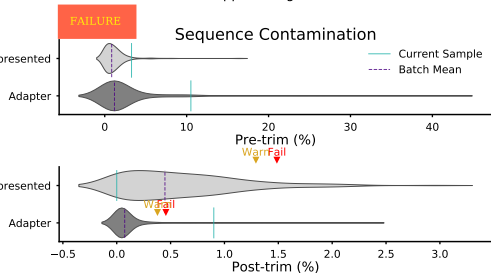
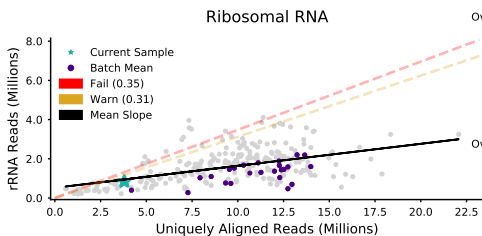
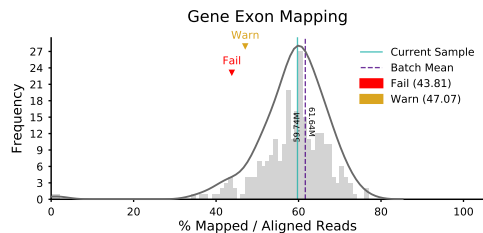
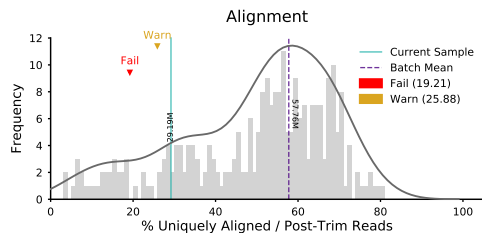
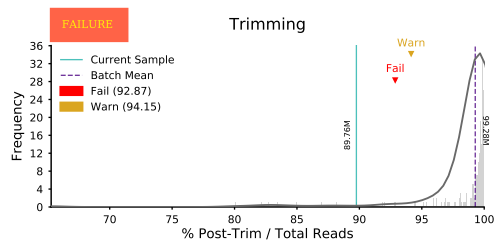
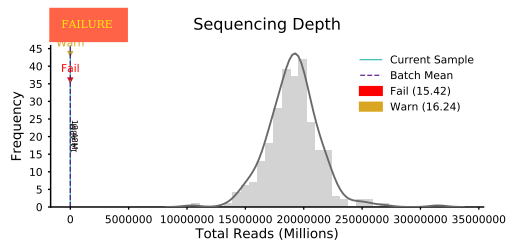
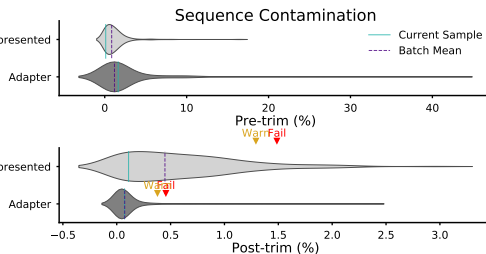
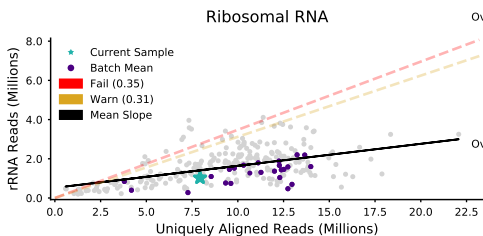
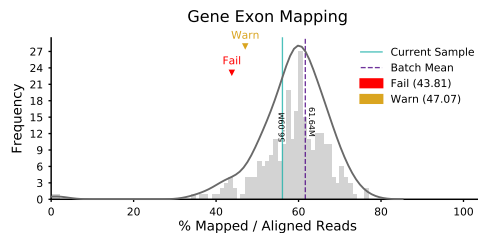
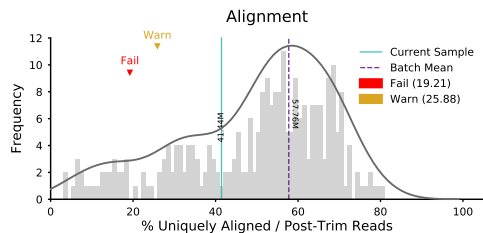
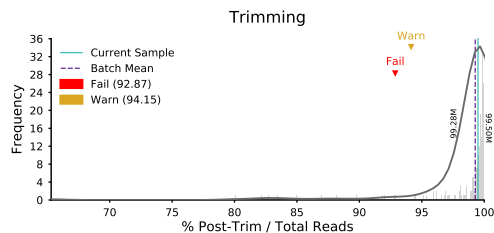
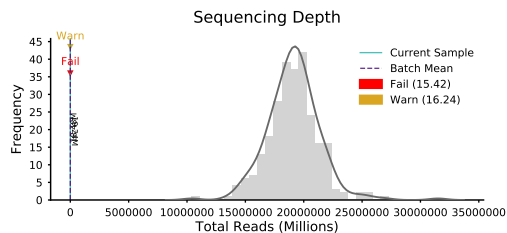


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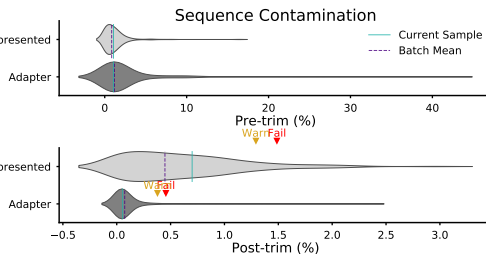
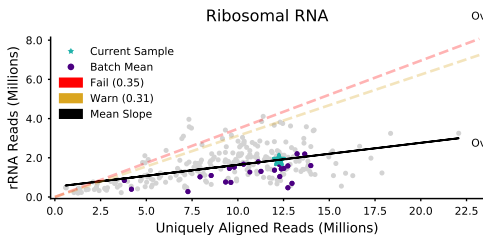
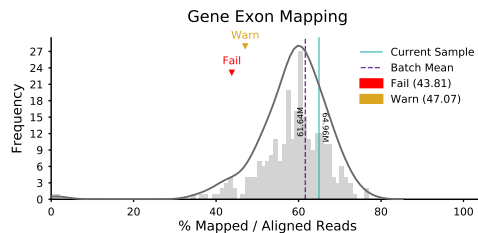
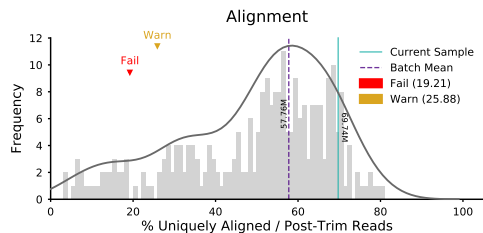
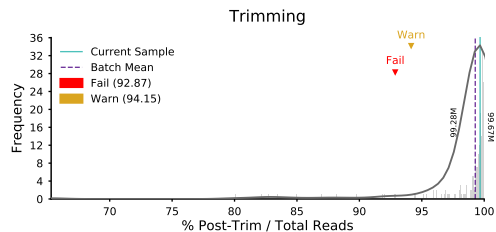
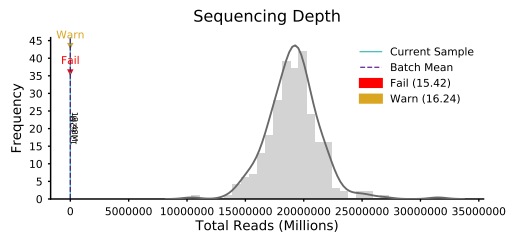


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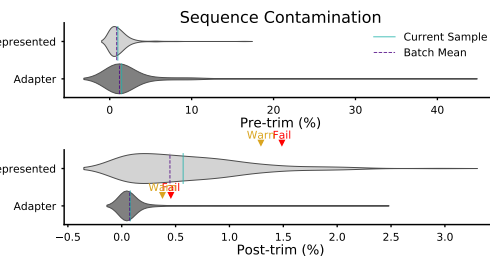
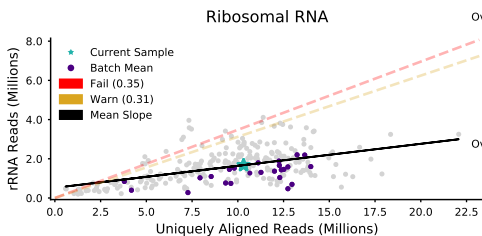
