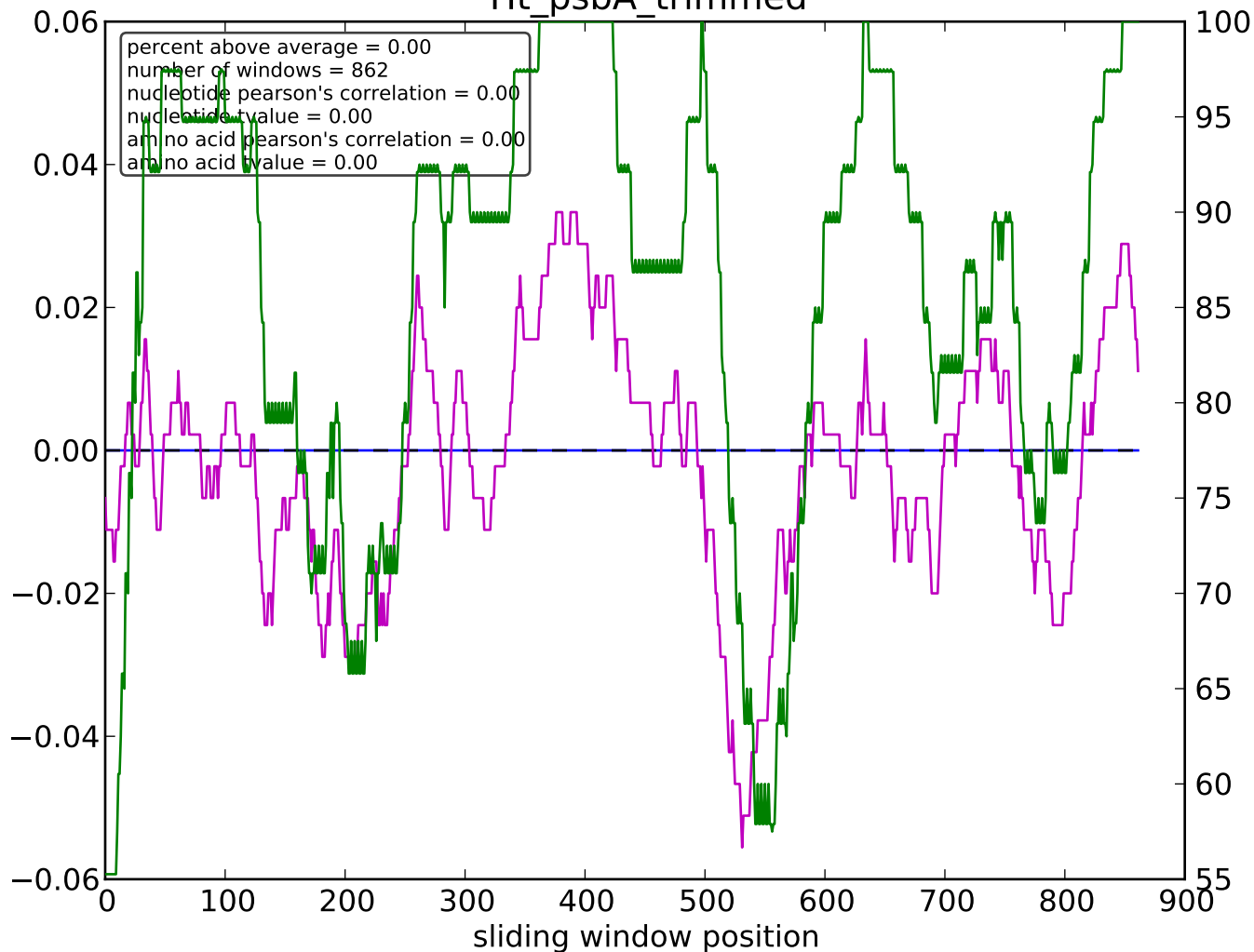


# Ht\_psbA\_trimmed

percent above average = 0.00  
 number of windows = 862  
 nucleotide pearson's correlation = 0.00  
 nucleotide tvalue = 0.00  
 amino acid pearson's correlation = 0.00  
 amino acid tvalue = 0.00

percent edited residues in RNA



percent sequence identity to reference  
 percent amino acid similarity to reference