Normalized Interface Residue Frequencies (Z-scores) for PDL1 ALA -1.1 -0.7 1.3 -0.7 -0.5 1.0 0.2 1.2 0.2 0.3 ARG -0.4 0.4 0.1 0.2 0.1 ASN -0.2 -0.7 -0.7 -0.7 -0.9 -0.1 -0.9 ASP -0.2 0.4 0.8 0.4 0.2 0.2 1.2 - 3 GLN --0.7 -0.2 -0.7 -1.2 -0.7 -0.9 -0.4 -0.8 GLU -0.2 -0.2 -0.2 -0.4 -0.3 -0.4 -0.1 GLY -1.1 -0.1 0.4 1.3 -0.3 -0.2 0.3 HIS -0.2 -0.2 -0.2 -1.2 -0.7 0.1 -0.4 -0.9 -0.6 - 2 нон --1.6 -1.3 -1.2 2.0 4.2 -1.4 4.4 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 -0.4 -0.3 -0.5 LYS --0.7 -0.2 -0.7 -0.1 -0.3 -0.5 -0.7 -0.1 MET --0.2 -0.2 -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 -0.4 -0.5 -0.7 SER -2.0 2.7 1.3 0.2 0.7 SO4 --1.6 -1.2 -1.2 -0.7 -1.4 -1.3 -0.5 -0.9 -1.1 THR --0.7 -0.2 -0.1 -0.3 0.1 -0.4 -0.4 TRP --0.7 -1.3 0.3 0.4 -0.4 -0.9 -0.4 TYR -1.1 2.8 2.9 0.2 2.3 2.1 2.0 1.6 VAL --0.7 -0.2 -0.2 0.4 0.2 -0.4 -0.2 -0.2 5781 Seil StiA PDB ID