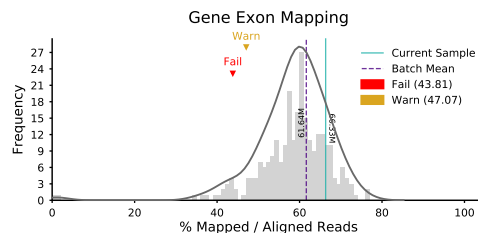
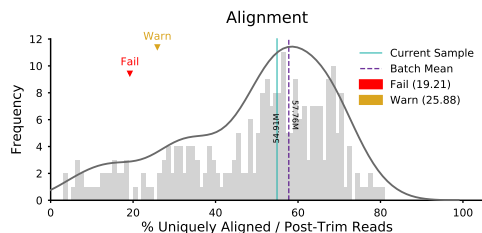
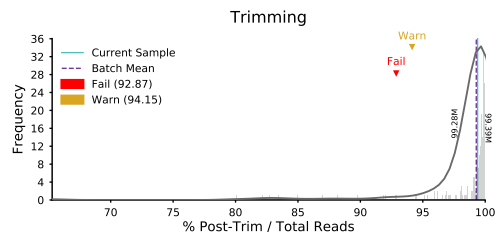
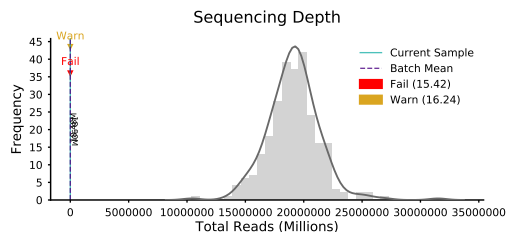
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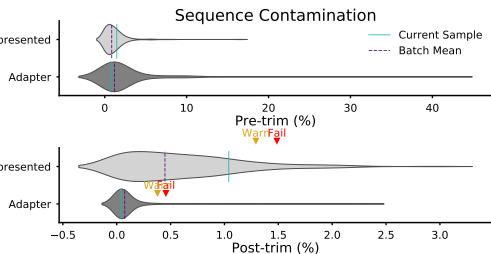
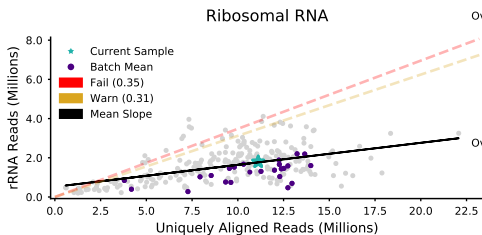
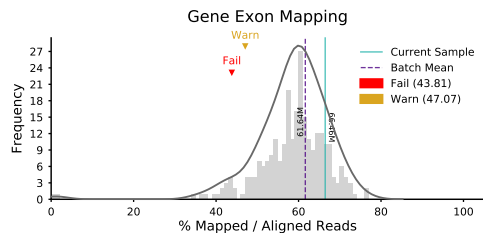
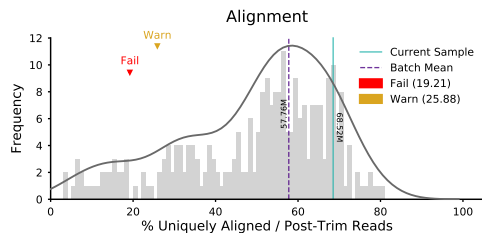
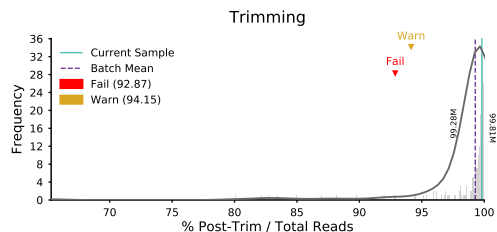
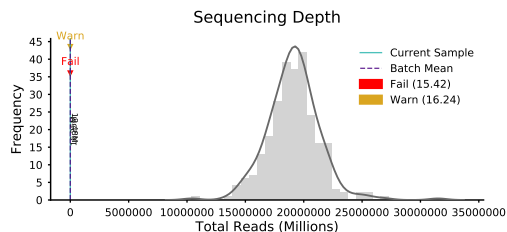


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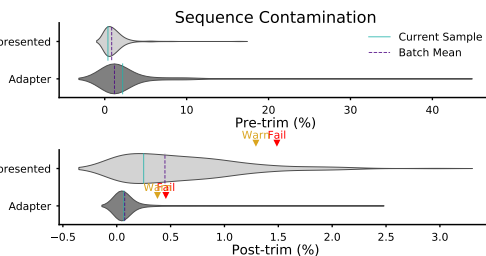
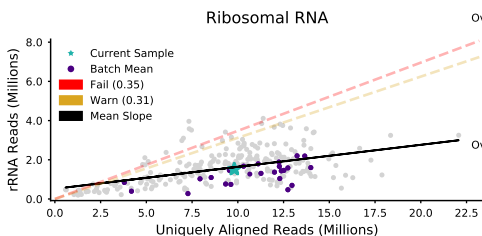
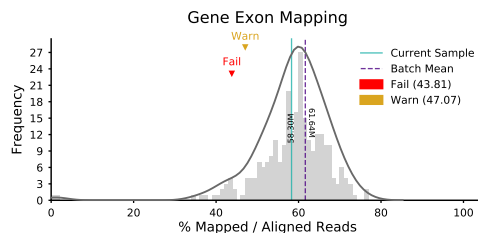
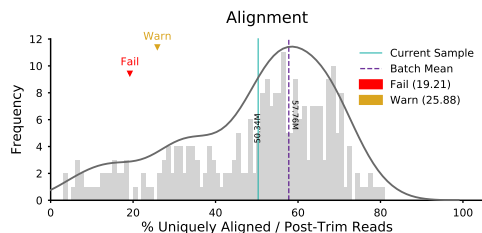
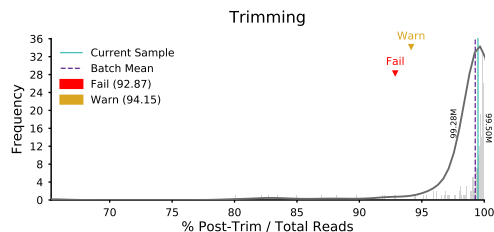
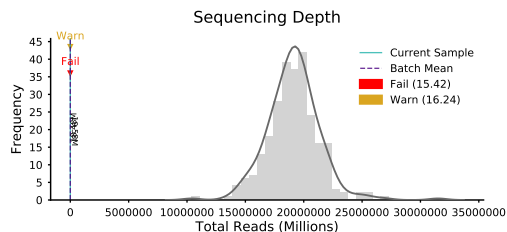


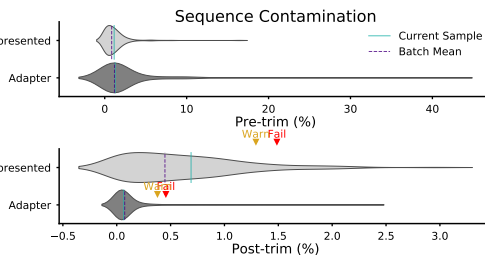
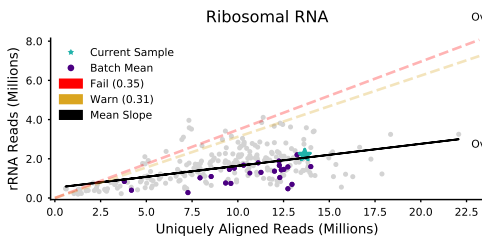
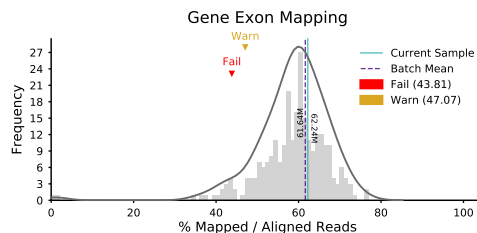
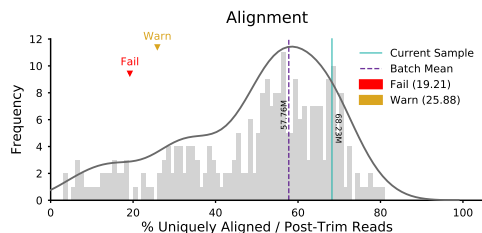
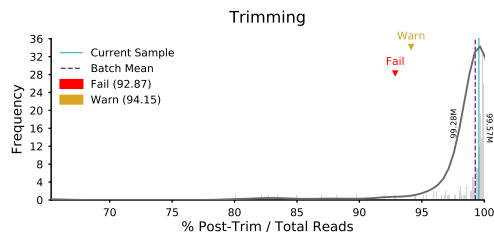
