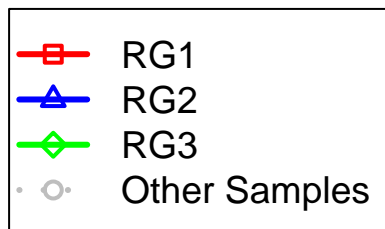
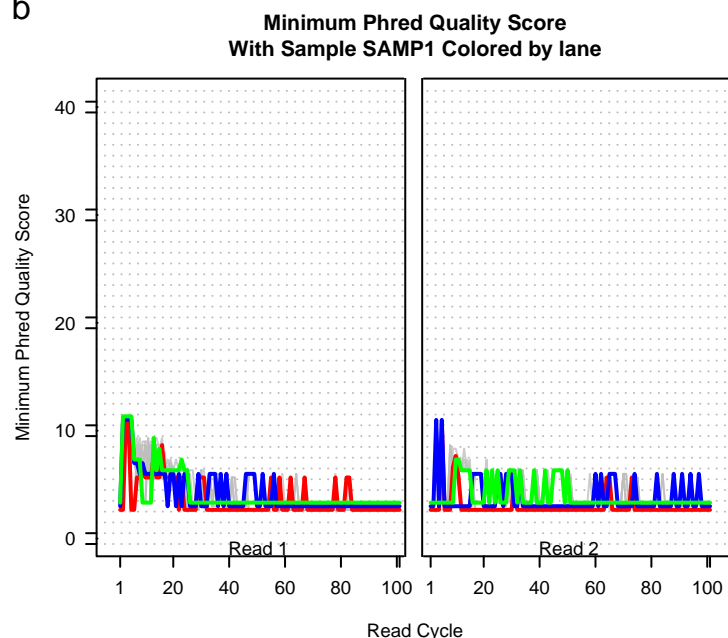


a

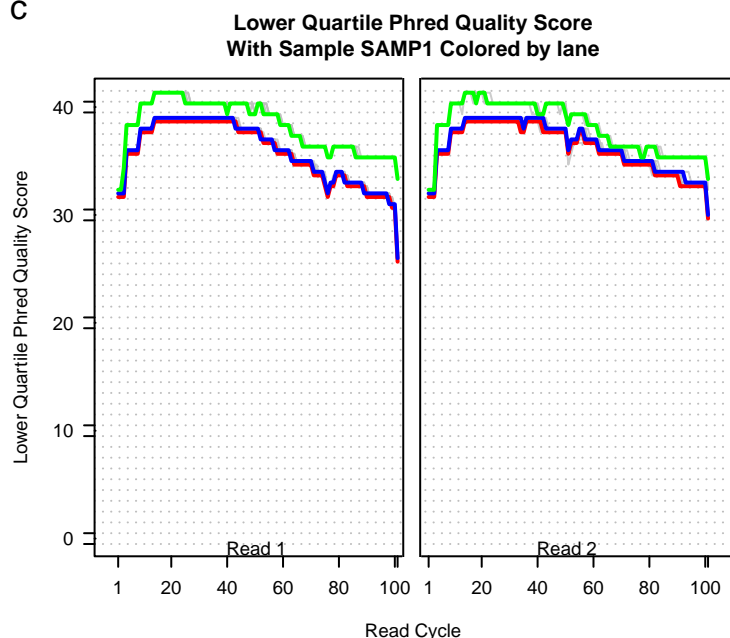
# Sample Highlight: SAMP1 Colored by Lane



