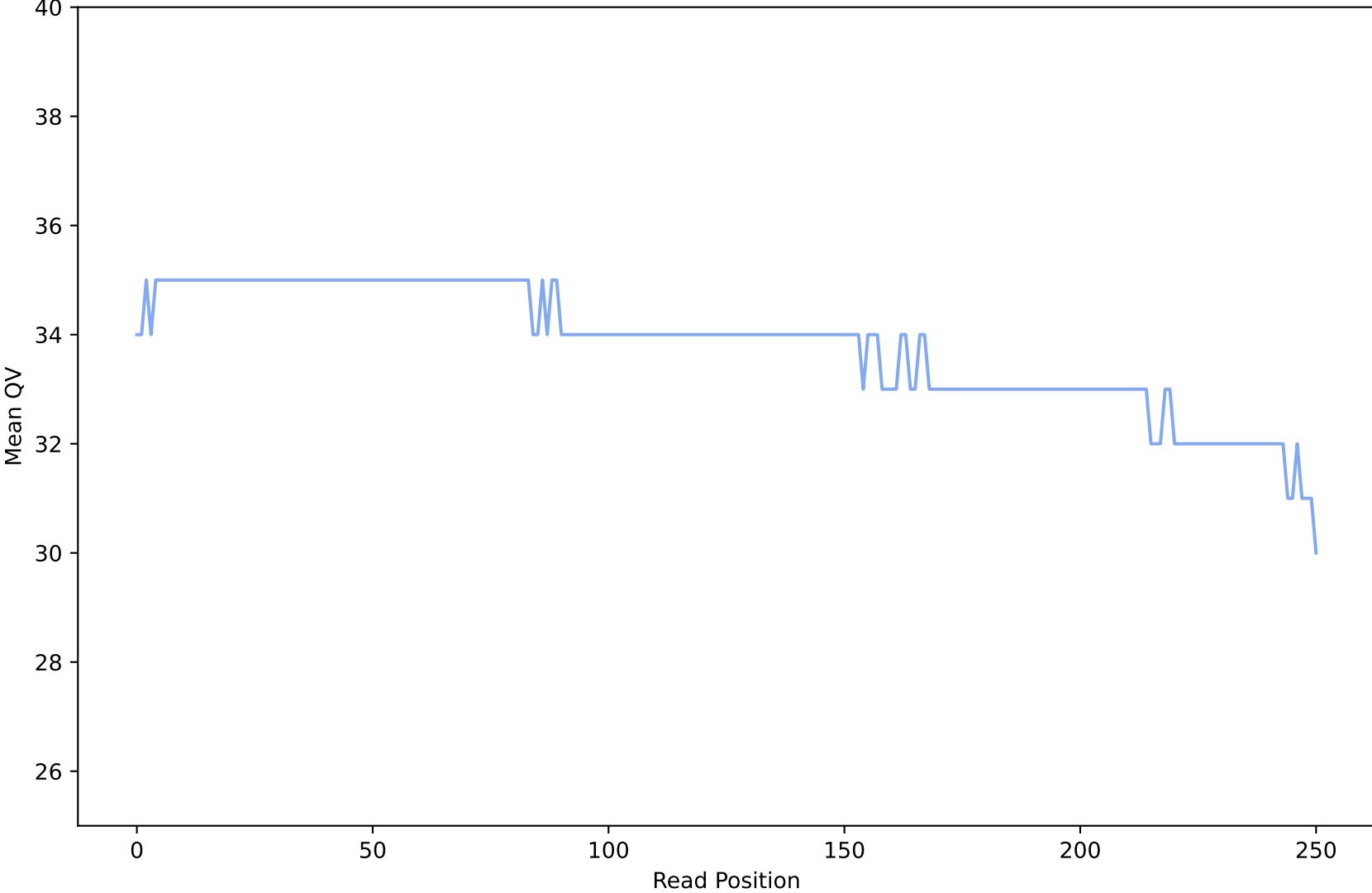
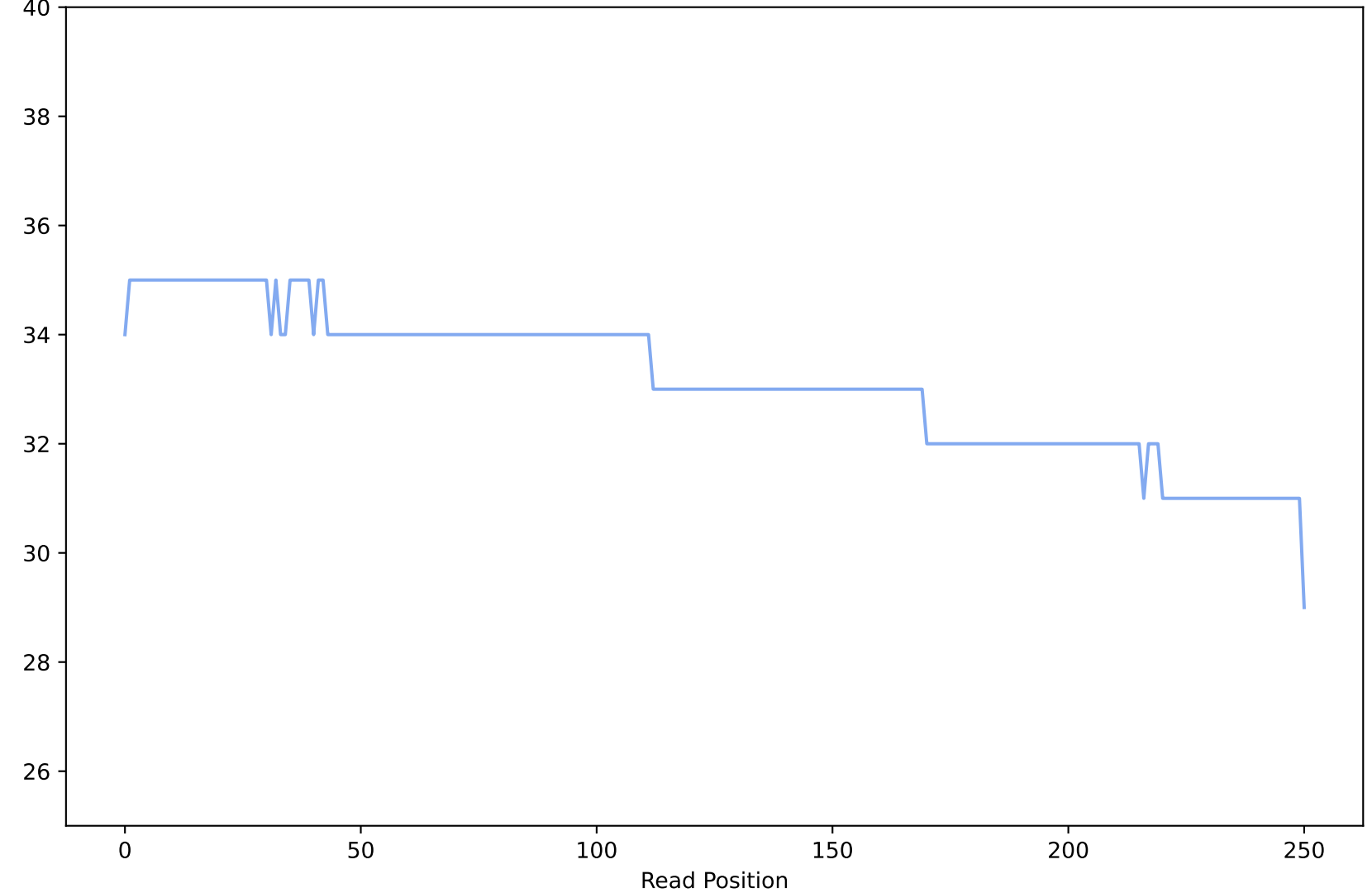


