Normalized Interface Residue Frequencies (Z-scores) for PDL1 ALA -1.1 -0.7 1.3 -0.7 -0.5 1.0 0.0 1.2 0.2 ARG -0.2 0.3 0.1 0.2 0.4 0.4 -0.0 0.0 0.1 ASN -0.2 -0.7 -0.7 -0.5 -0.9 -0.1 -0.9 -0.7 -0.5 ASP -0.2 0.8 0.4 0.2 0.5 1.2 0.4 0.2 0.5 - 3 GLN --0.7 -0.2 -0.7 -1.2 -0.7 -0.9 -0.4 -0.9 -0.8 GLU -0.2 -0.2 -0.2 1.5 0.4 -0.4 -0.3 -0.4 -0.1 GLY -1.1 0.4 1.3 -0.1 -0.3 1.0 -0.2 1.2 0.3 HIS -0.2 -0.2 0.1 -0.9 -0.2 -1.2 -0.7 -0.4 -0.6 - 2 HOH --1.6 -1.3 -1.2 2.0 4.2 -1.4 4.4 1.2 3.6 ILE -2.0 1.0 -0.7 -1.2 -0.5 0.1 -0.4 -0.4 -0.4 LEU --0.7 -0.7 -0.7 -0.7 -0.5 -0.4 -0.3 1.2 -0.5 LYS --0.7 -0.2 -0.7 -0.1 -0.0 -0.9 -0.3 -0.9 -0.5 MET --0.7 -0.2 -0.2 -0.1 -0.0 -0.4 -0.3 -0.9 -0.4 PHE -0.2 -0.2 -1.2 -0.1 -0.3 -0.9 -0.3 -0.4 -0.5 PRO --0.7 -0.7 -0.2 -1.2 -0.7 0.1 -0.5 -0.4 -0.7 SER -2.0 2.7 1.3 1.5 -0.0 1.5 0.2 0.7 0.9 SO4 --1.6 -1.3 -1.2 -1.2 -0.7 -1.4 -0.5 -0.9 -1.1 THR --0.7 1.0 -0.2 -0.1 -0.3 0.1 -0.5 -0.4 -0.4 TRP --0.7 -1.3 0.3 0.4 -0.0 0.5 -0.4 -0.9 -0.4 TYR -1.1 2.1 2.8 2.0 0.6 2.9 0.2 2.3 1.6 -0.7 VAL --0.2 -0.2 0.4 0.2 -0.4 -0.2 -0.9 -0.2 57817 574 soil PDB ID