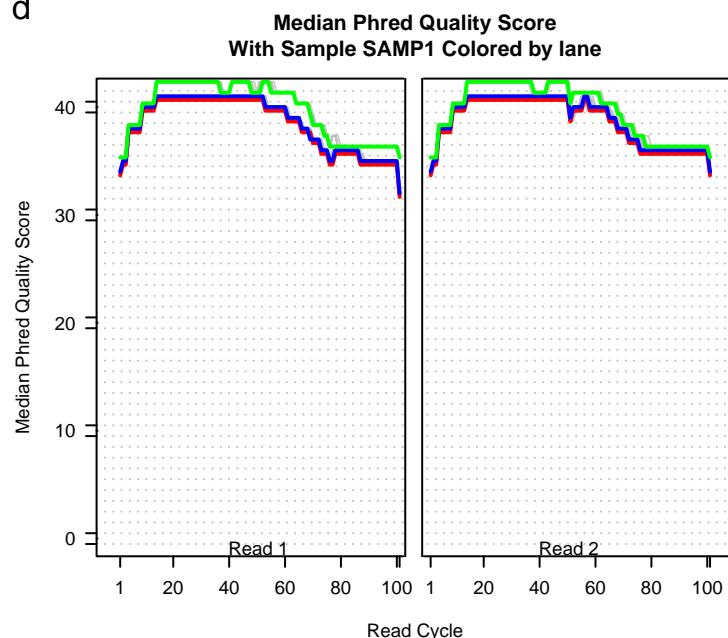
b



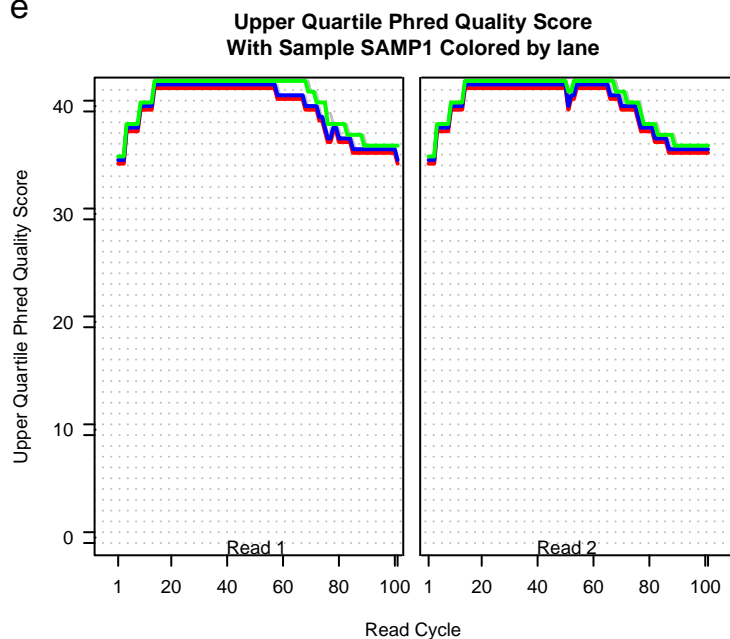
c



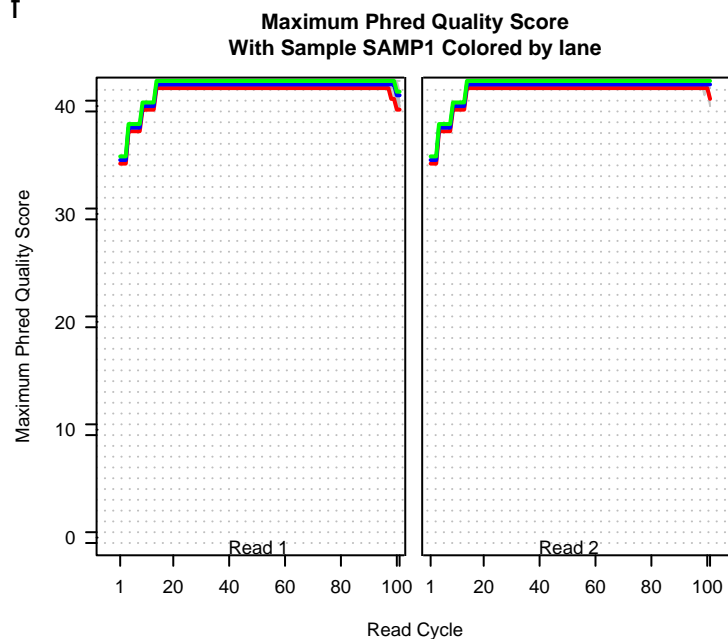
d

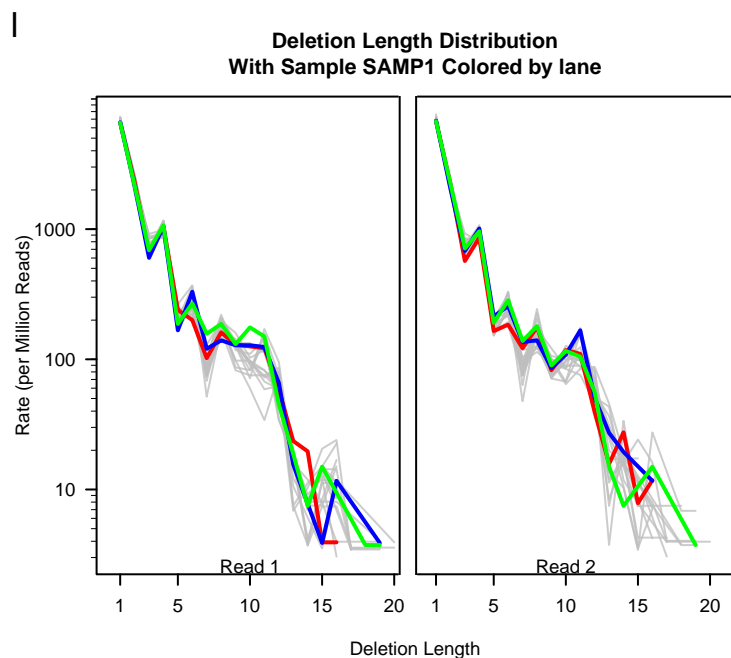