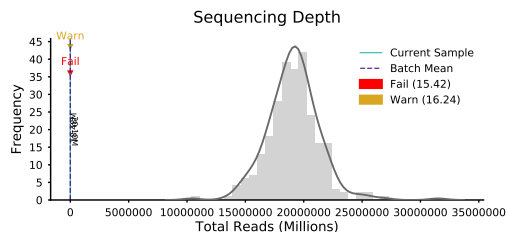
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# Sample : RNA\_B11\_06 Batch : batch\_11

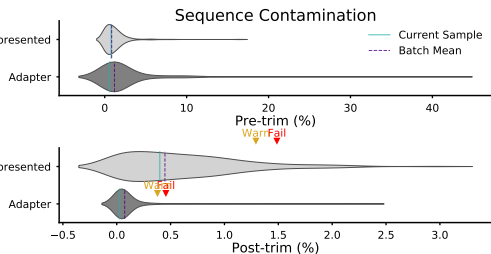
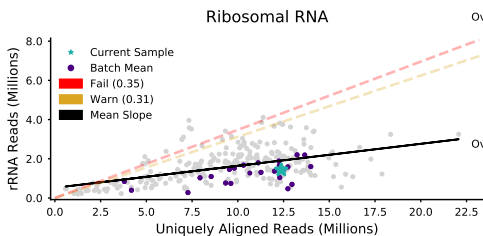
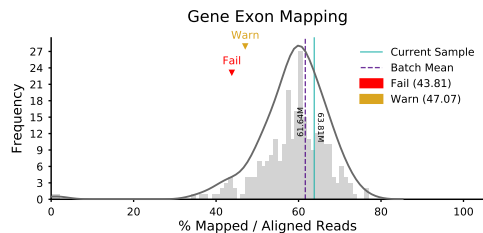
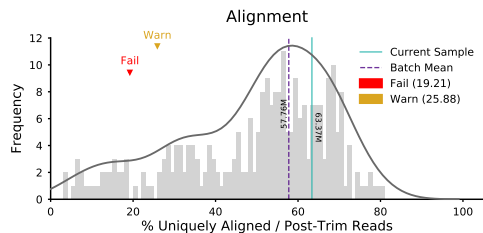
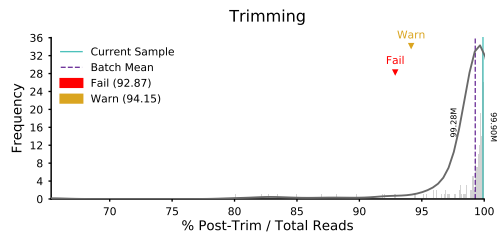
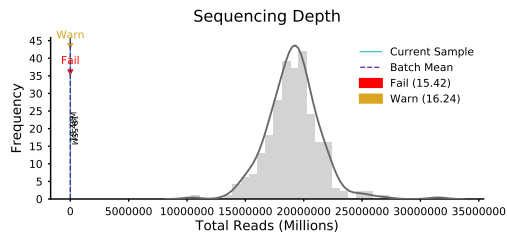
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# Sample : RNA\_B11\_08 Batch : batch\_11

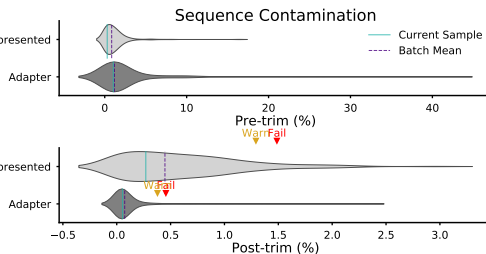
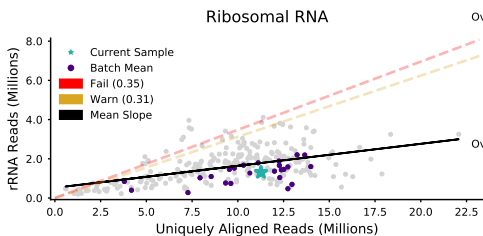
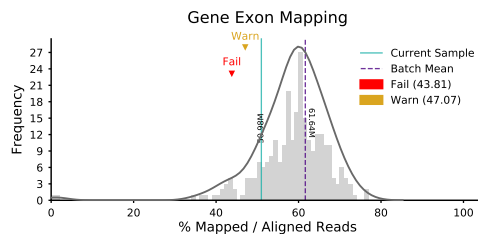
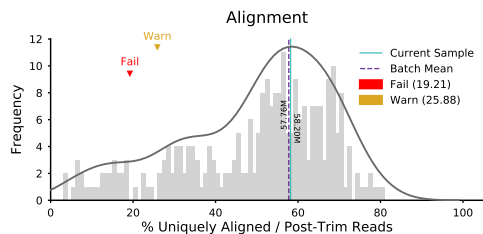
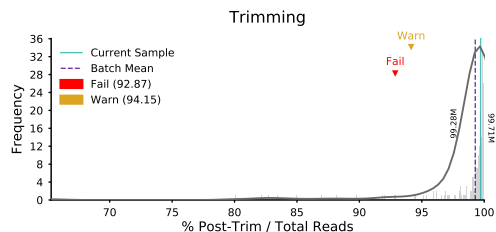
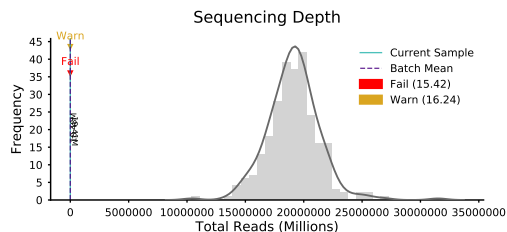
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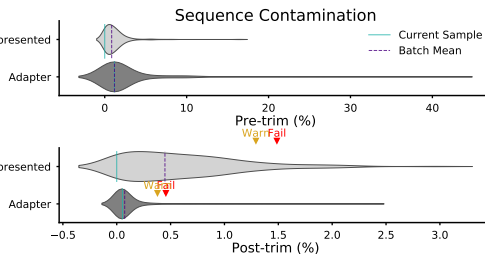
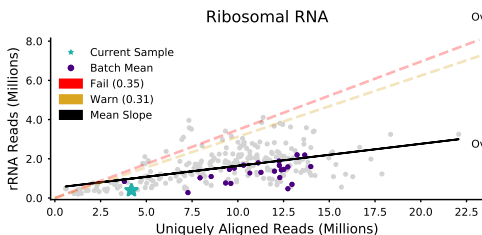
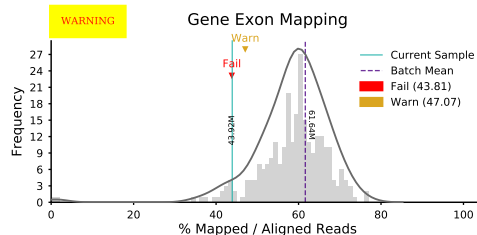
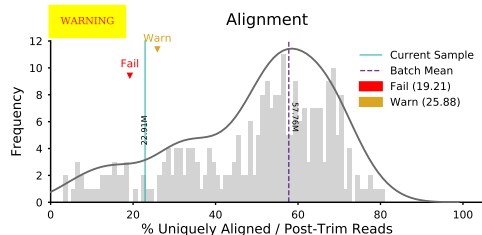
