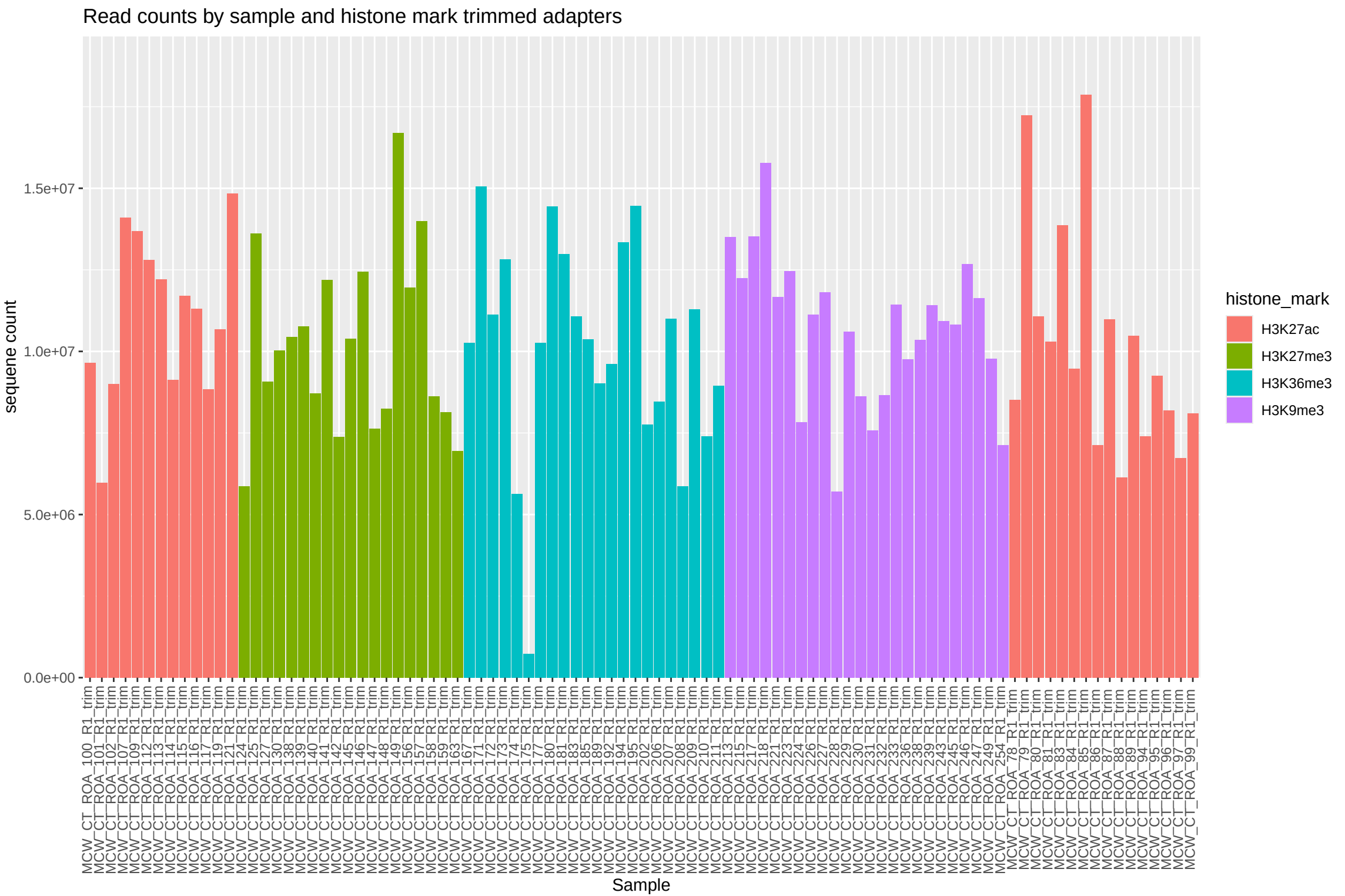
Read counts by sample and histone mark trimmed adapters
sequene count
1.5e+07
1.0e+07
5.0e+06
0.0e+00
Sample
histone_mark
H3K27ac
H3K27me3
H3K36me3
H3K9me3