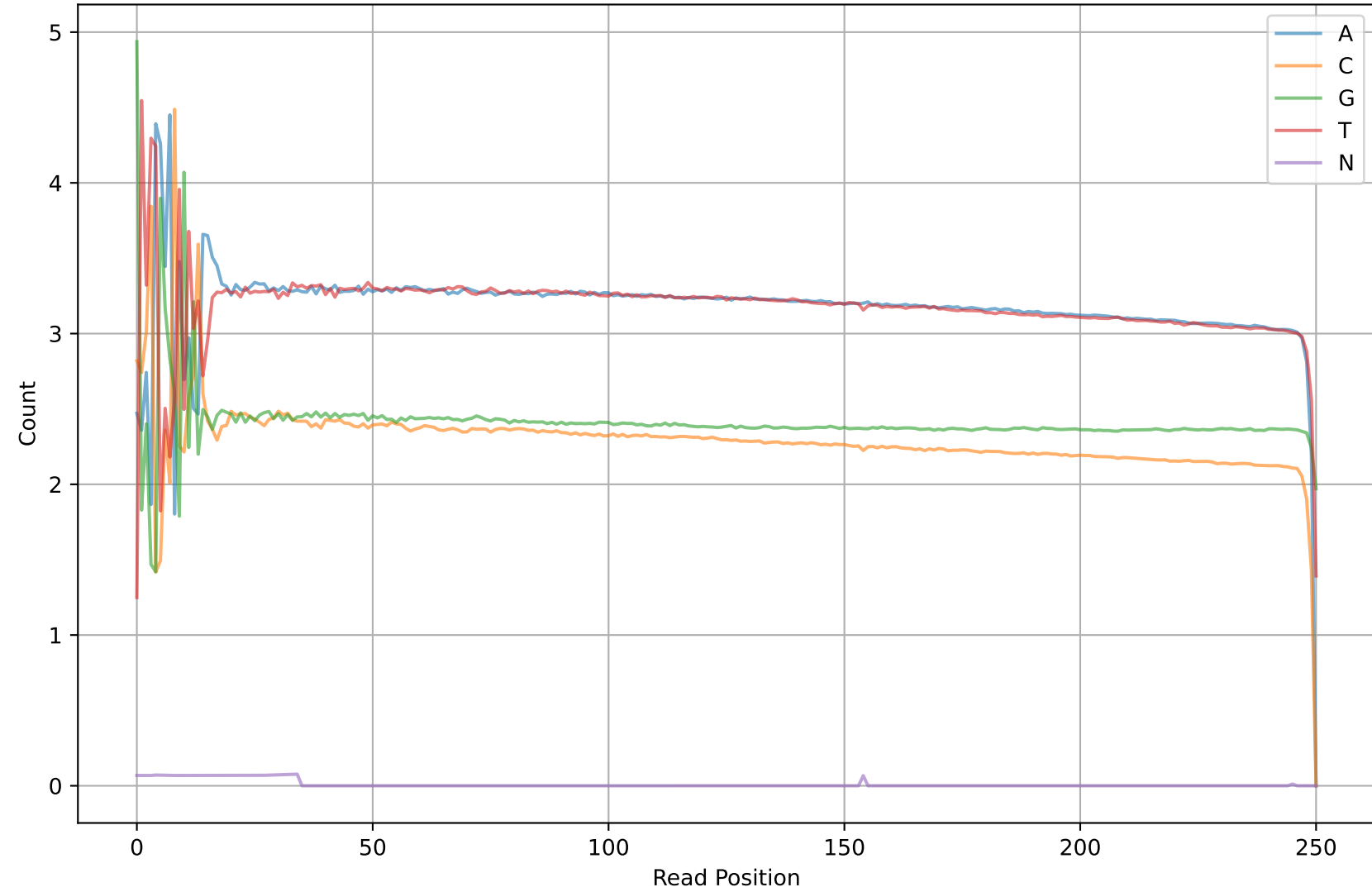
Mean QV at Read Position (R1)



