

	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	Green	Green	Red	Green	Red	Red	Green	Orange	Green	Red	Green	Red
Original_CtrlM201A2_S16_L001_R2	Green	Green	Orange	Green	Red	Red	Green	Orange	Orange	Red	Green	Red
Aligned_CtrlM201A2_S16_L001_R1	Green	Green	Red	Green	Red	Green	Green	Orange	Green	Green	Green	Red
Aligned_CtrlM201A2_S16_L001_R2	Green	Green	Orange	Green	Red	Green	Green	Orange	Orange	Green	Green	Red
Unaligned_CtrlM201A2_S16_L001_R1	Green	Green	Red	Green	Red	Red	Orange	Orange	Red	Red	Green	Red
Unaligned_CtrlM201A2_S16_L001_R2	Green	Green	Orange	Green	Red	Red	Orange	Orange	Red	Red	Green	Red