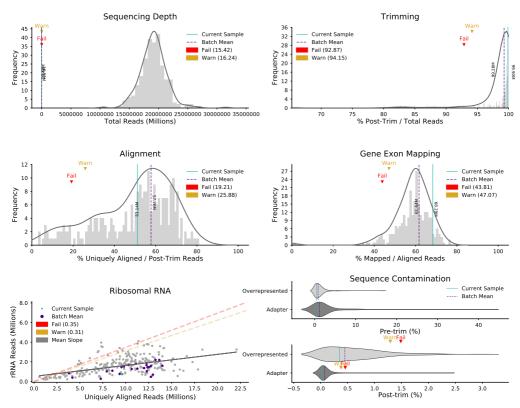


Warm cutoffs: Default alpha = 0.8| Sequencing Depth = 1.6.241476381729186| Trimming = 94.146229040725| Alignment = 25.883432206791934| Gene Exon Mapping = 47.068386642306564| Ribosomal RNA = 0.312| Sequence Contamination = 0.37816252422038655| Gene Body Coverage = 0.8| Life Fall cutoffs: Default alpha = 0.9| Sequencing Depth = 15.422129911096862| Trimming = 92.86918155320676| Alignment = 19.211411756396192| Gene Exon Mapping = 43.813950731314556| Ribosomal RNA = 0.348| Sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Dis

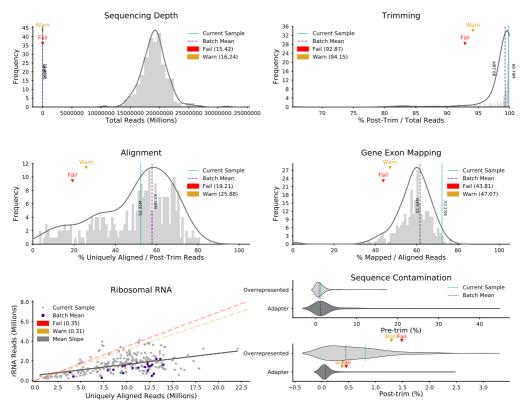


Warm cutoffs: Default alpha = 0.8| Sequencing Depth = 1.6.241476381729186| Trimming = 94.146229040725| Alignment = 25.883432206791934| Gene Exon Mapping = 47.068386642306564| Ribosomal RNA = 0.312| Sequence Contamination = 0.37816252422038655| Gene Body Coverage = 0.8| Life Fall cutoffs: Default alpha = 0.9| Sequencing Depth = 15.422129911096862| Trimming = 92.86918155320676| Alignment = 19.211411756396192| Gene Exon Mapping = 43.813950731314556| Ribosomal RNA = 0.348| Sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Dis

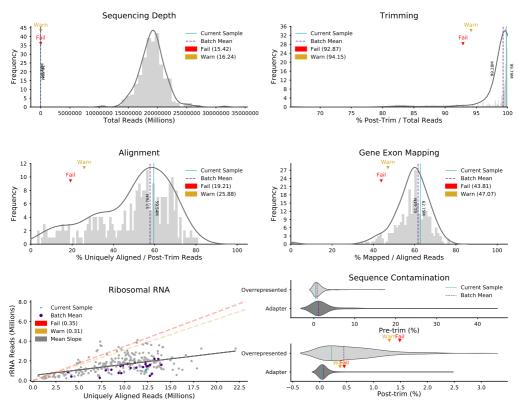




Warm cutoffs: Default alpha = 0.8| Sequencing Depth = 1.6.241476381729186| Trimming = 94.146229040725| Alignment = 25.883432206791934| Gene Exon Mapping = 47.068386642306564| Ribosomal RNA = 0.312| Sequence Contamination = 0.37816252422038655| Gene Body Coverage = 0.8| Life Fall cutoffs: Default alpha = 0.9| Sequencing Depth = 15.422129911096862| Trimming = 92.86918155320676| Alignment = 19.211411756396192| Gene Exon Mapping = 43.813950731314556| Ribosomal RNA = 0.348| Sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Dis



Warm cutoffs: Default alpha = 0.8| Sequencing Depth = 1.6.241476381729186| Trimming = 94.1462290640725| Alignment = 25.883432206791934| Gene Exon Mapping = 47.068386642306564| Ribosomal RNA = 0.312| Sequence Contamination = 0.37816525422033655| Gene Body Coverage = 0.8| Default alpha = 0.9| Sequencing Depth = 15.422129911096862| Trimming = 92.86918155320676| Alignment = 19.211411756396192| Gene Exon Mapping = 43.813950731314556| Ribosomal RNA = 0.348| Sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Di



Warm cutoffs: Default alpha = 0.8| Sequencing Depth = 1.6.241476381729186| Trimming = 94.146229040725| Alignment = 25.883432206791934| Gene Exon Mapping = 47.068386642306564| Ribosomal RNA = 0.312| Sequence Contamination = 0.37816252422038655| Gene Body Coverage = 0.8| Life Fall cutoffs: Default alpha = 0.9| Sequencing Depth = 15.422129911096862| Trimming = 92.86918155320676| Alignment = 19.211411756396192| Gene Exon Mapping = 43.813950731314556| Ribosomal RNA = 0.348| Sequence Contamination = 10.860615219281971| Gene Body Coverage = 0.9| Dis

