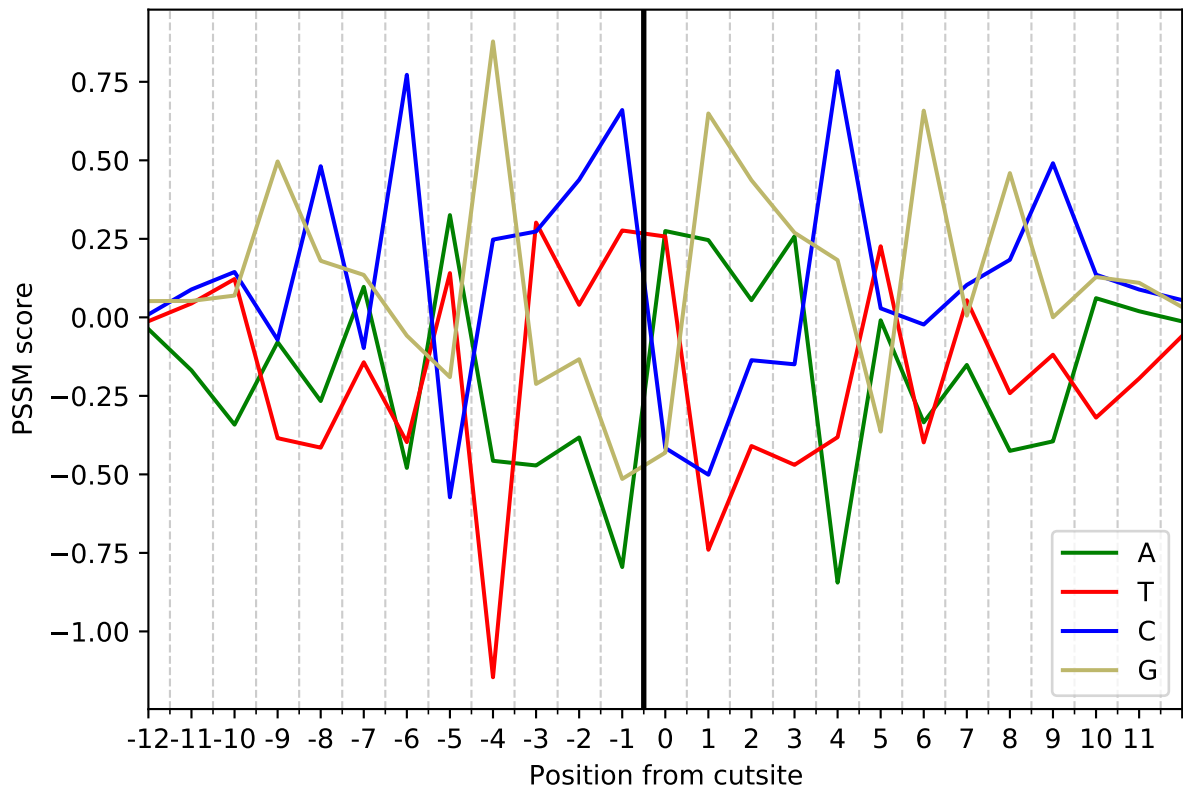
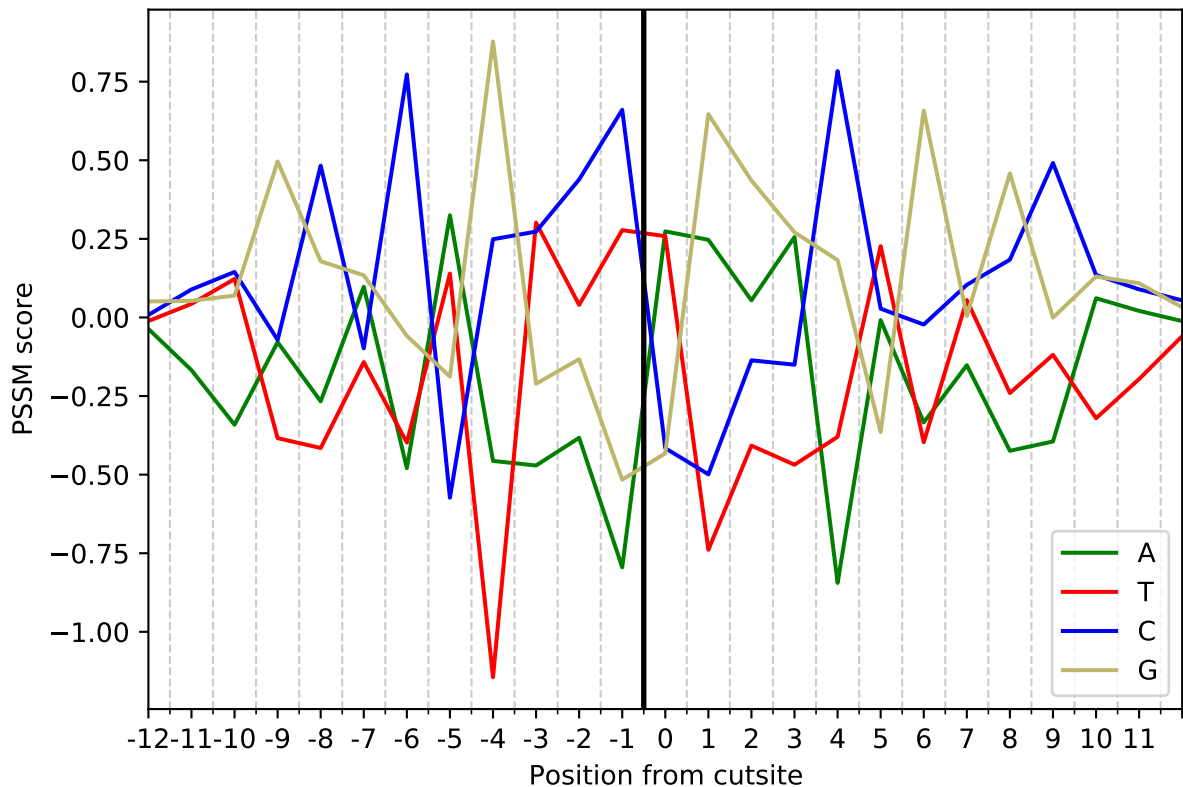


