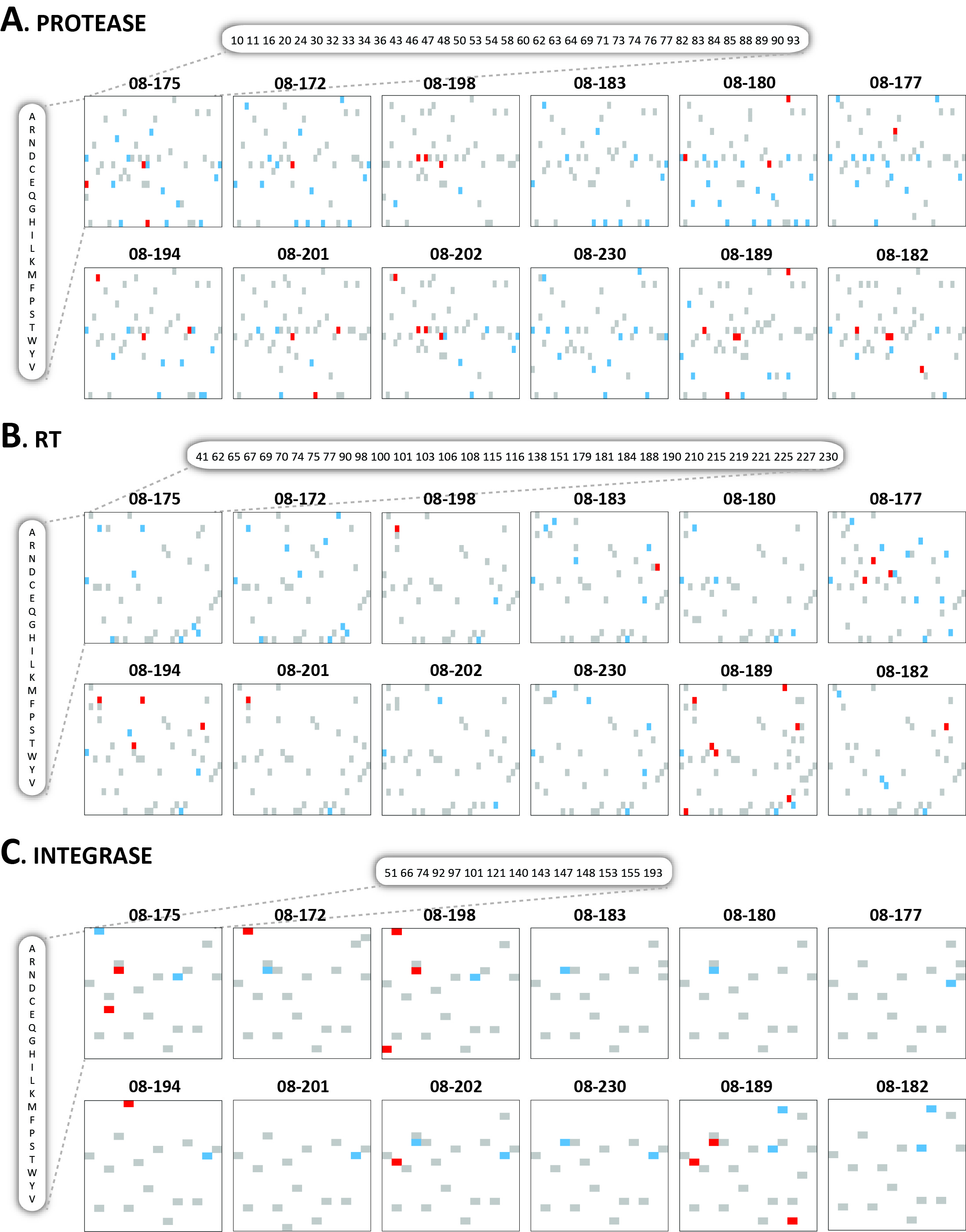
**Figure S1.** Amino acids detected in codons associated with drug resistance in the protease, RT, and integrase regions according to the Stanford University HIV Drug Resistance Database (<http://hivdb.stanford.edu>)



Drug resistance mutations with a frequency ≥20% (blue), >1% to <20% (red) or any other amino acid changes (grey) are indicated. Only amino acid substitutions with a frequency >1% are depicted.