



Supplemental Figure 4. Phylogenetic tree of exon 2 sequences from all rhesus macaque DRB genes and human DRB9 and DRB6. The tree is constructed with the Neighbor-joining method and the Jukes-Cantor genetics distance model. The clustering of human and rhesus macaque *DRB6* (blue) and *DRB9* (green) exon 2 sequences indicates their genetic homology.