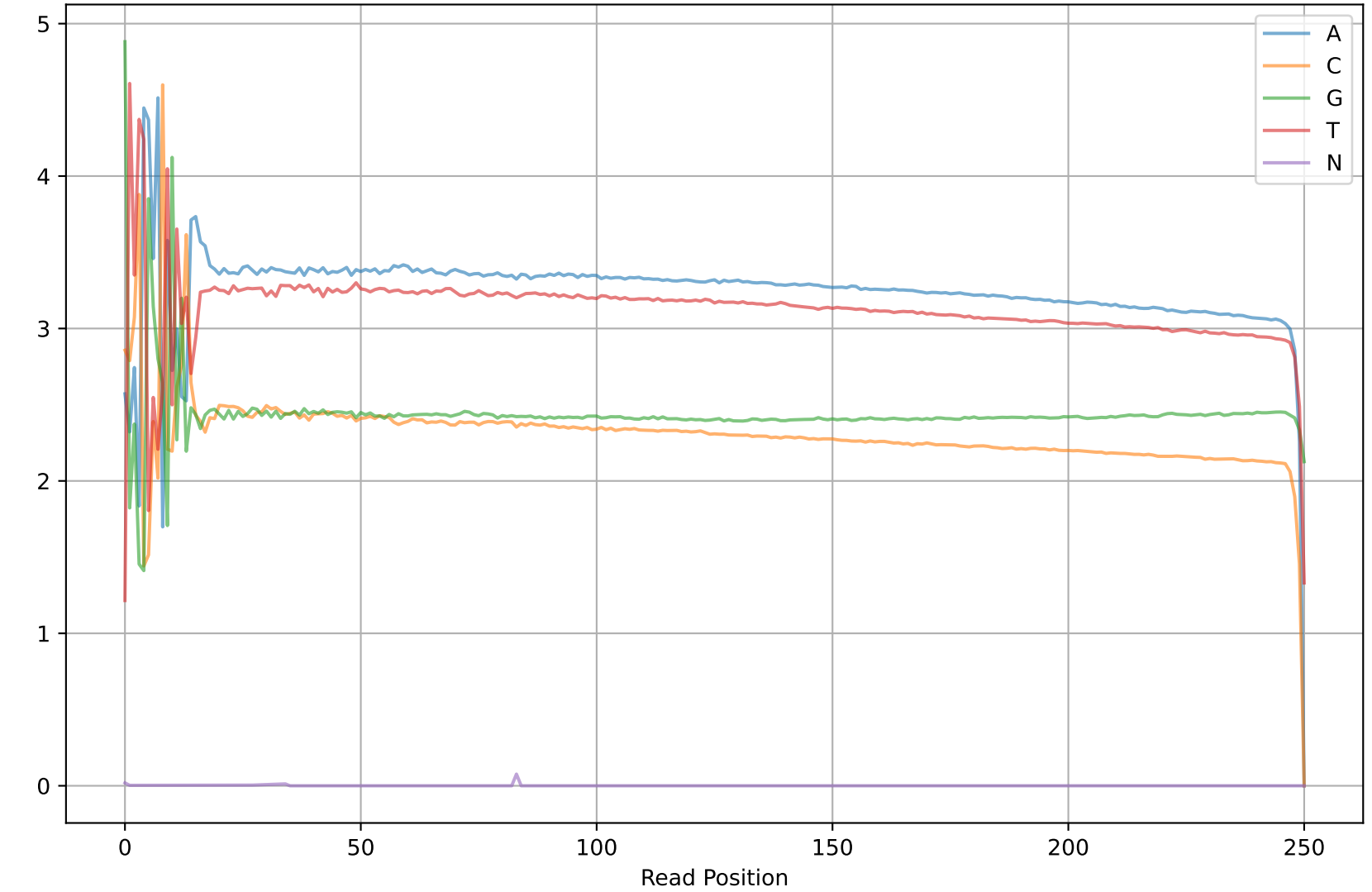
Mean QV at Read Position (R2)