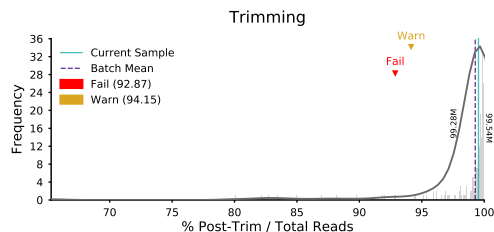
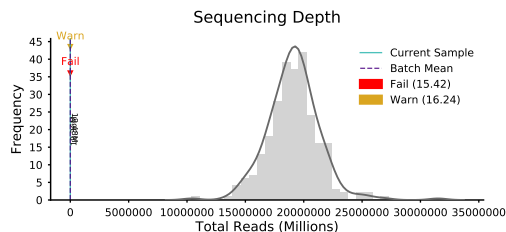
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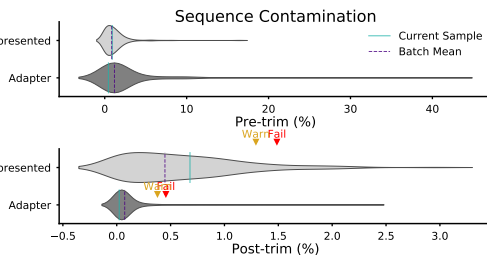
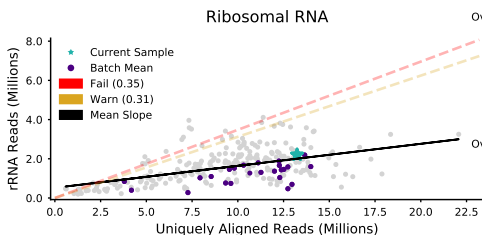
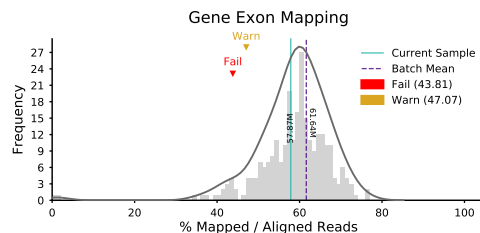
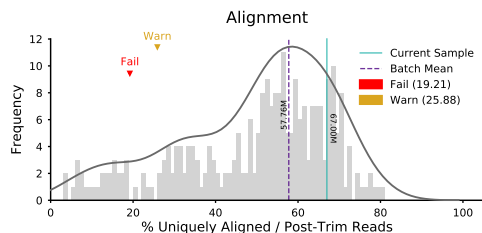
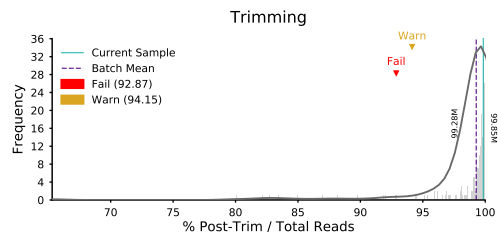
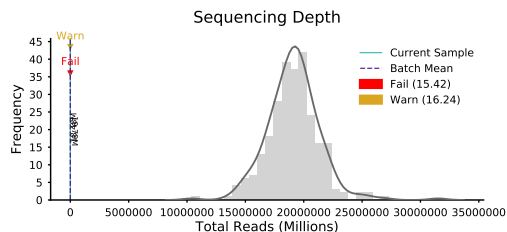


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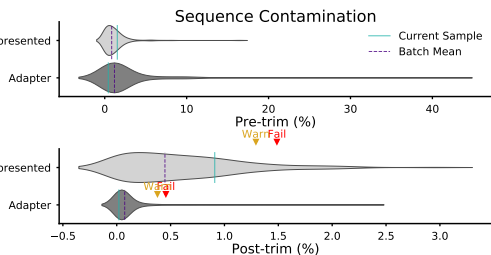
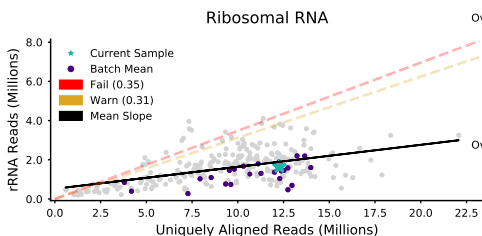
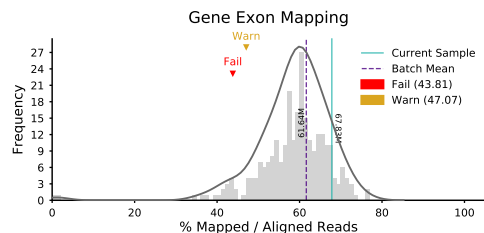
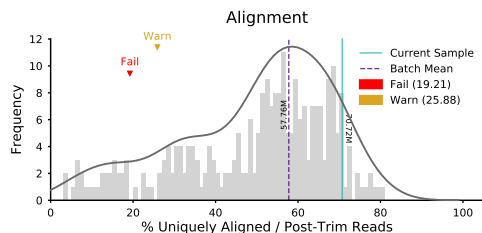
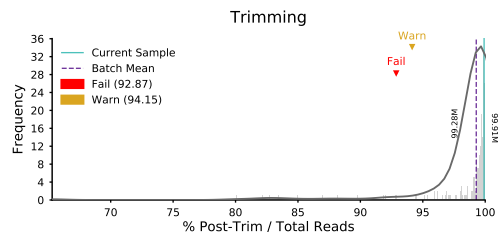
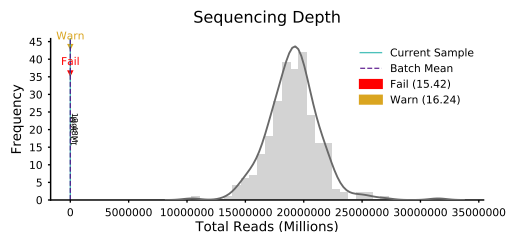
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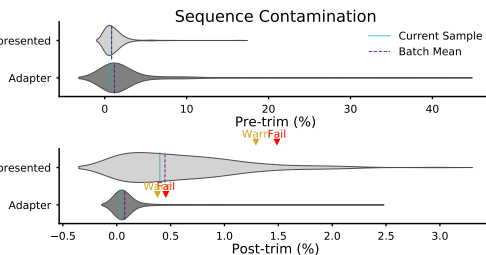
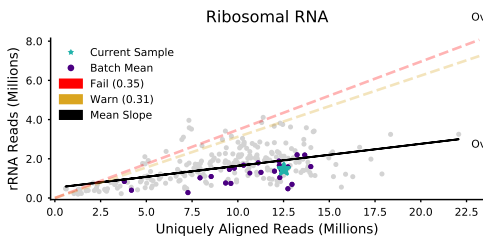
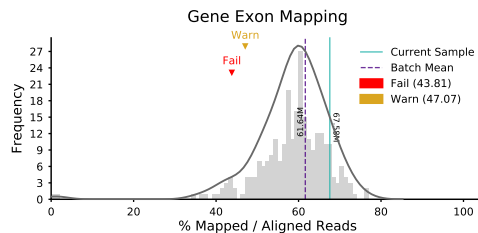
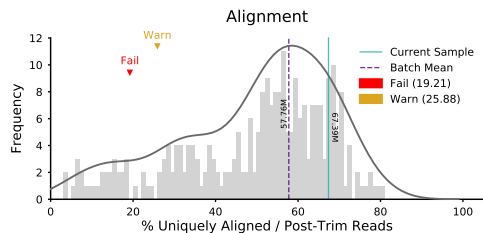
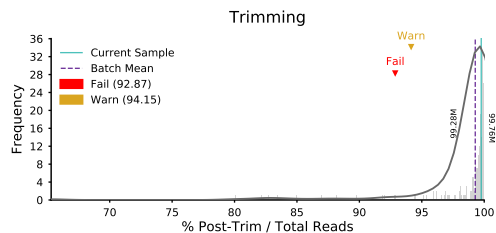