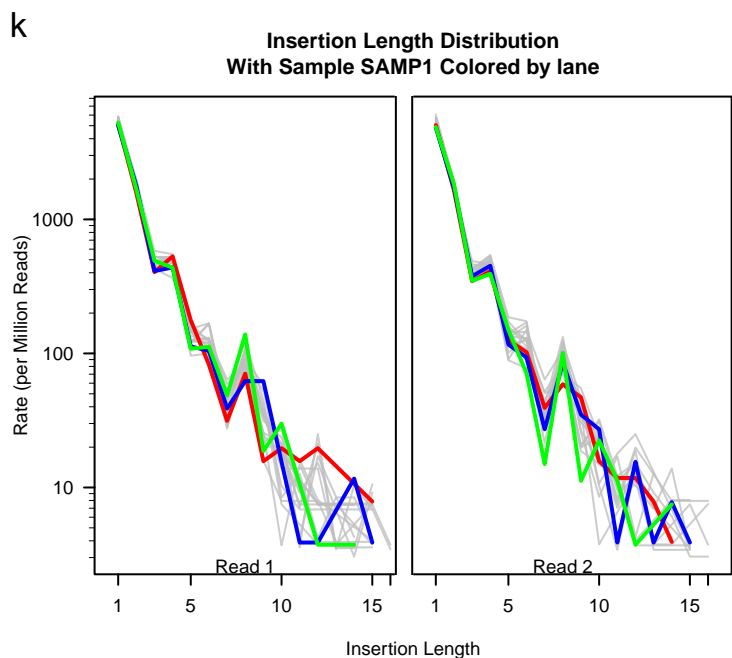
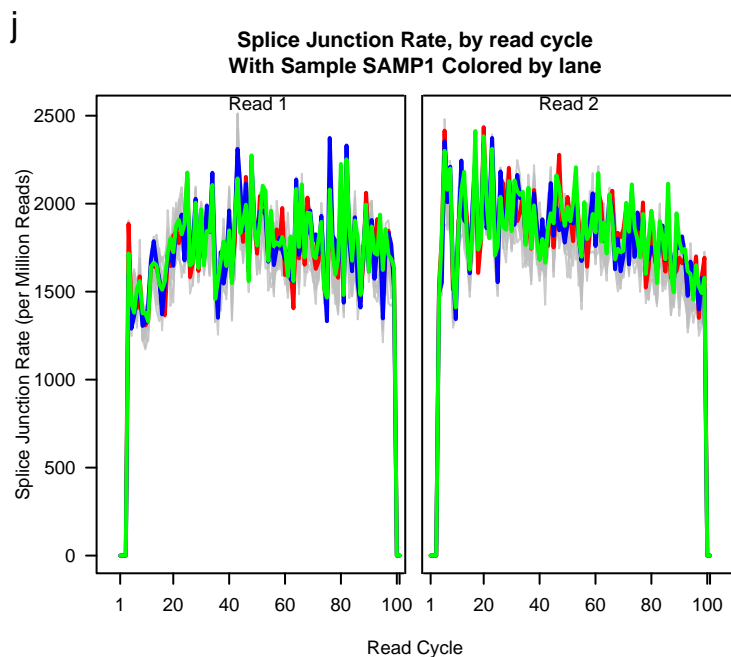
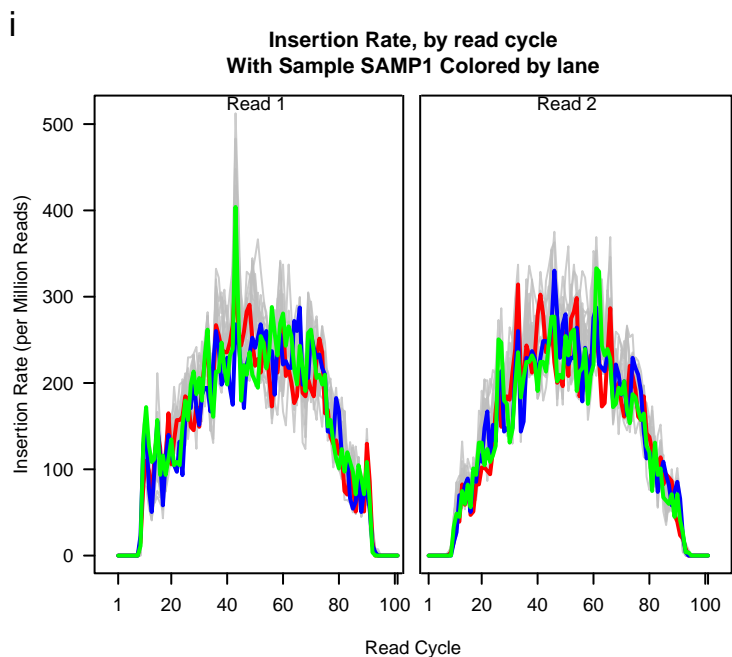
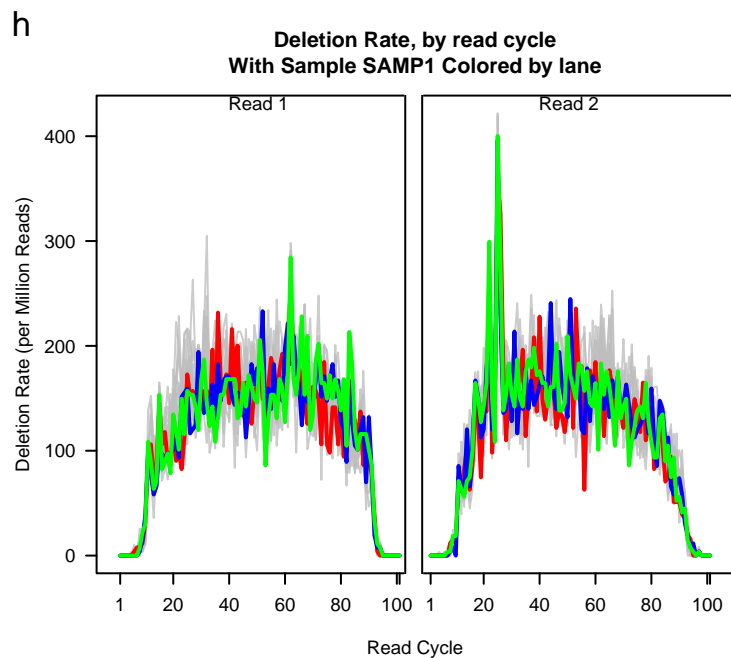
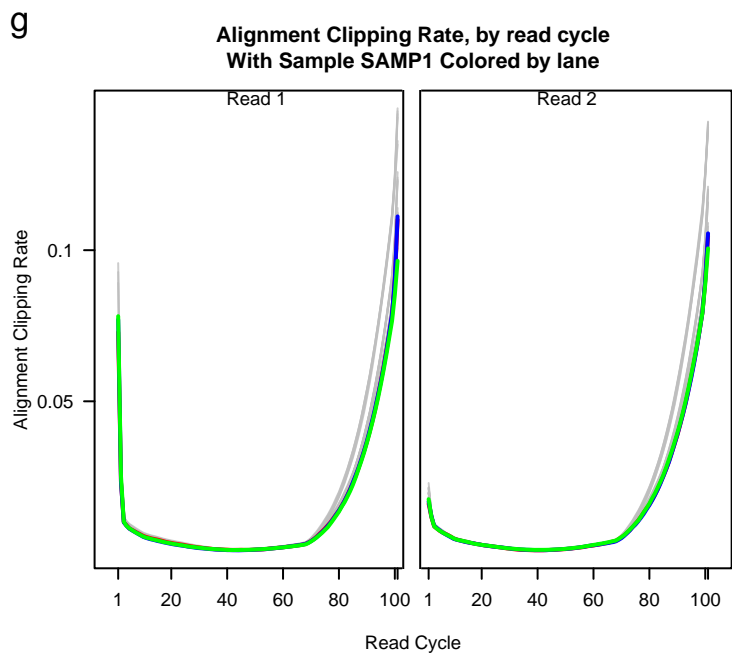