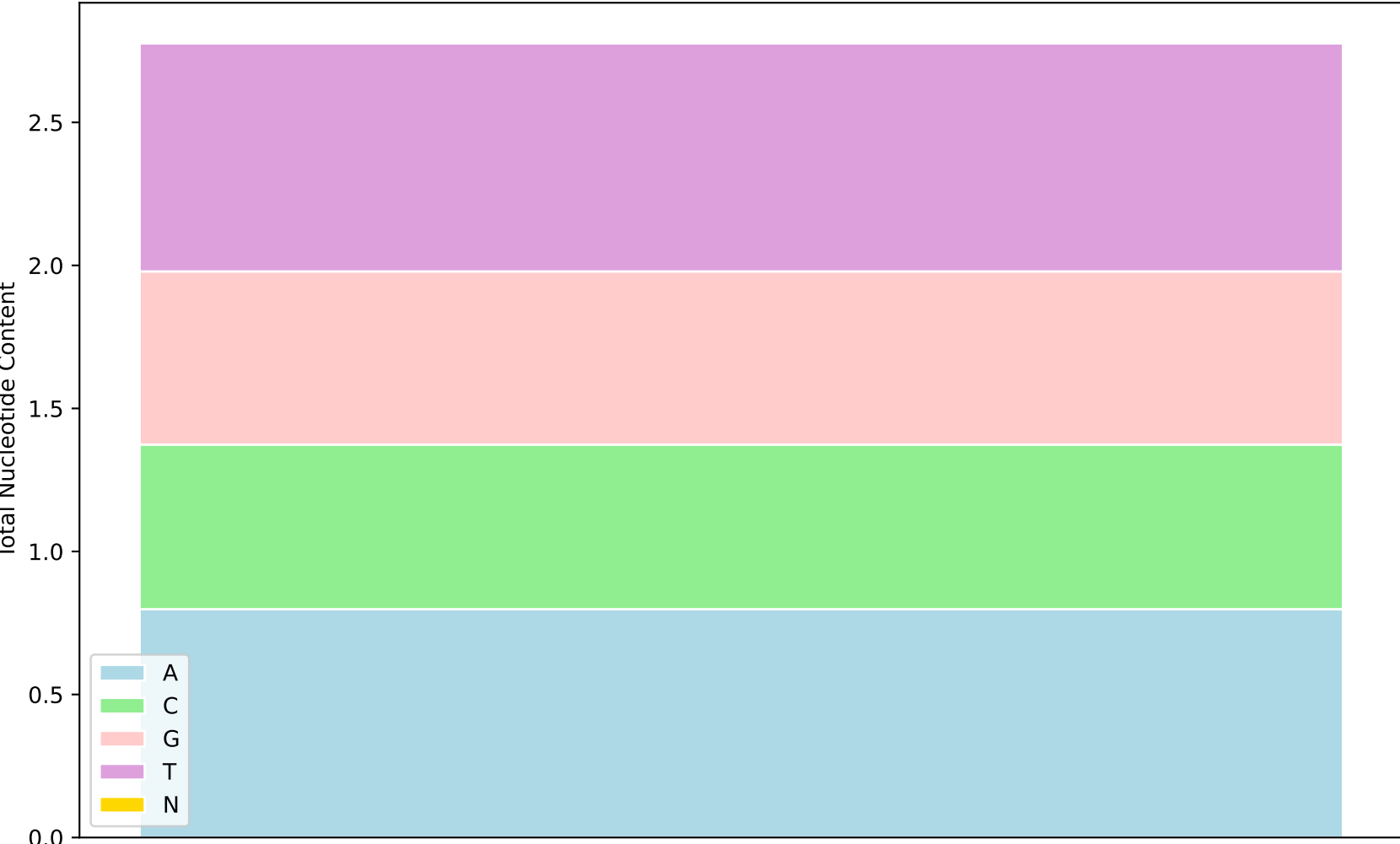
Nucleotide Counts for R1 File



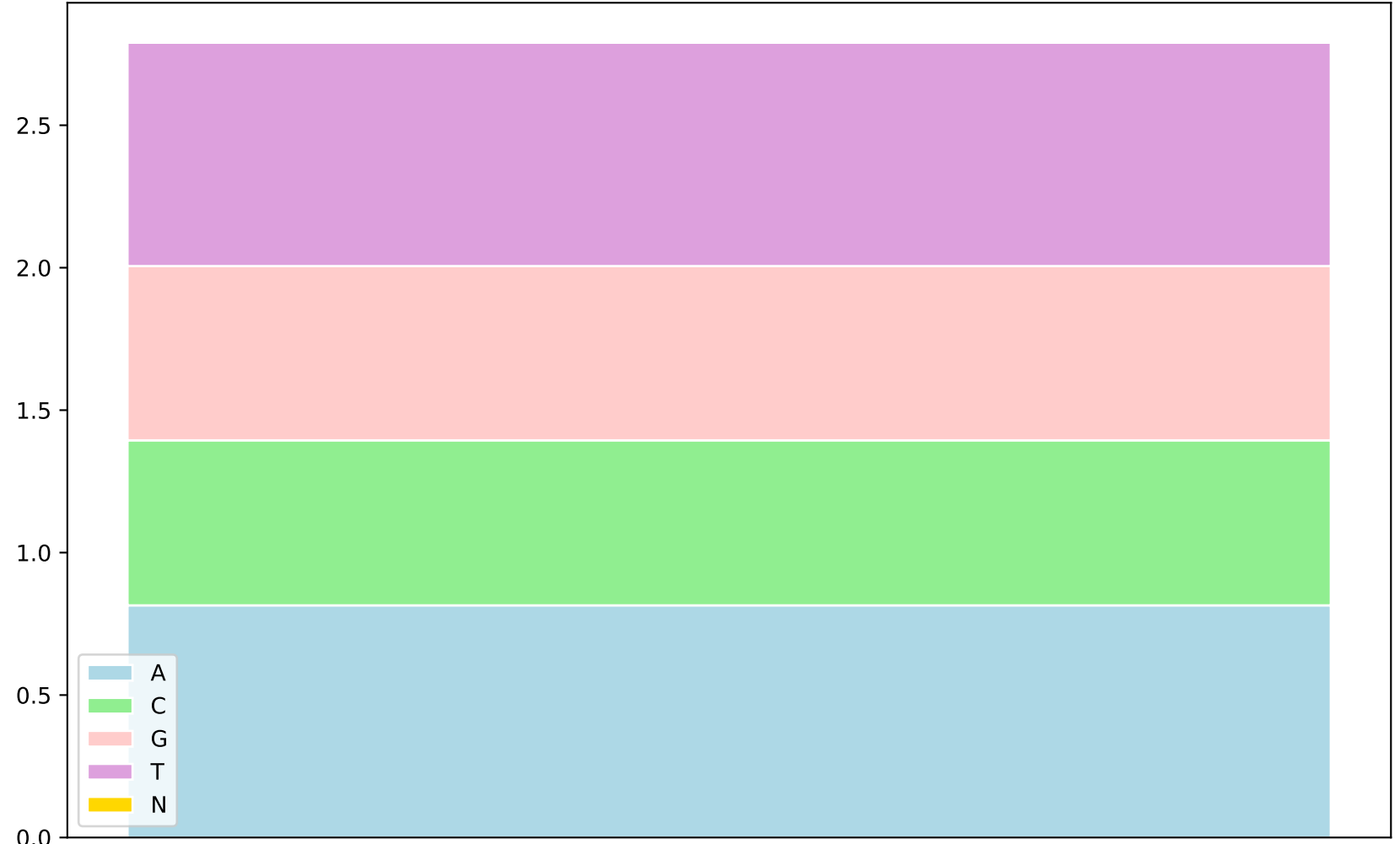