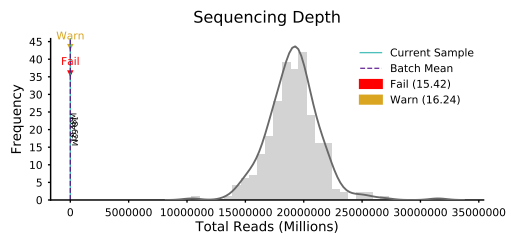
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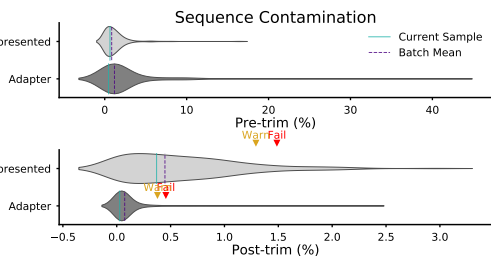
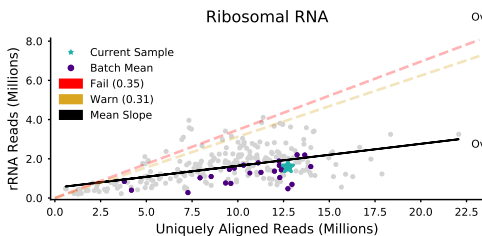
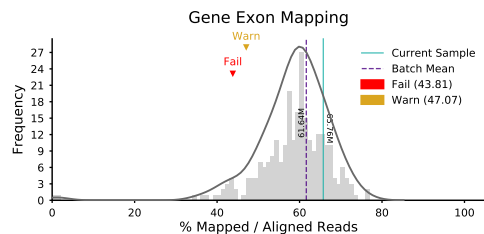
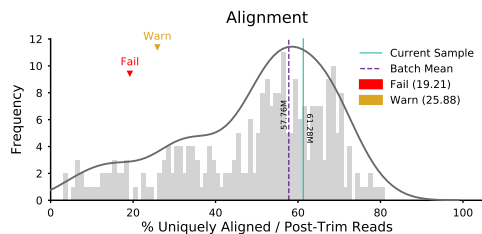
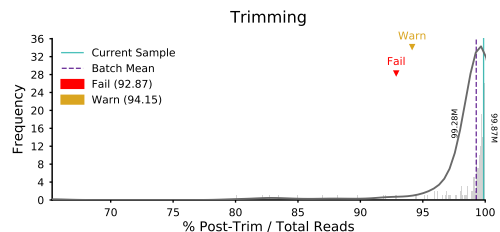
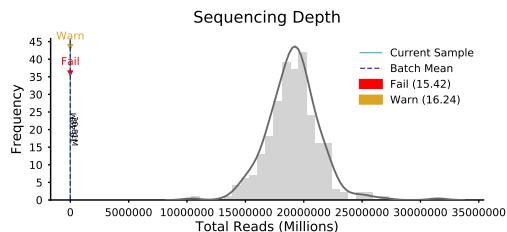
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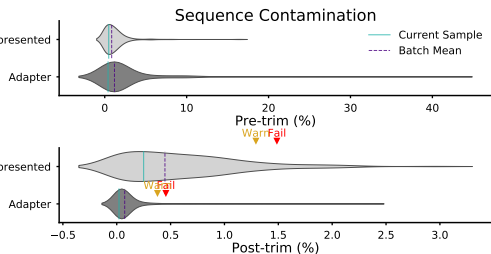
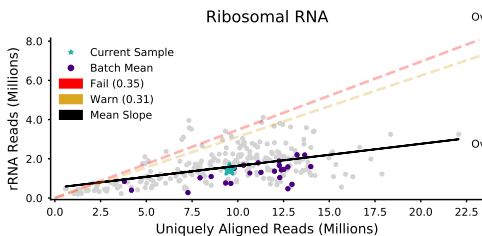
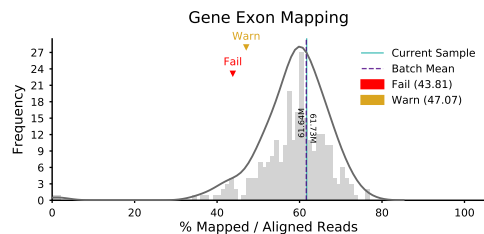
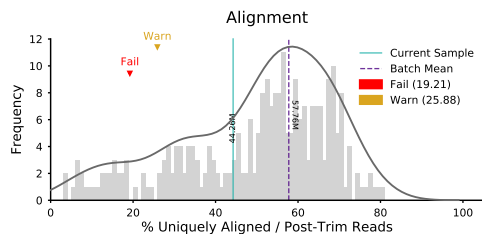
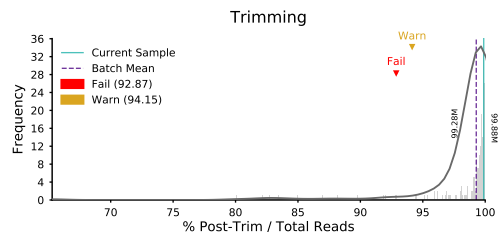
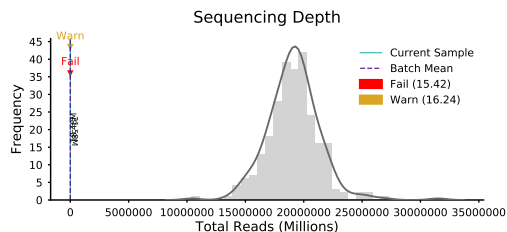
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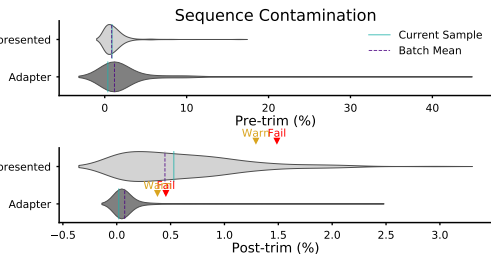
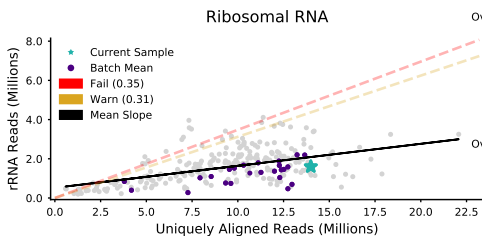
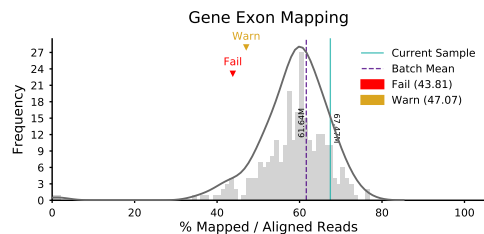
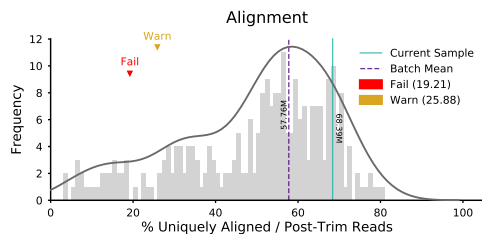
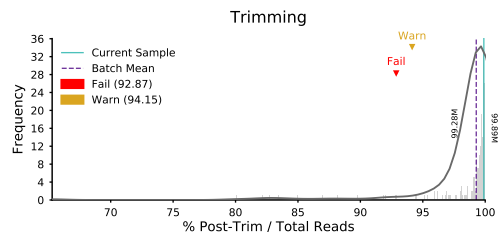
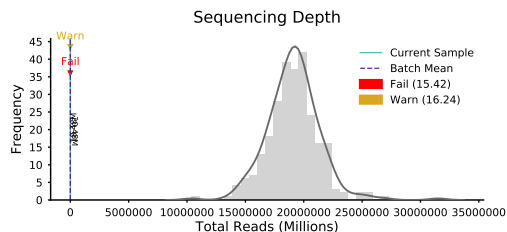
