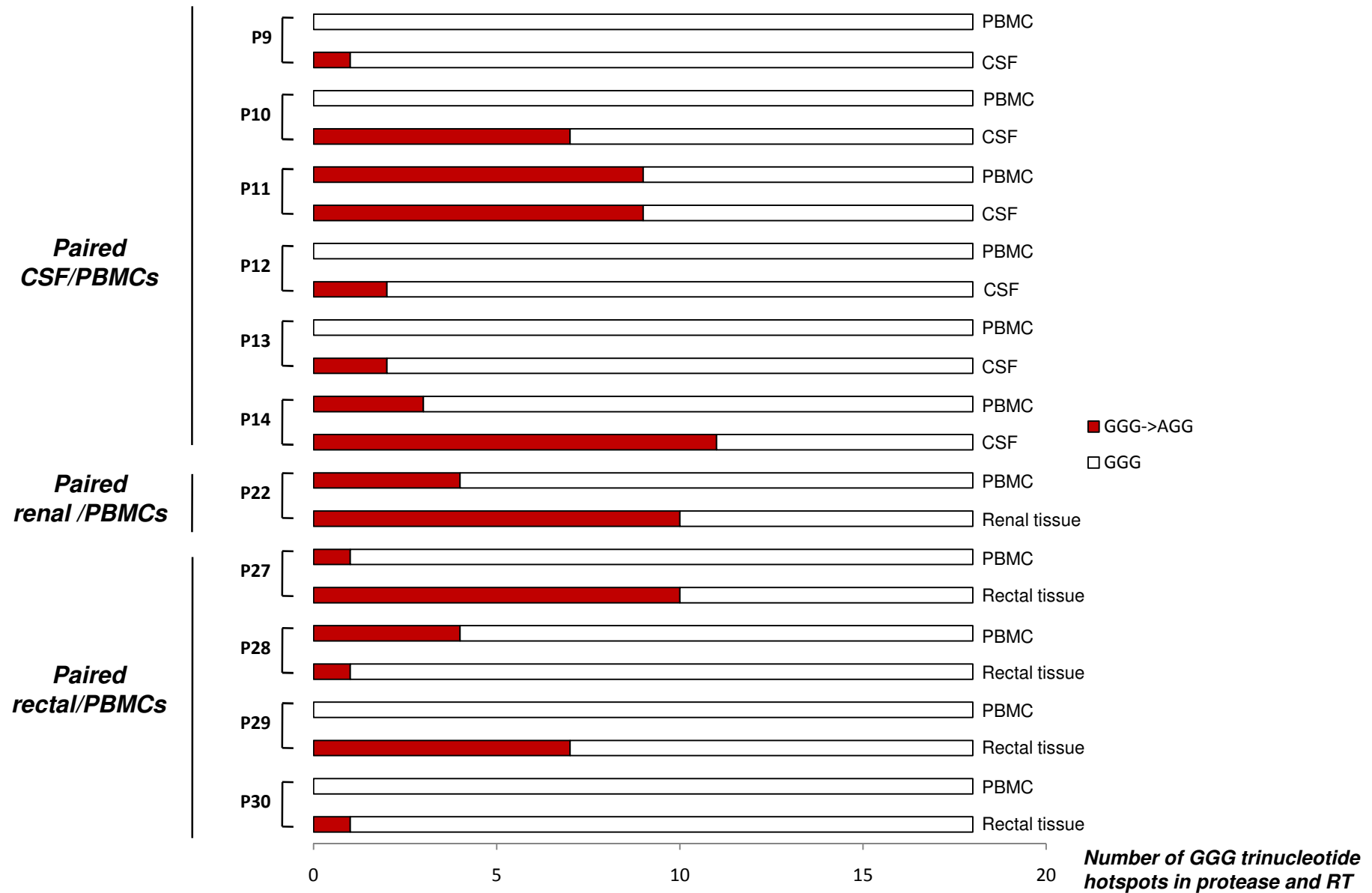


# Supplemental file 1



1 Supplementary file 1. Schematic representation G-to-A mutation in GGG trinucleotide  
2 context (strong preference for APOBEC3G editing). Overall, 18 GGG trinucleotide hotspots  
3 have been identified in protease and the analyzed region of RT (7 in protease and 11 in RT).  
4 GGG-to-AGG mutations were sought in protease and RT genes recovered from specific  
5 compartments of each patient. Red lines indicate the presence of GGG-to-AGG mutations;  
6 white lines indicate absence de GGG-to-AGG changes.  
7  
8