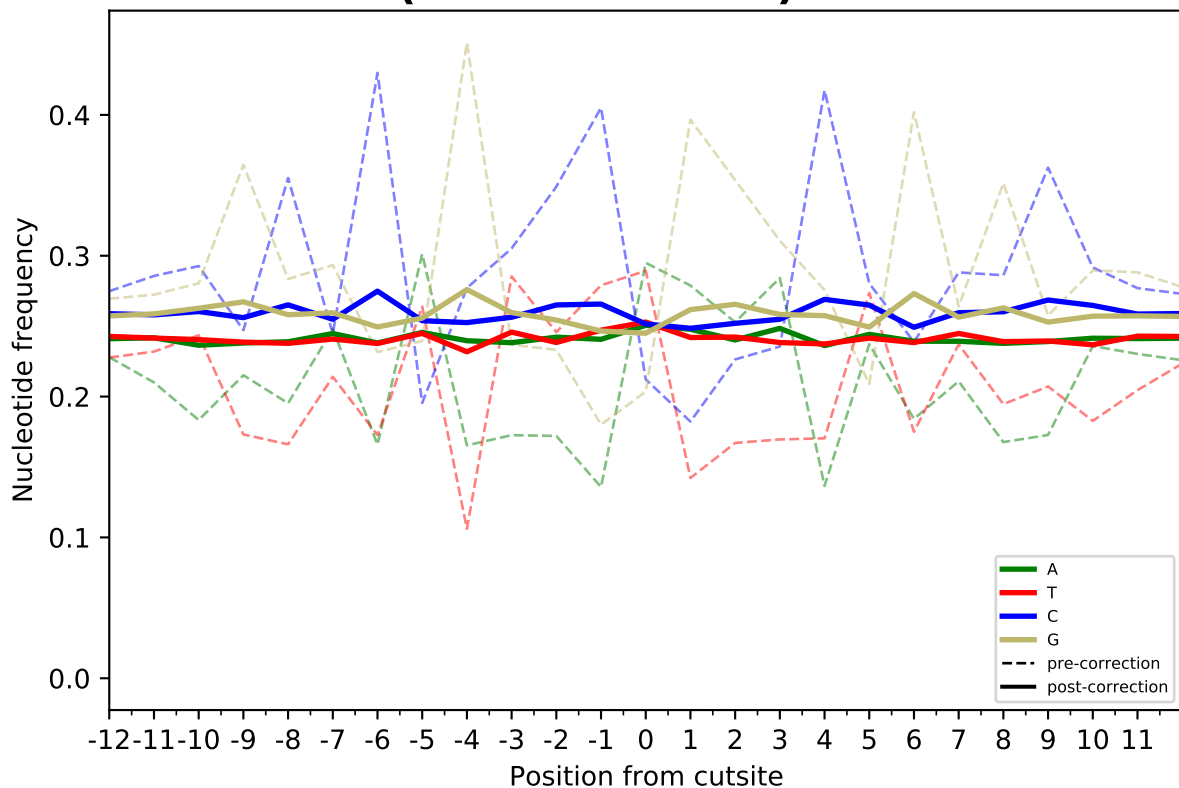
# Tn5 insertion bias of reads (forward)



# Tn5 insertion bias of reads (reverse)



# Nucleotide frequencies in corrected reads (forward strand)



# Nucleotide frequencies in corrected reads (reverse strand)

