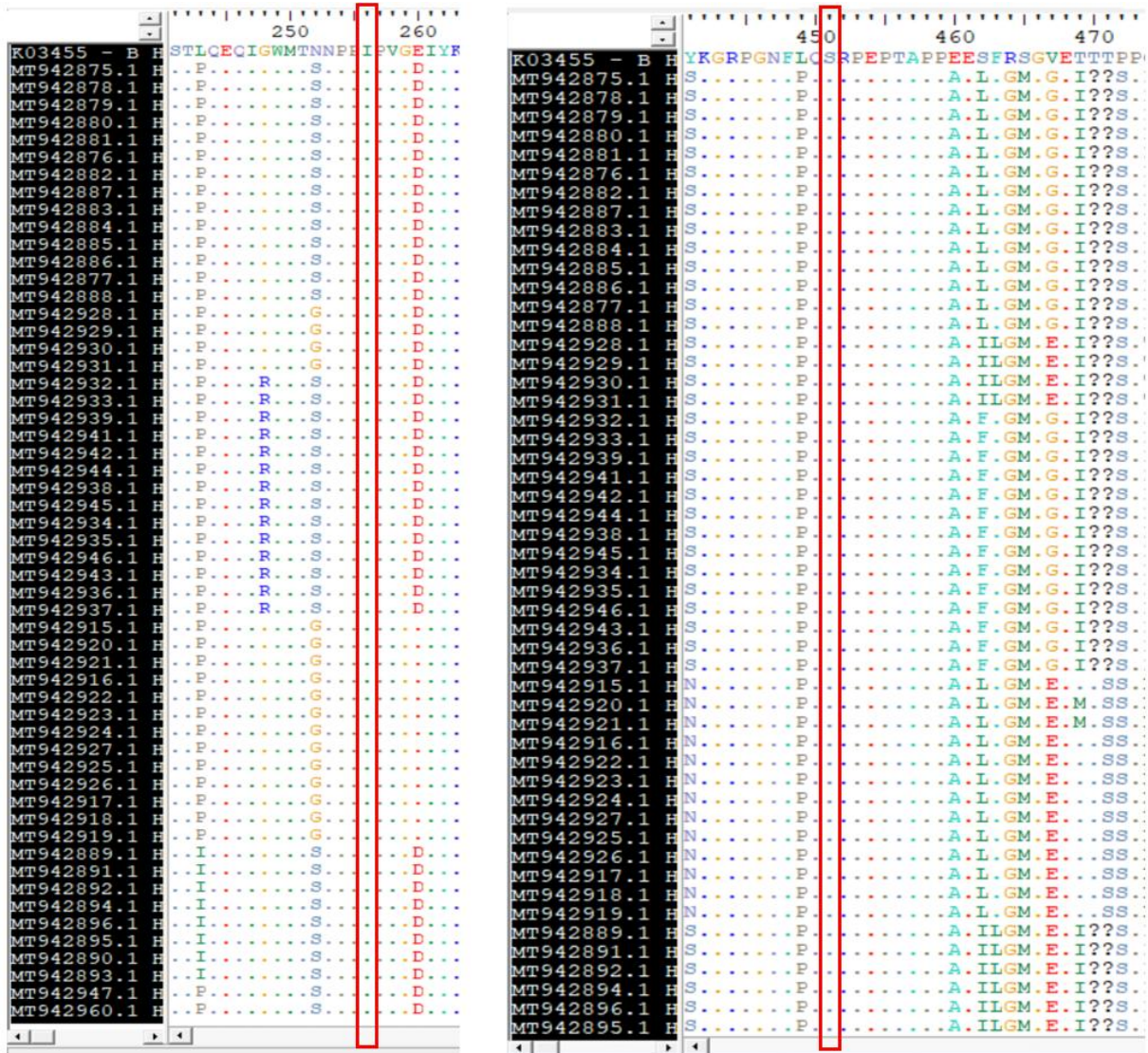


Informative sites I256 and S451 on HIV-1 subtype A recent reference sequences.



Supplementary figure 4: Performance of the informative sites in subtype A reference sequences. S54 amino acid was not conserved, and E55 amino acid only had a few mutations. Both I256 and S451 amino acids were highly conserved and thus show good performance for prediction of recent infection in HIV-1 subtype A reference sequences. HIV-1 – Human Immunodeficiency virus type 1.