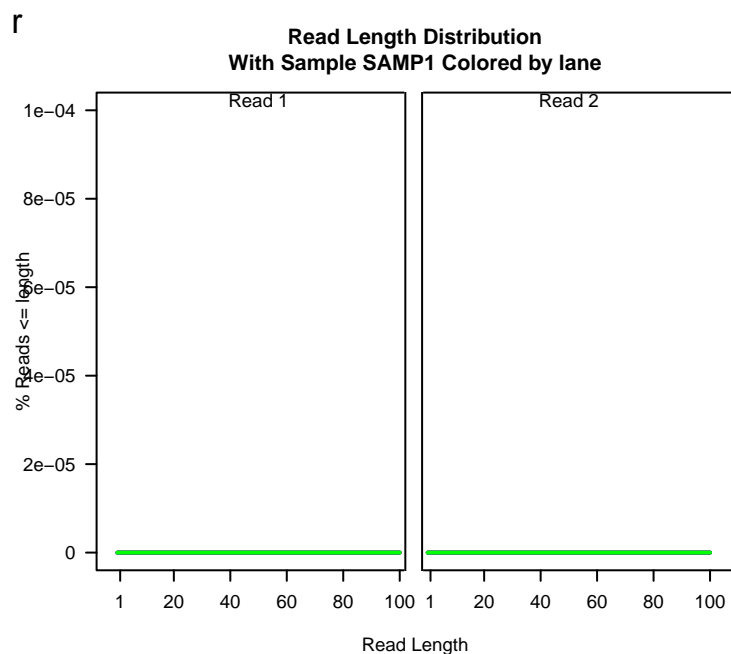
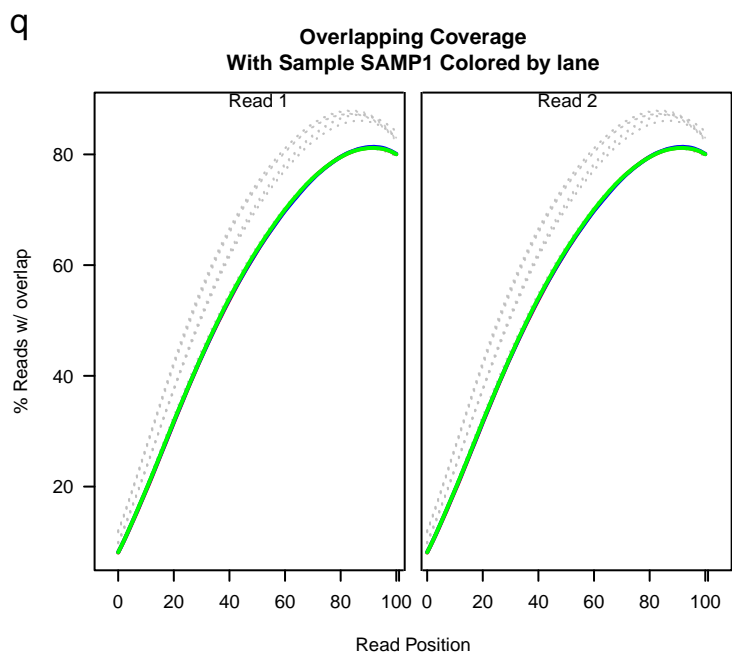
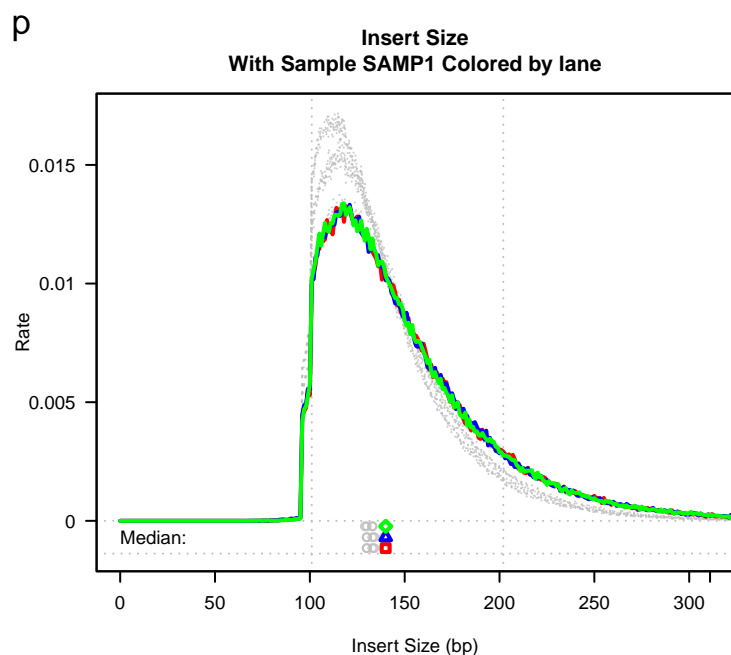
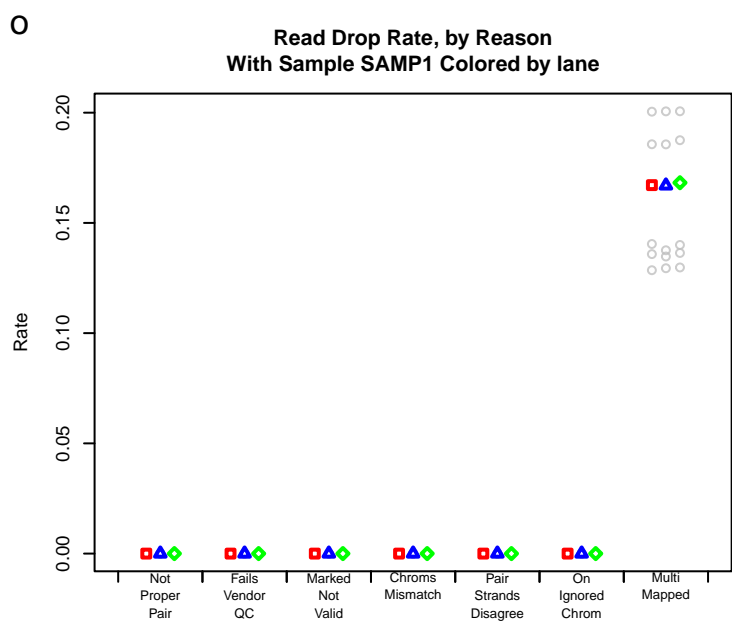
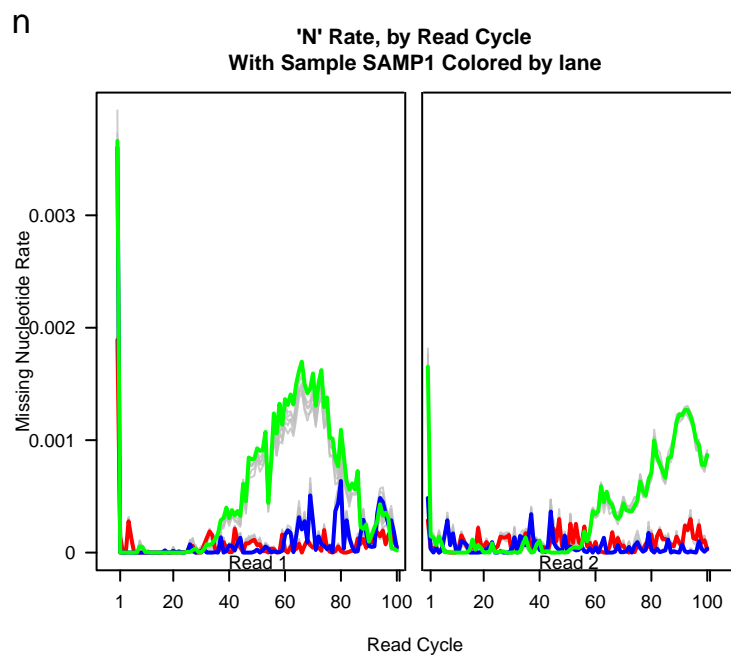
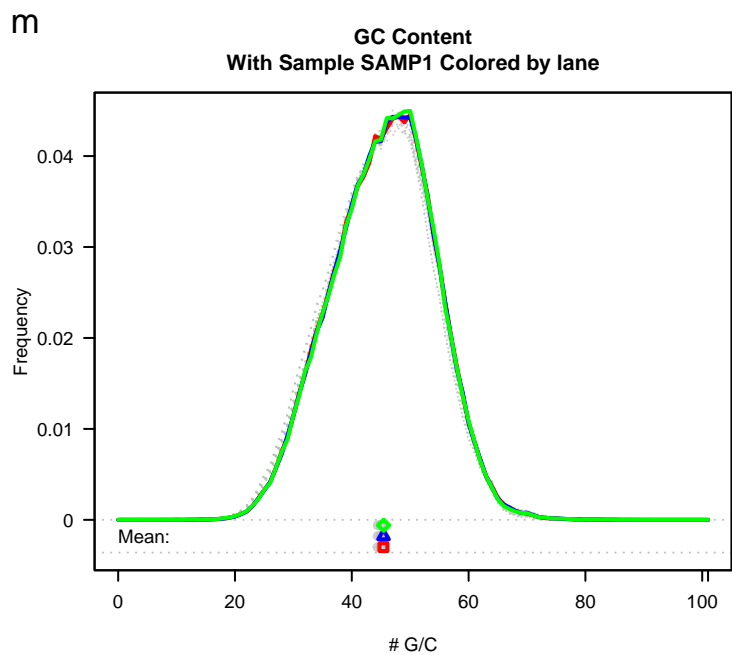
e