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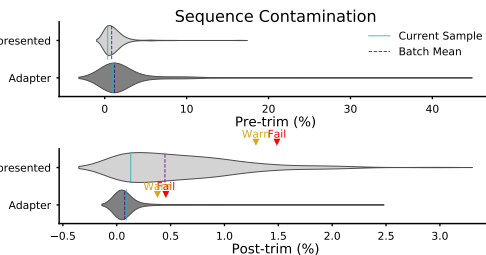
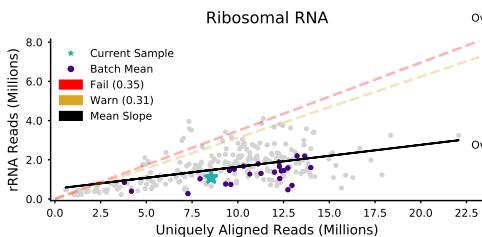
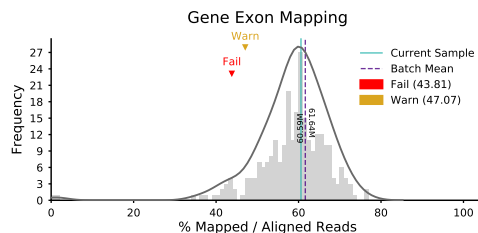
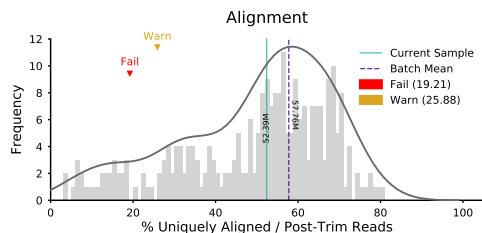
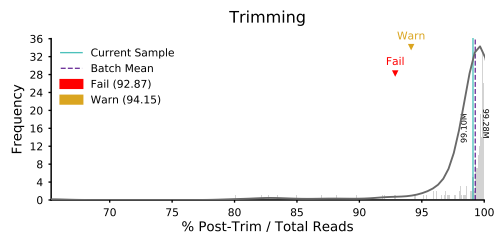
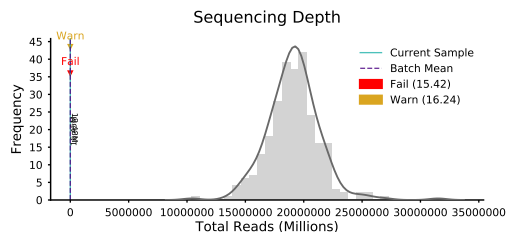


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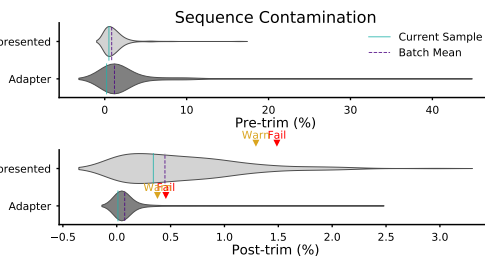
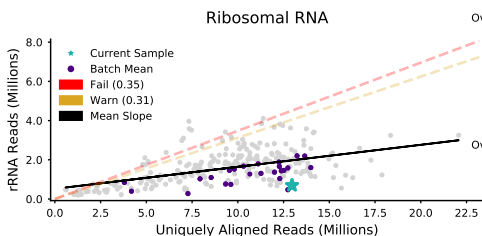
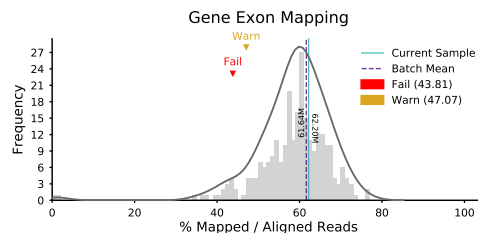
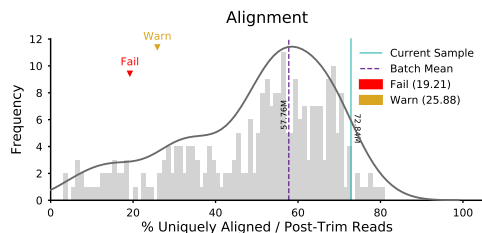
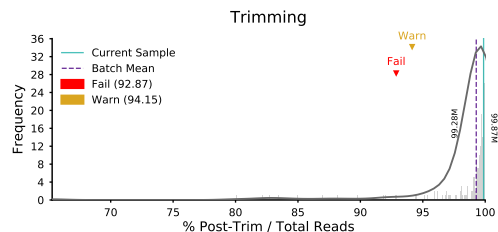
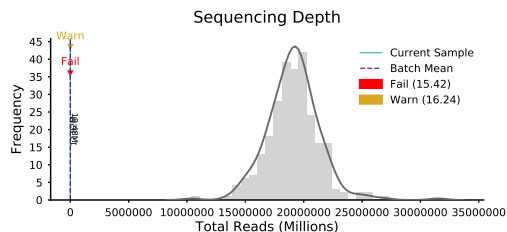


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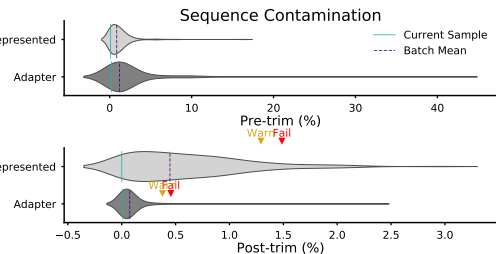
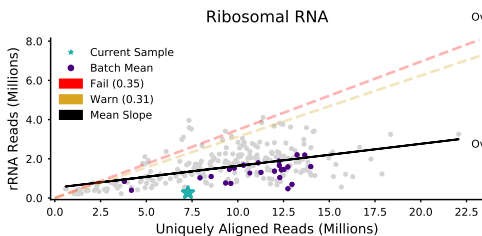
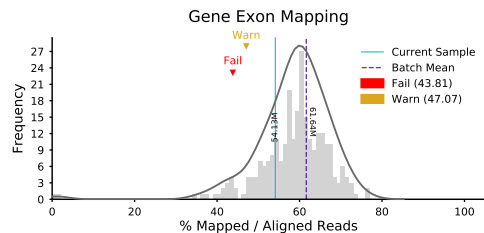
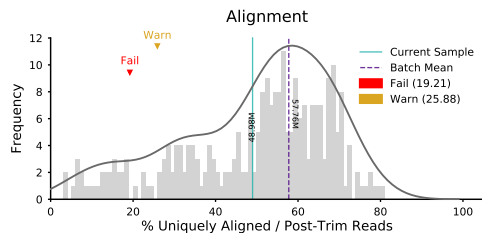
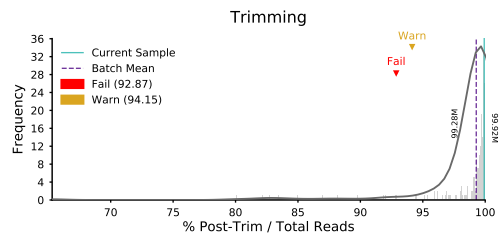
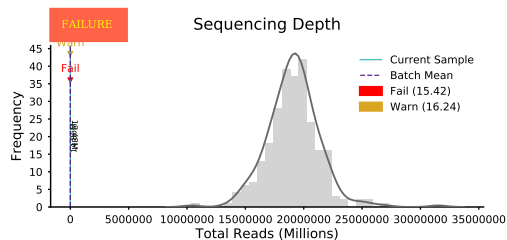


