## Normalized Epitope Residue Frequencies (Z-scores) for CTLA4 -1.1 ALA --0.4 -0.2 -1.1 -1.2 0.1 0.1 -0.4 0.1 -1.1 -1.1 -0.7 ARG --0.4 -0.2 -0.2 -0.2 -0.8 0.1 -0.2 -0.4 0.1 -0.2 -0.4 -1.0 -0.4 -1.2 ASN --0.4 -1.1 -1.2 -0.8 0.1 -1.1 -1.4 -0.9 -1.1 -1.1 0.0 -1.0 ASP --1.2 -1.2 -1.1 -1.2 -0.8 0.1 -1.1 -1.4 -0.9 -1.1 -1.1 -1.1 - 3 CYS --1.2 -1.1 -0.8 -1.1 -1.1 -1.2 -1.2 -1.2 0.1 -1.4 -0.9 -1.1 -1.0 GLN --0.2 -1.1 -0.2 -0.8 -0.9 -0.2 -0.9 -0.2 0.4 -0.3 0.5 0.0 GLU -1.8 -0.8 2.2 1.6 1.1 1.8 1.9 1.6 GLY --0.2 -0.2 -0.2 0.1 -0.9 -0.2 -0.4 -0.9 -0.2 -0.4 -1.0 -0.4 - 2 HIS --1.2 -0.2 -1.1 -0.2 0.1 -0.9 -1.1 -0.4 -0.9 -0.2 -1.1 0.0 -0.8 Residue ILE -0.7 0.1 -0.9 0.7 0.1 0.8 -0.4 0.4 1.6 1.6 LEU -2.3 1.1 0.1 2.1 1.8 -1.0 1.4 LYS --0.2 -0.2 0.1 -0.9 -0.2 -0.4 0.1 -0.2 -0.4 -0.1 MET -0.5 0.7 0.8 0.1 1.1 0.7 1.1 0.8 1.2 -1.0 0.7 PHE --1.2 -1.1 -1.2 -1.1 -1.2 -0.8 -1.4 -0.9 -1.1 -0.4 -1.1 -0.9 0.0 PRO -1.4 1.6 1.8 2.9 0.1 0.8 2.9 -0.9 SER --0.4 -0.2 -0.2 -0.2 0.1 -0.9 -0.2 -0.4 -0.9 -1.1 -1.1 -0.6 THR --1.2 -1.2 -0.2 -0.8 0.7 -0.2 1.9 TYR -1.4 1.6 1.5 1.8 2.0 2.2 1.6 1.5 2.1 1.8 1.0 2.0 VAL --0.4 -0.2 -0.2 -0.2 -0.8 0.1 -0.2 -0.4 0.1 -0.2 0.4 -1.0 -0.3 543 SIN 7014 PDB ID