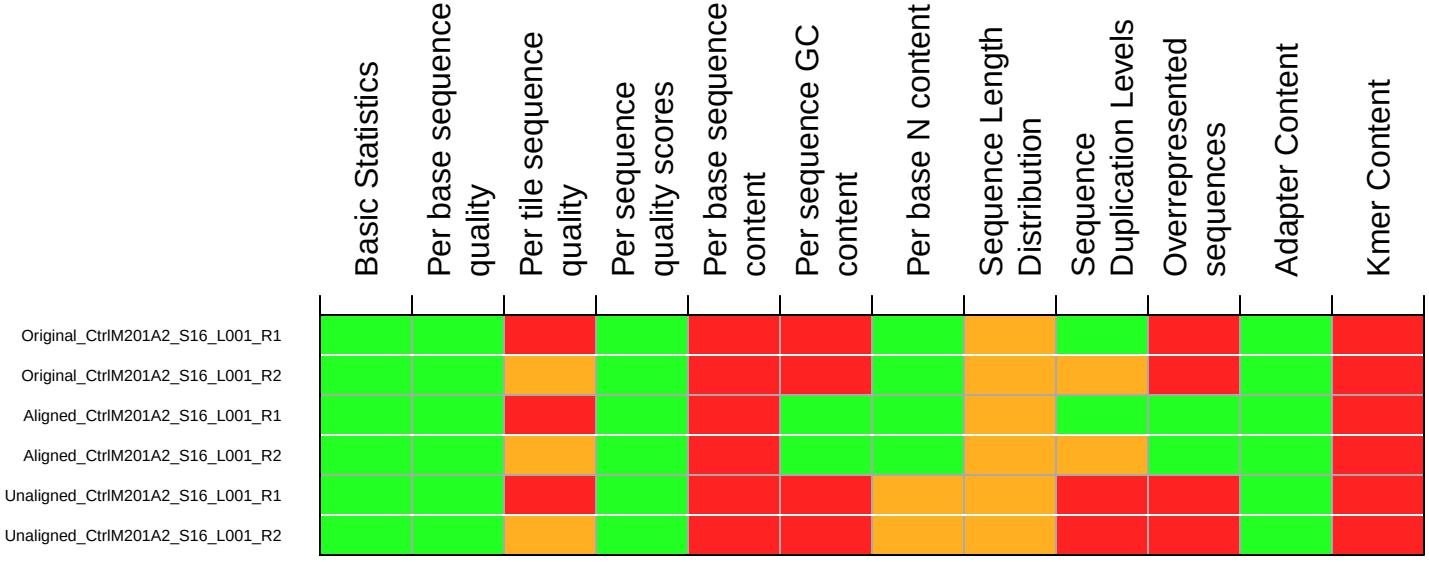
Basic Statistics
Per base sequence quality
Per tile sequence quality
Per sequence quality scores
Per base sequence content
Per sequence GC content
Per base N content
Sequence Length Distribution
Sequence Duplication Levels
Overrepresented sequences
Adapter Content
Kmer Content
Original_CtrlM201A2_S16_L001_R1
Original_CtrlM201A2_S16_L001_R2
Aligned_CtrlM201A2_S16_L001_R1
Aligned_CtrlM201A2_S16_L001_R2
Unaligned_CtrlM201A2_S16_L001_R1
Unaligned_CtrlM201A2_S16_L001_R2