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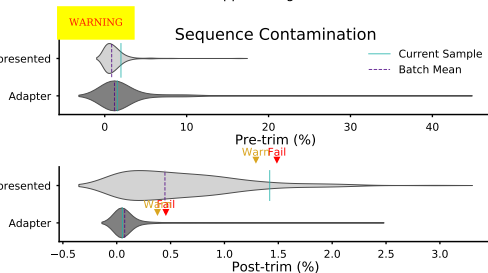
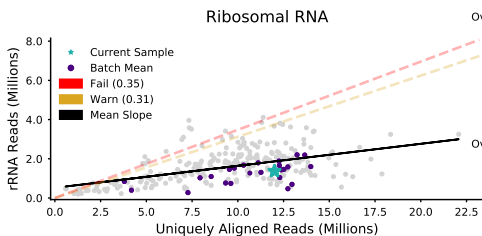
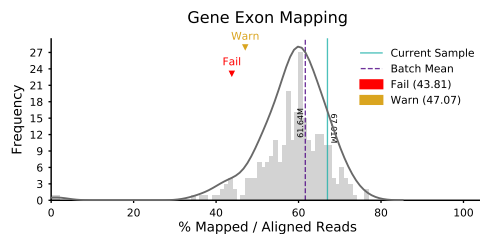
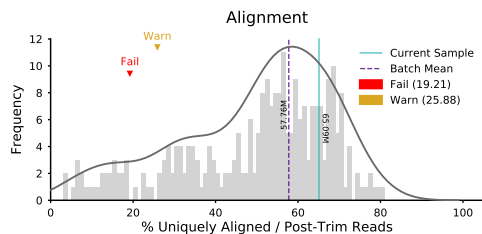
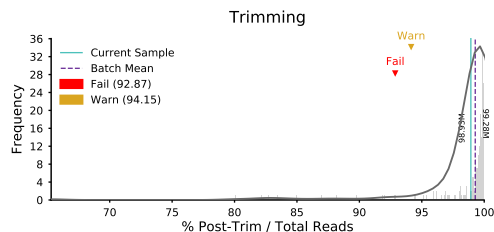
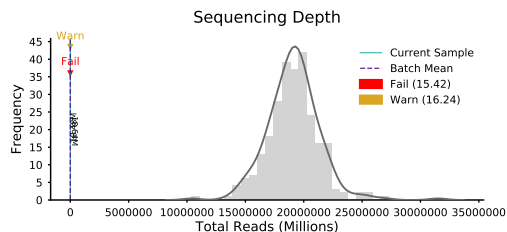
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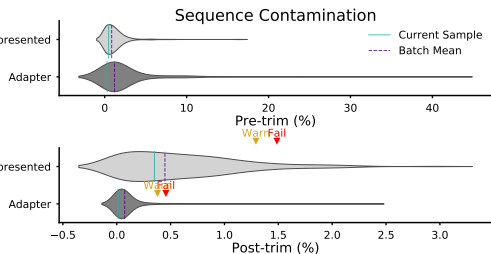
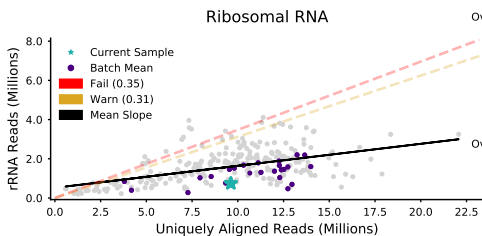
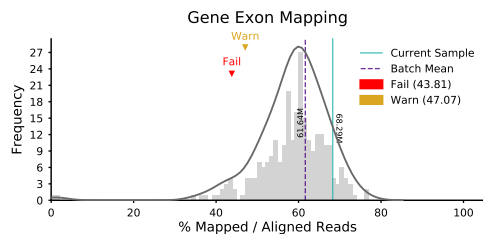
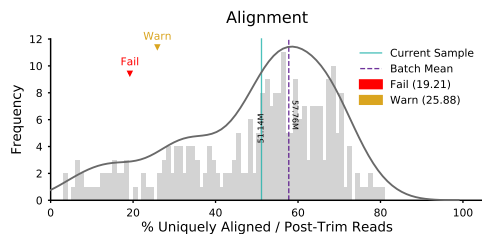
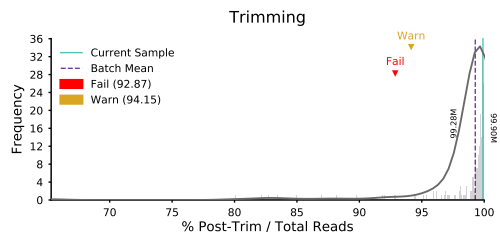
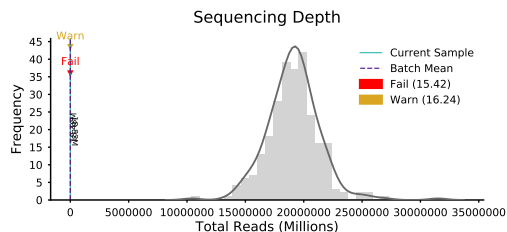
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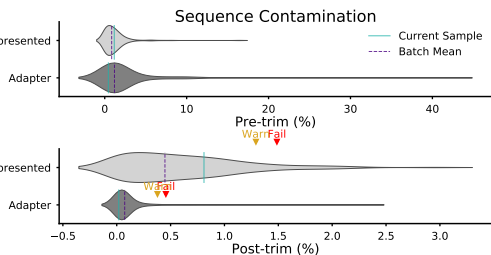
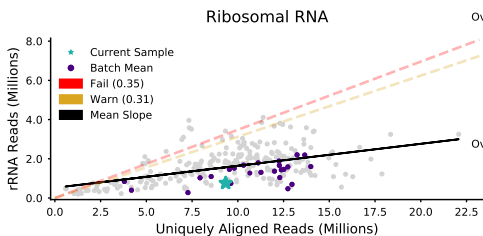
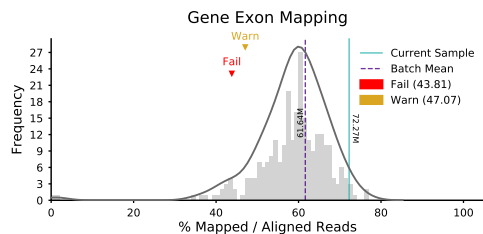
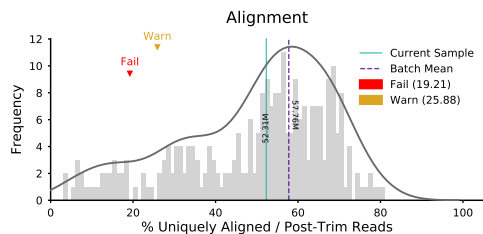
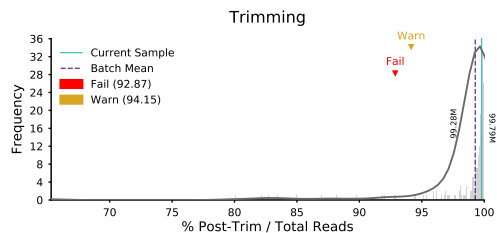
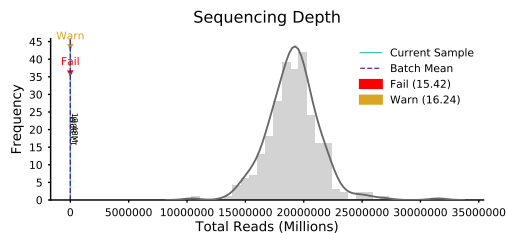
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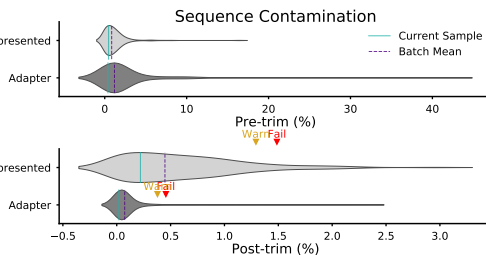