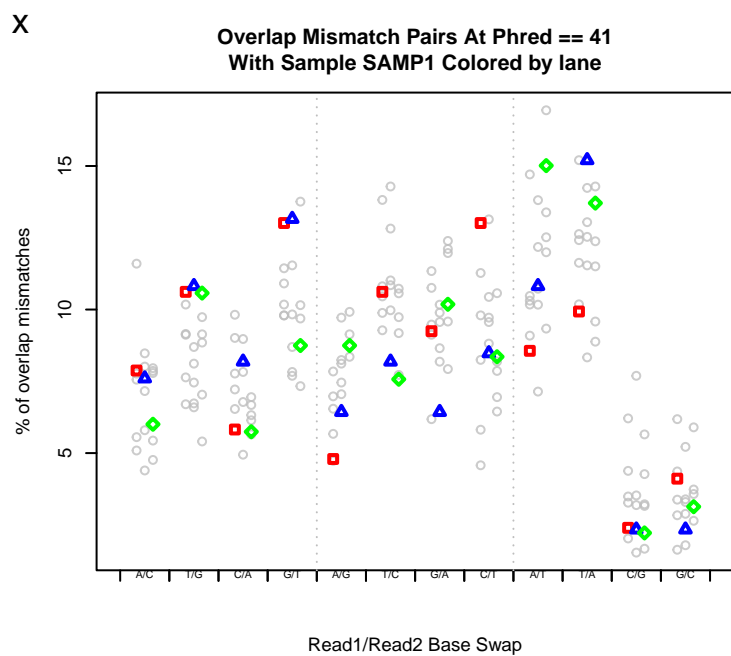
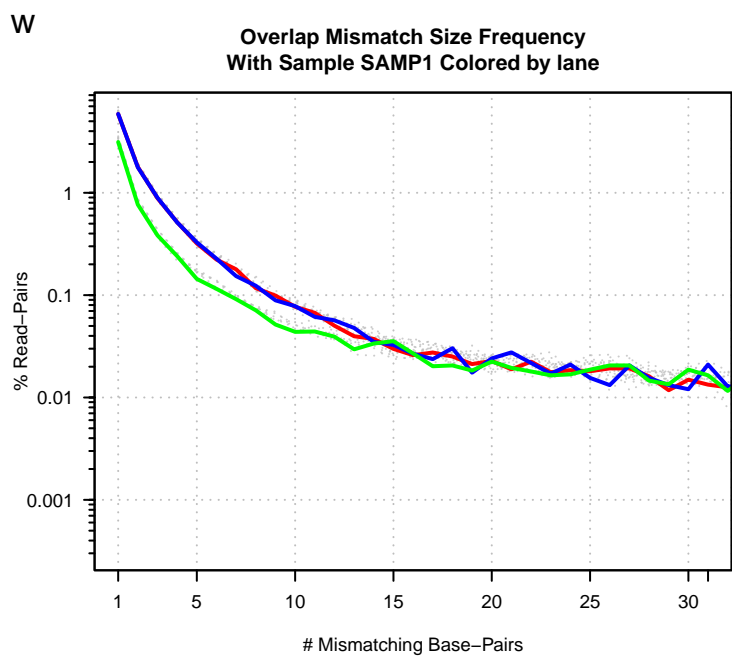
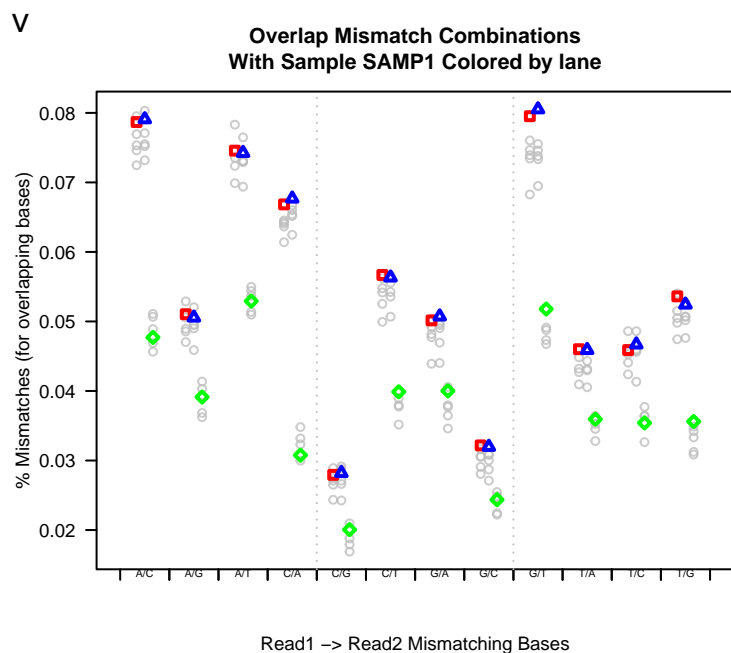
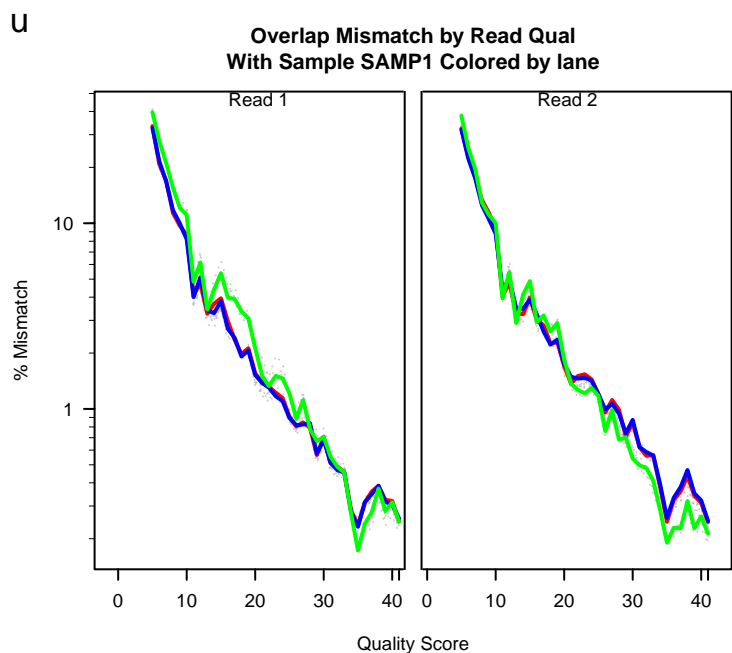
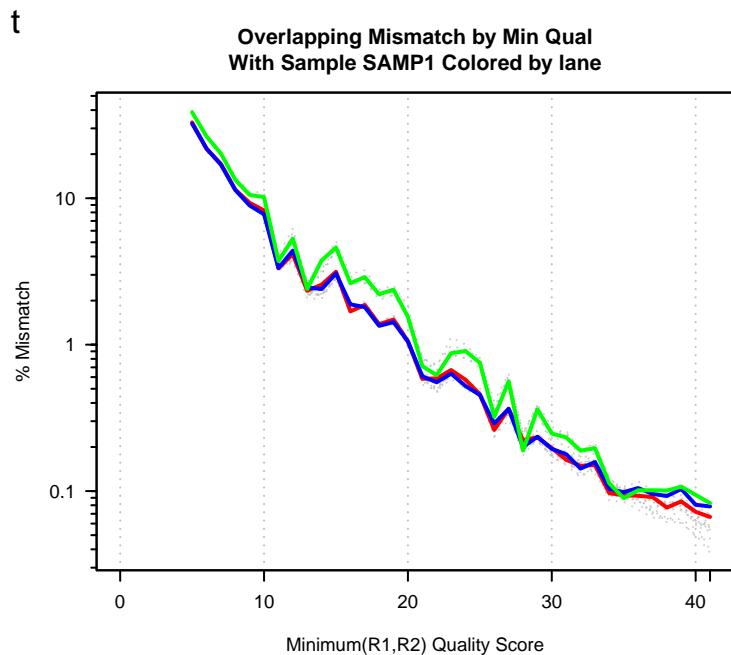
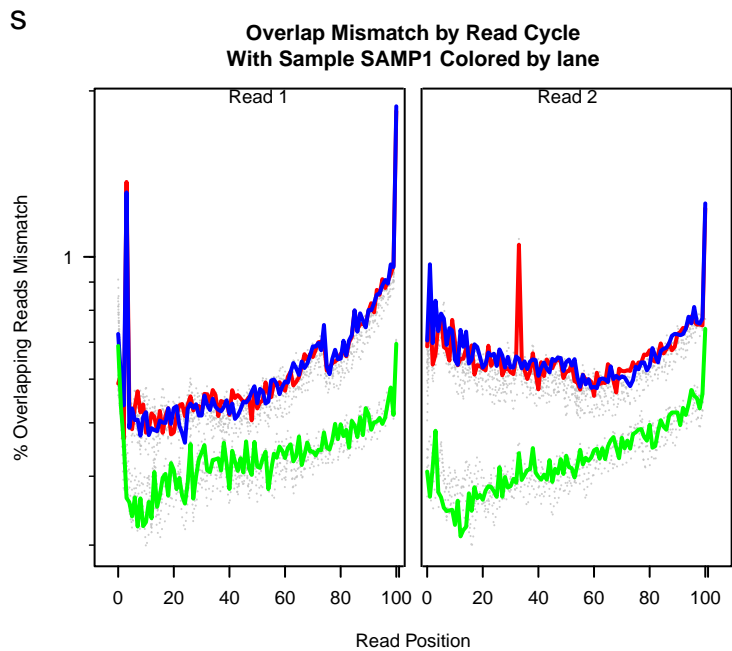


