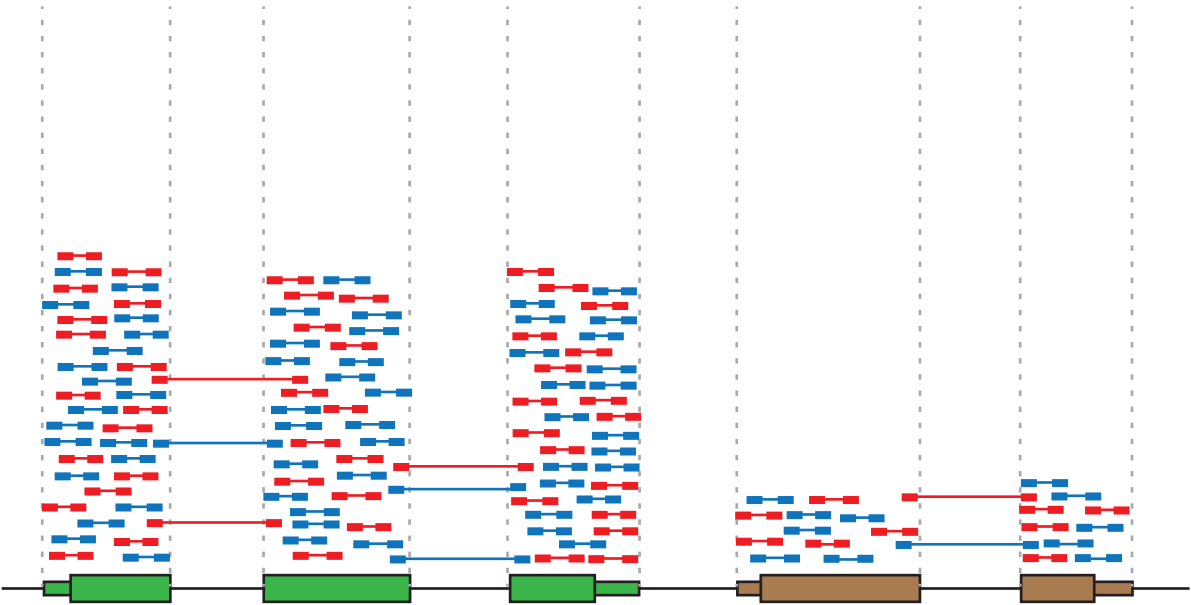
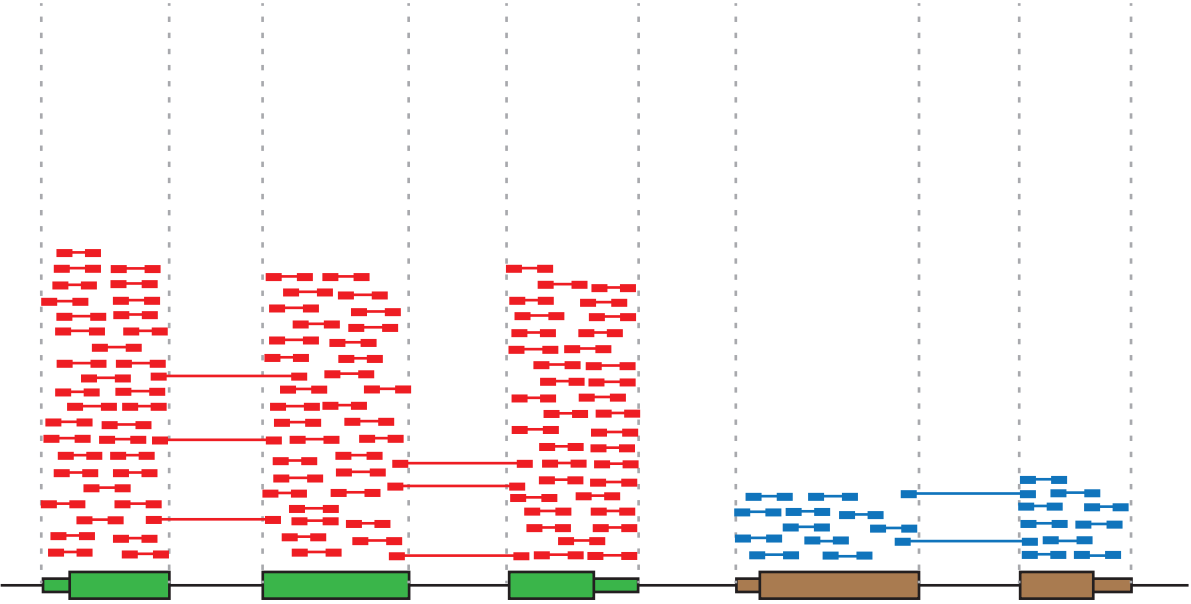


# Unstranded



# Stranded



# Analysis software settings for strand

Library Kit	5' to 3' IGV	TopHat	HTSeq	Picard	Stranded
TruSeq Strand Specific Total RNA	F2R1	<u>fr-firststrand</u>	reverse	SECOND_READ_TRANSCRIPTION_STRAND	Yes
NuGEN Encore	F1R2	<u>fr-secondstrand</u>	yes	FIRST_READ_TRANSCRIPTION_STRAND	Yes
NuGEN OvationV2	F2R1 or F1R2	fr-unstranded	no	NONE	No

# Viewing strand in IGV

