Normalized Paratope Residue Frequencies (Z-scores) for PDL1 -0.2 -0.9 -1.1 -0.3 ALA --1.1 -0.8 -0.2 -0.5 ARG -0.1 -0.3 -0.2 -0.2 -0.2 -0.3 -0.2 -0.2 ASN -0.1 -0.8 -0.9 -0.2 0.3 -1.1 -0.6 ASP --1.1 -0.3 -0.2 -0.9 -0.2 -0.3 -0.2 -0.3 - 3 GLU --1.1 -0.8 1.6 -0.9 -1.1 -0.5 GLY -0.1 -0.3 HIS -0.1 -0.8 -0.2 -0.9 -1.1 0.3 -1.1 -0.7 ILE -0.3 -0.9 -0.9 -1.1 -0.9 -0.9 -1.1 -0.7 Residue - SAT e -1.1 -0.3 -0.2 -0.2 -0.2 -0.3 -0.3 -0.3 -0.8 -0.9 -0.2 0.7 -0.9 -0.2 0.1 -0.9 -0.6 PHE -0.3 -0.2 -0.2 -0.2 -0.3 -0.3 PRO -0.1 -0.3 -0.9 -1.1 -0.9 -0.2 -0.3 SER -2.4 2.7 2.0 2.7 2.1 2.2 1.7 2.7 SO4 --1.1 -0.8 -0.9 -0.9 -1.1 -0.9 -0.9 -0.2 -1.1 THR -0.1 -0.9 -1.1 0.2 TRP -1.3 -0.2 0.1 -0.8 -0.3 TYR -2.0 2.1 2.2 2.7 2.3 VAL --1.1 -0.3 -0.9 -0.2 -0.2 -0.9 -0.3 -1.1 -0.7 1485 58il 442 PDB ID