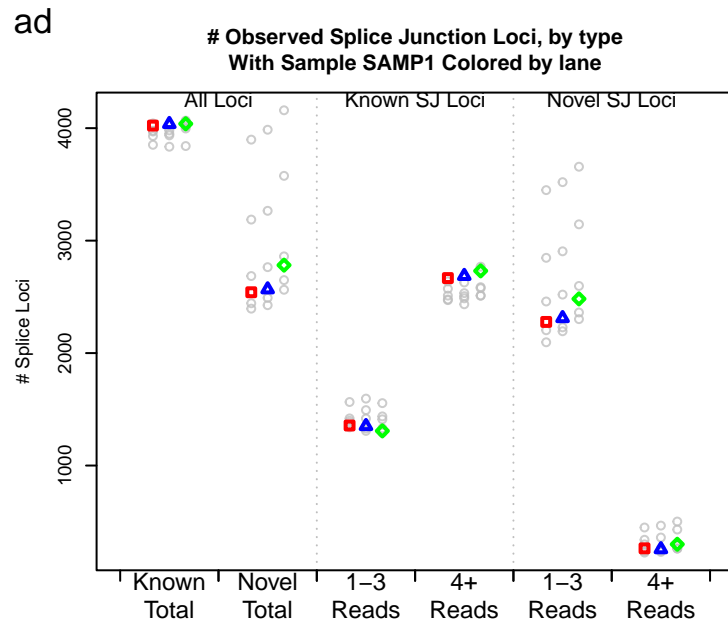
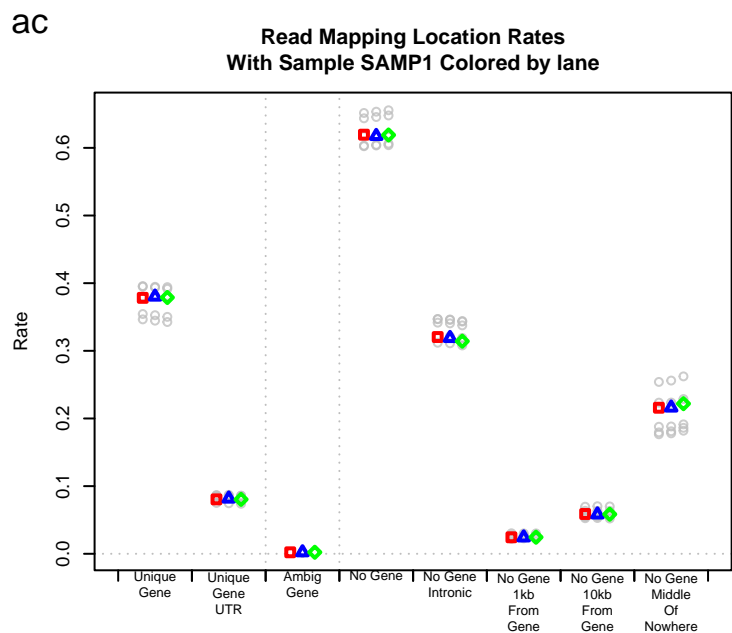
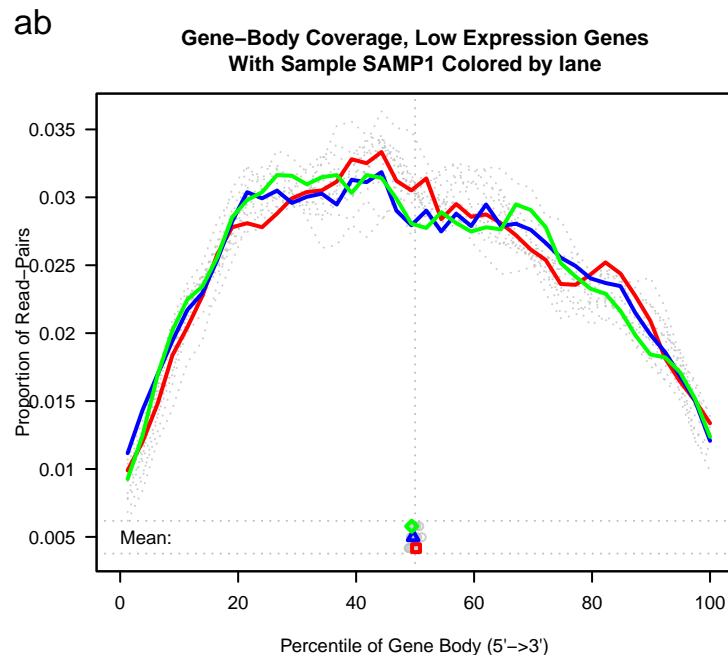
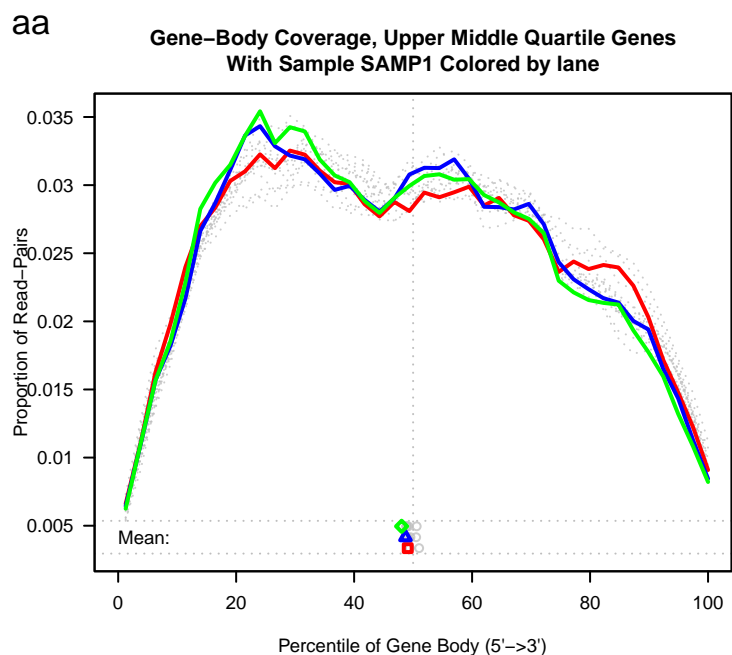
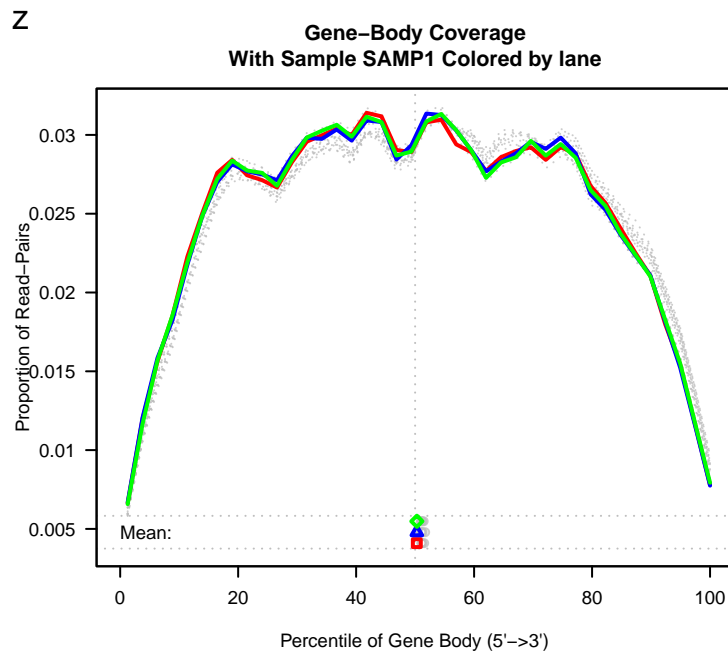
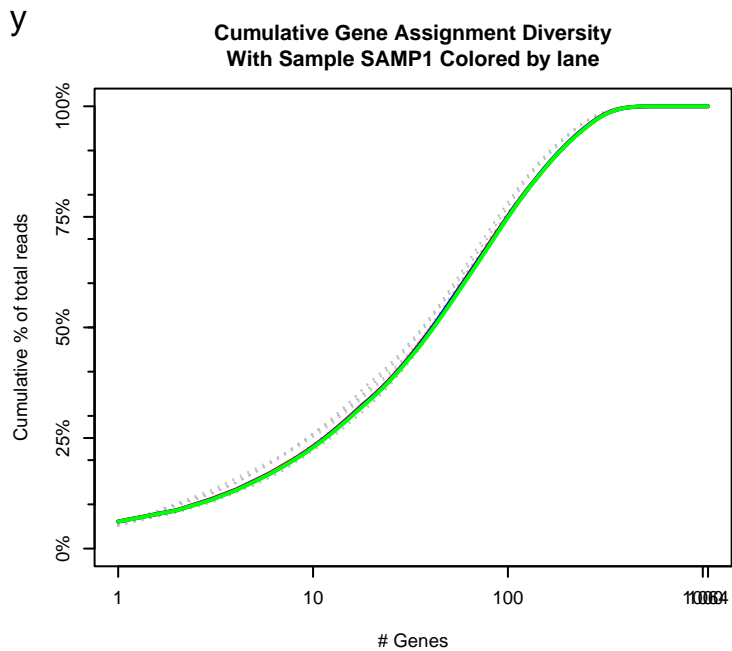
f