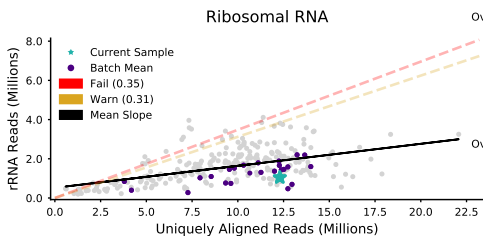
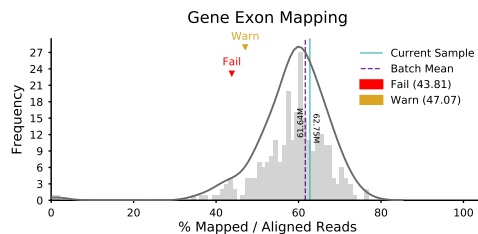
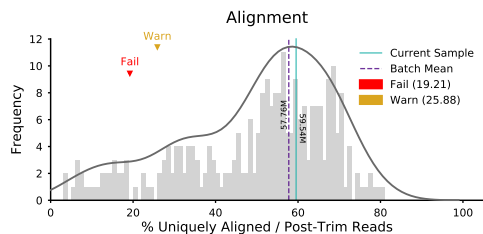
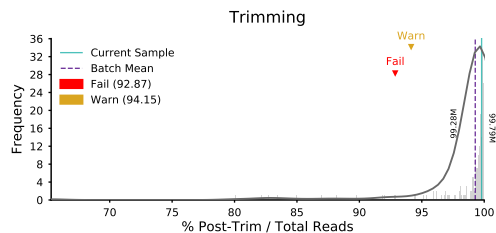
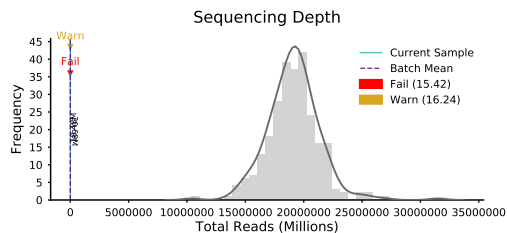
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Warn cutoffs: Default alpha =0.8| Sequencing Depth = 16.241476381729186| Trimming = 94.1462290640725| Alignment = 25.883432206791934| Gene Exon Mapping = 47.068386642306564| Ribosomal RNA = 0.312| Sequence Contamination = 0.37816525422038655| Gene Body Coverage = 0.8|  
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Warn 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 = 0.8100000000000001  
 Fail cutoffs: Default alpha = 0.9 Sequencing Depth = 15.422129911096862 Trimming = 92.86918155320676 Alignment = 19.21411756396192 Gene Exon Mapping = 43.813950731314556 Ribosomal RNA = 0.348 Sequence Contamination = 10.860615219281971 Gene Body Coverage = 0.9100000000000001

