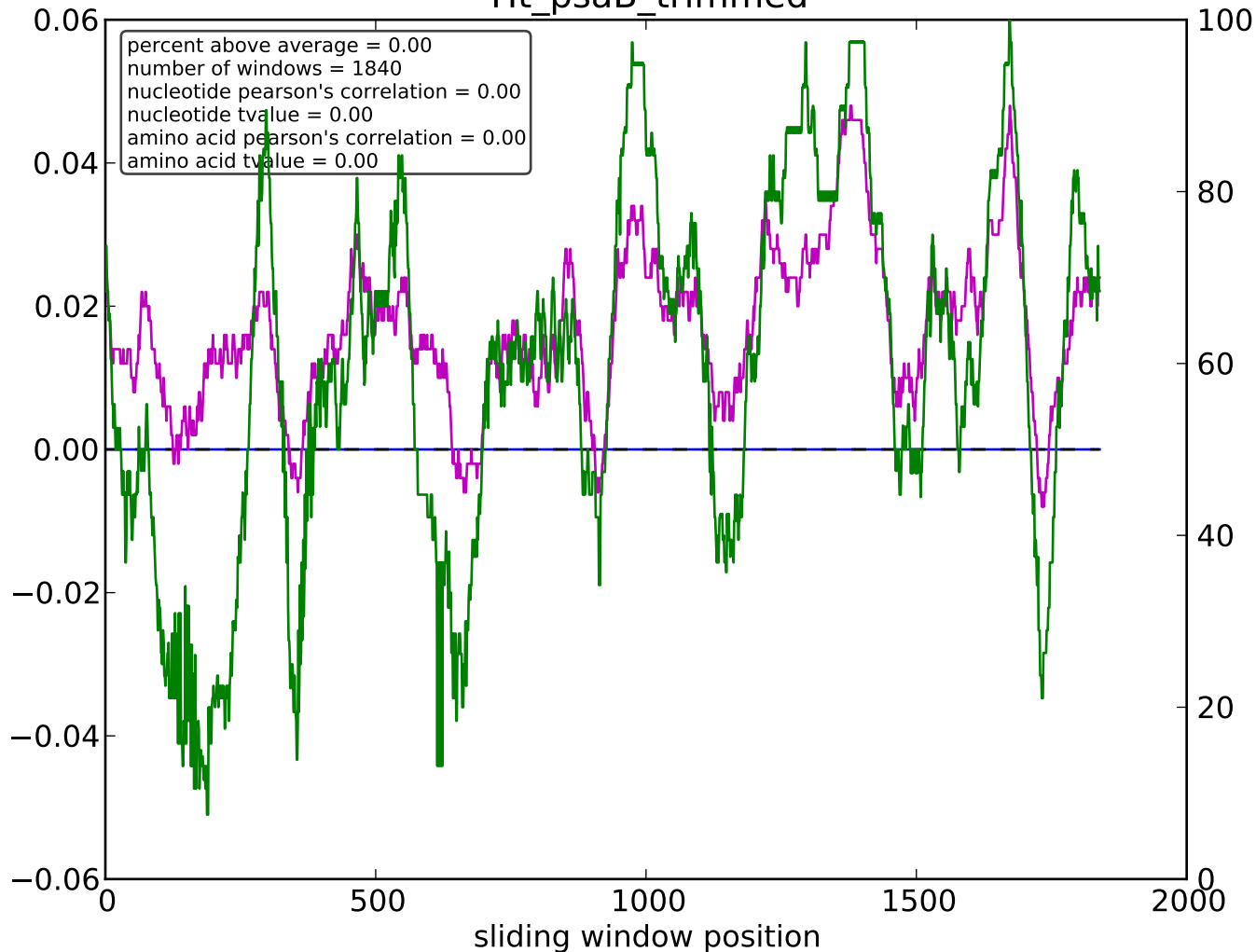


# Ht\_psaB\_trimmed

percent above average = 0.00  
number of windows = 1840  
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