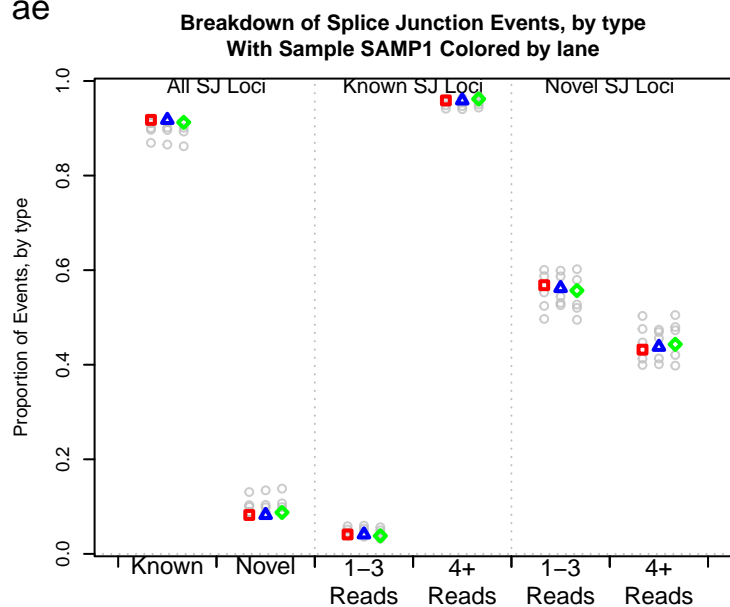




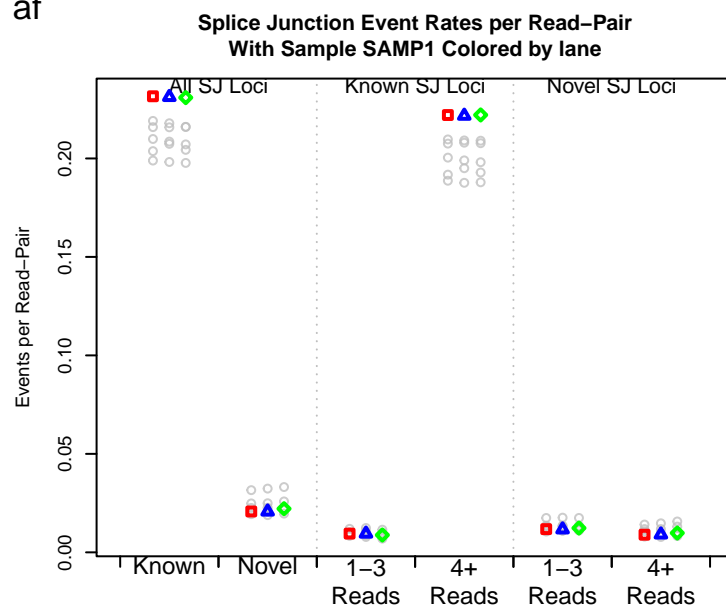




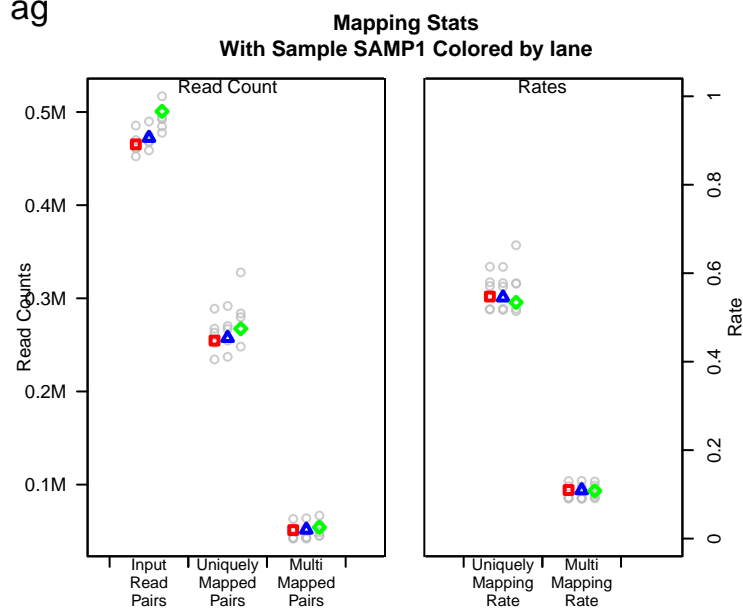
ae



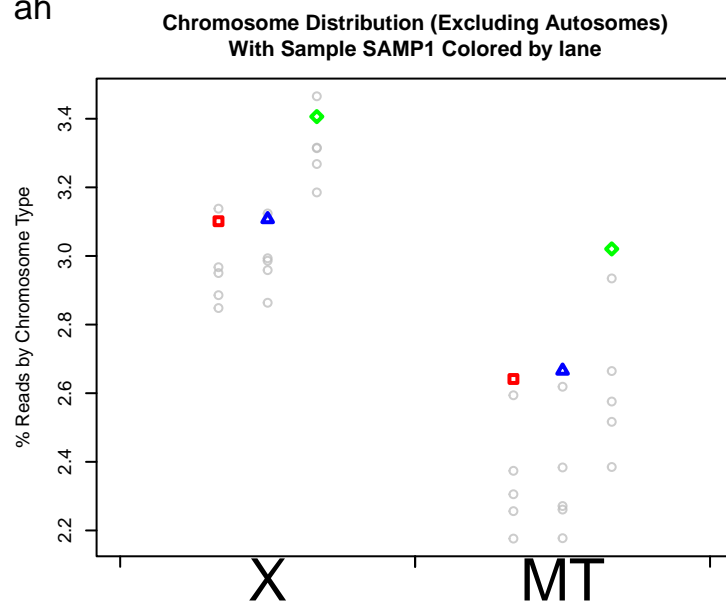
af



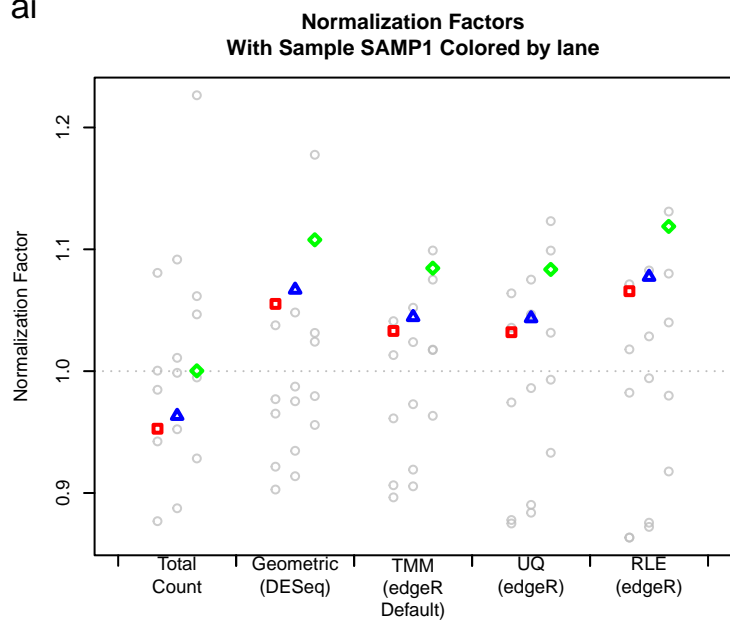
ag



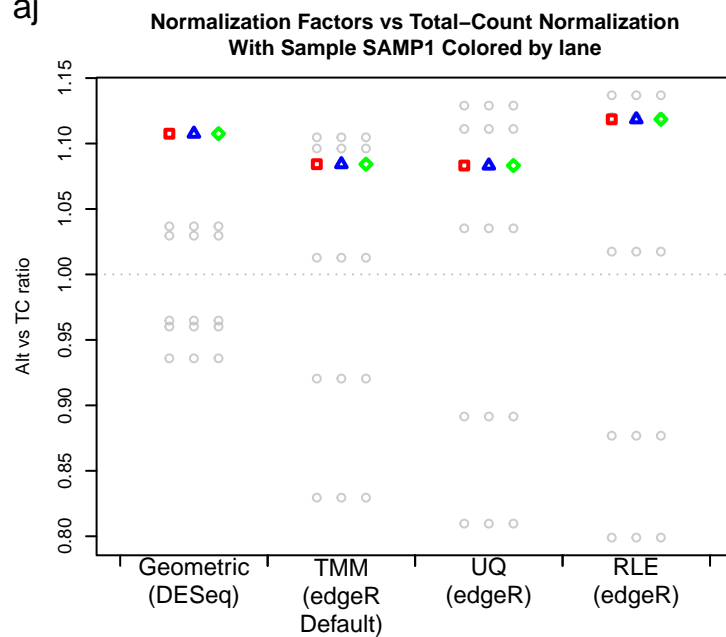
ah



