Normalized Paratope Residue Frequencies (Z-scores) for PDL1 -0.2 -0.9 -1.1 -0.3 ALA --1.1 -0.8 -0.2 ARG -0.1 -0.3 -0.2 -0.2 -0.2 -0.3 -0.2 0.1 -0.8 -1.1 ASN --0.9 -0.2 -0.2 -0.9 0.3 ASP --1.1 -0.3 -0.2 -0.9 -0.2 -0.3 -0.2 - 3 GLU --1.1 -0.8 -0.9 -0.9 -0.9 -1.1 GLY -0.1 -0.3 HIS -0.1 -0.8 -0.2 -0.9 -1.1 0.3 -0.9 -1.1 ILE -0.3 -0.9 -1.1 -0.9 -1.1 Residue - Nan -1.1 -0.3 -0.2 -0.2 -0.2 -0.3 -0.3 -0.8 -0.9 -0.2 0.7 -0.9 0.1 -0.9 -0.2 PHE -0.3 -0.2 -0.2 -0.3 -0.9 -0.2 PRO -0.1 -0.3 0.5 -0.9 -1.1 -0.9 -0.2 SER -2.4 2.7 2.0 2.1 2.7 2.2 1.7 SO4 --1.1 -0.8 -1.1 -0.9 -0.2 THR -0.1 -0.9 -1.1 -0.9 TRP -0.1 -0.8 -0.3 -0.2 TYR -1.2 2.0 2.1 2.2 2.7 VAL --1.1 -0.3 -0.9 -0.2 -0.2 -0.9 -0.3 -1.1 STIA Seil