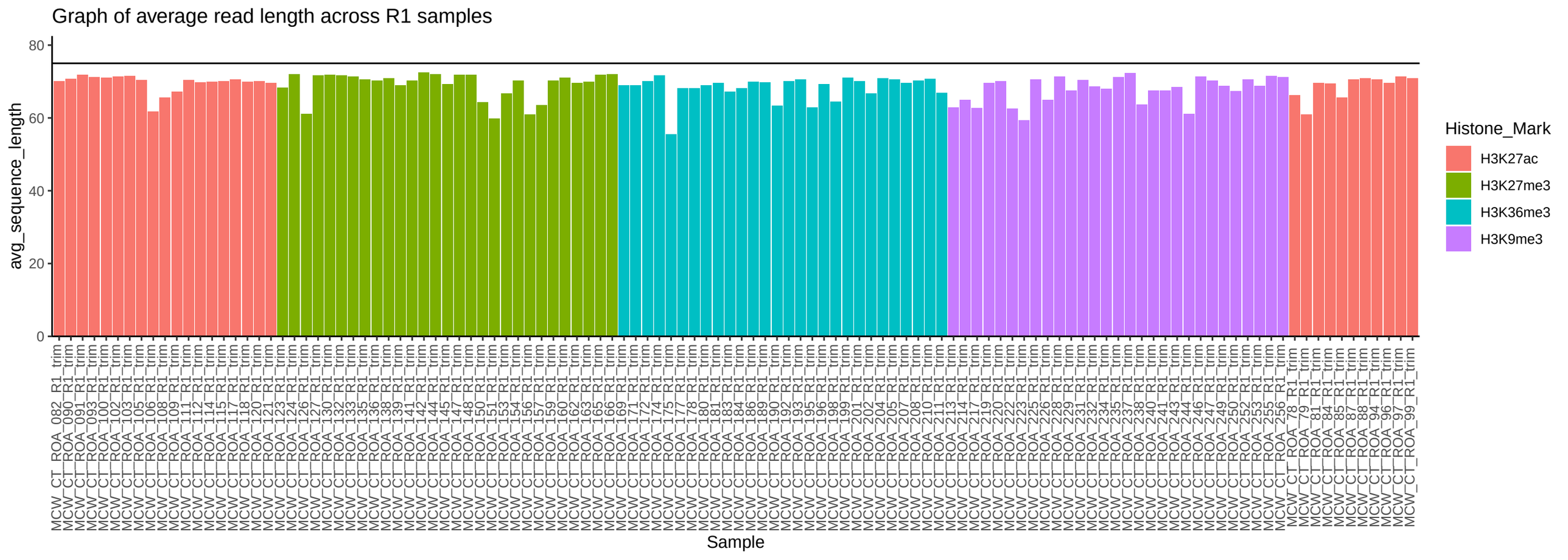
Graph of average read length across R1 samples
avg_sequence_length
80
60
40
20
0
Sample
Histone_Mark
H3K27ac
H3K27me3
H3K36me3
H3K9me3