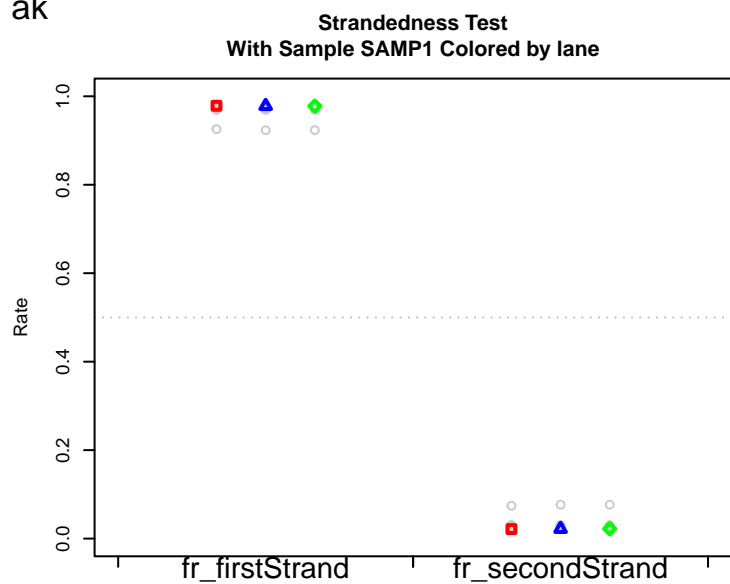
ai



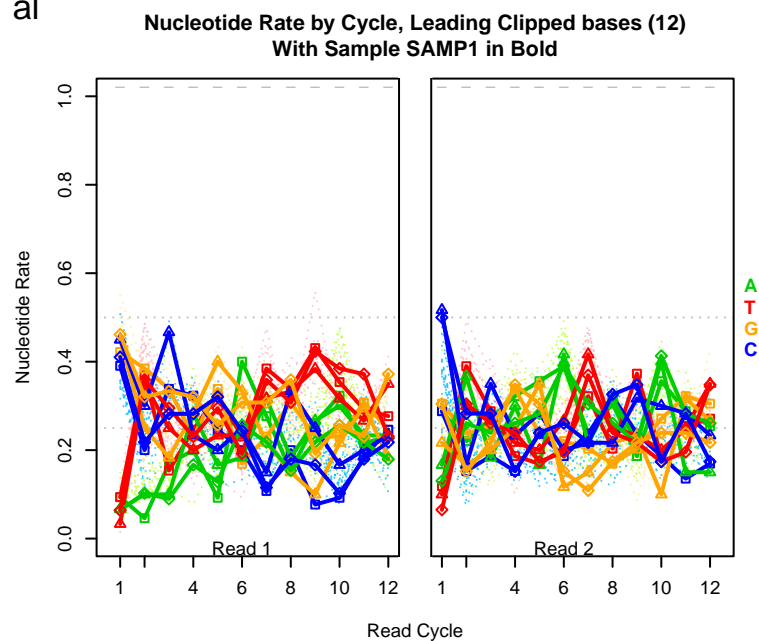
aj



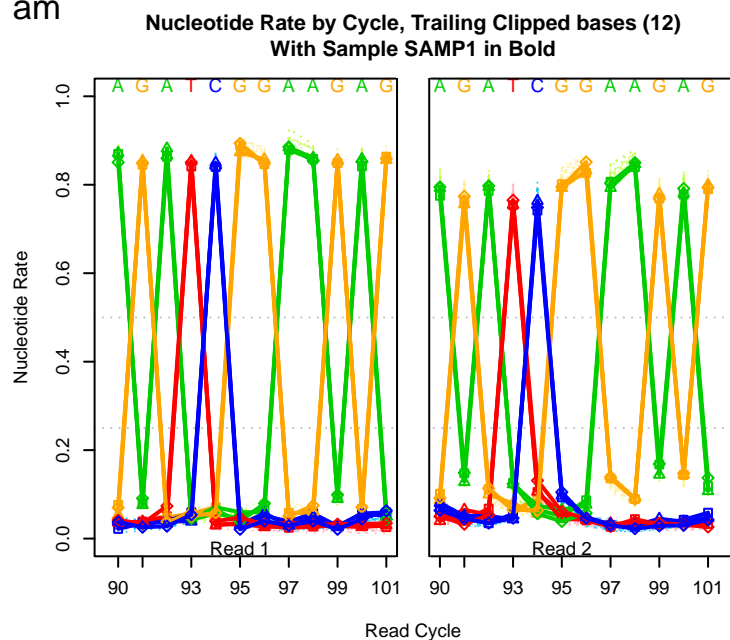
ak



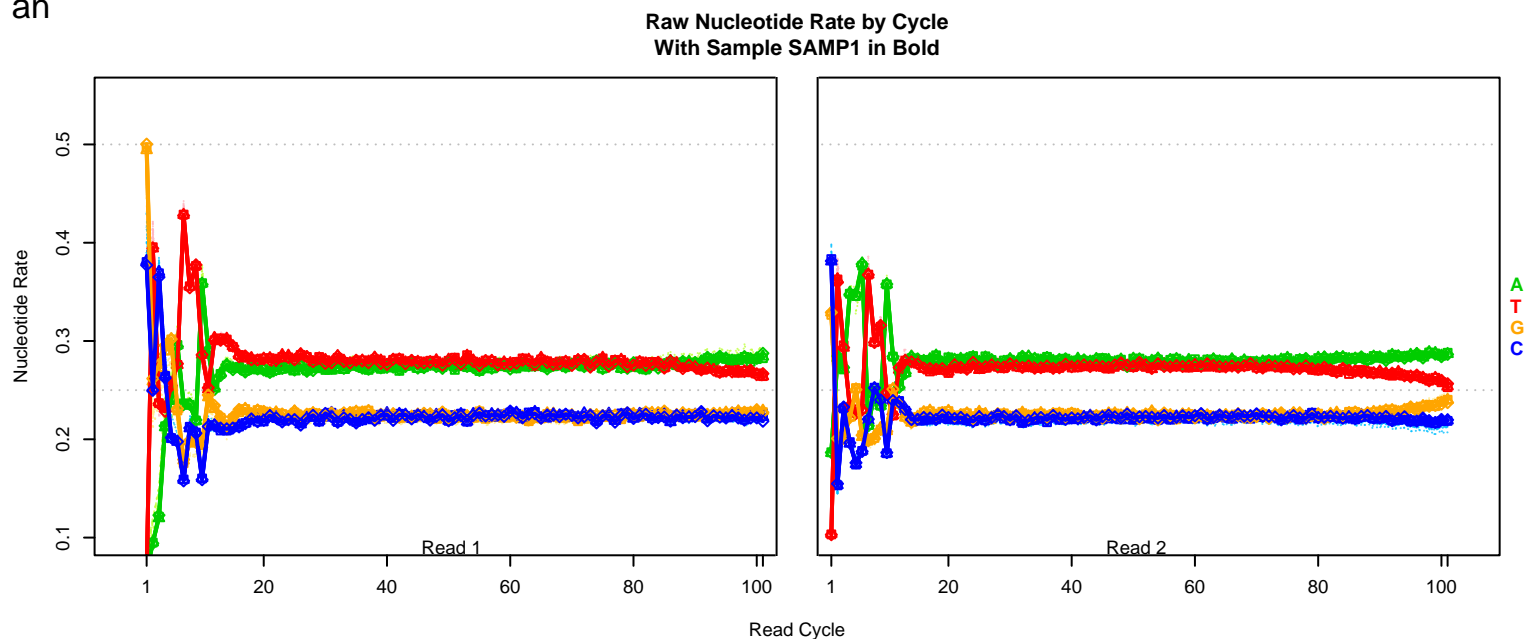
al



am



an



ao

Nucleotide Rate by Cycle, Aligned bases only  
With Sample SAMP1 in Bold