Nucleotide Counts for R2 File



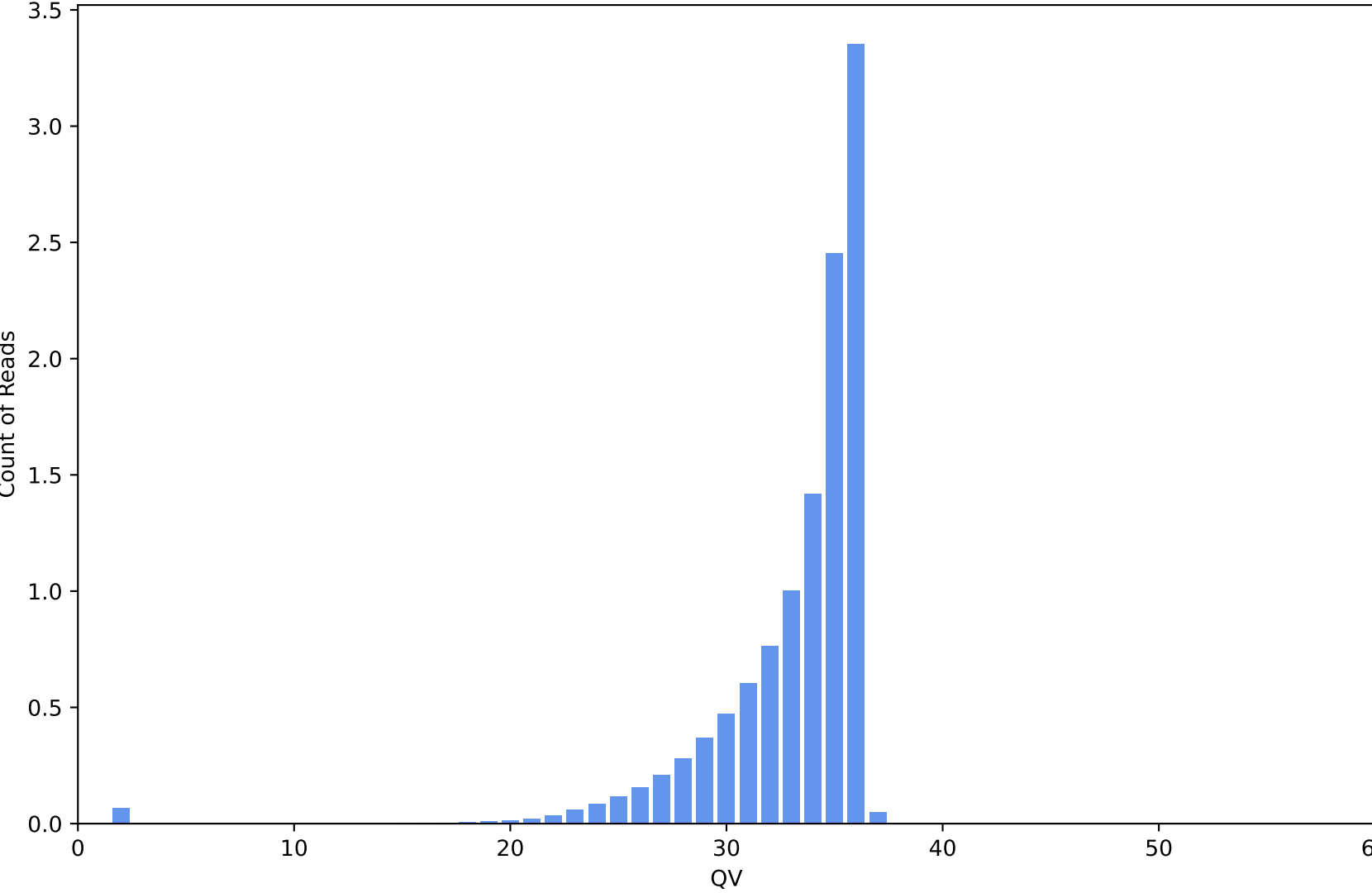
Total Nucleotide Content for R1 File



Total Nucleotide Content for R2 File



Mean QV Per Read for R1 File (QV 0 to 60)



Mean QV Per Read for R2 File (QV 0 to 60)

