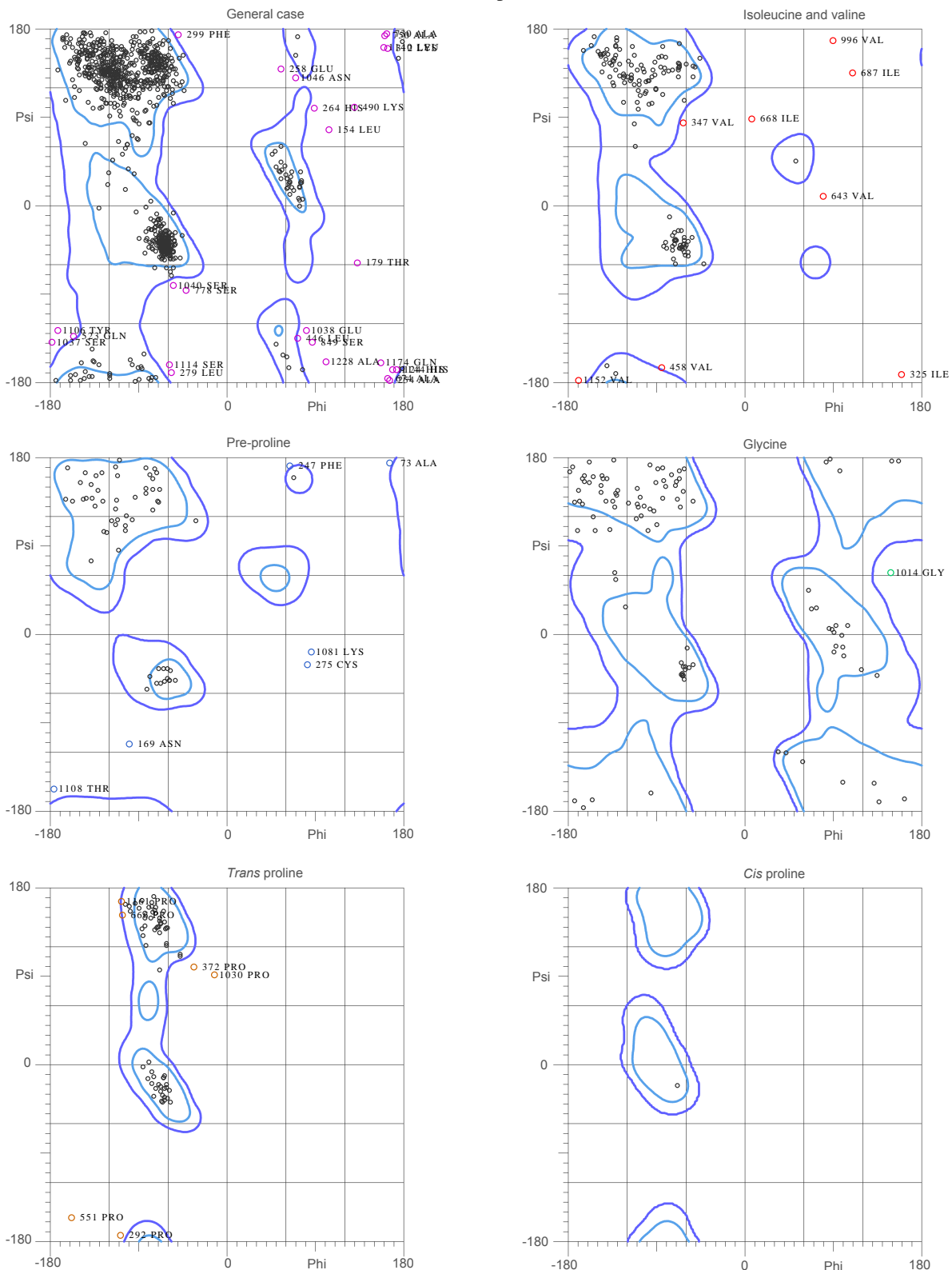


MolProbity Ramachandran analysis

6abu.B99990021.pdb, model 1



90.6% (172/192) of all residues were in favored (90%) regions.
96.7% (124/128) of all residues were in allowed (96.7%) regions.

There were 48 outliers (phi, psi):

71 ALA (166.9, 175.9)	666 PRO (-107.6, 155.2)
134 LEU (104.7, 79.1)	668 SER (-2.1, 99.9)
100 ASN (106.6, 112.6)	678 ALA (165.9, -174.2)
170 THR (111.9, -78.8)	687 SER (110.5, 176.8)
107 THR (161.4, 77.9)	778 SER (-42.1, -86.5)
104 ALA (166.8, -178.1)	849 SER (87.3, -139.8)
106 GLU (156.4, 169.9)	896 VAL (106.4, 165.5)
104 SER (161.4, 169.9)	1014 GLY (119.8, 83.9)
104 THR (161.4, 169.9)	1015 PRO (-13.1, -91.1)
104 LEU (161.4, 169.9)	1017 SER (179.7, 139.3)
104 VAL (161.4, 169.9)	1018 GLY (181.1, -177.2)
104 ILE (161.4, 169.9)	1019 SER (-76.8, -81.7)
104 MET (161.4, 169.9)	1044 ASN (79.4, 131.1)
104 ASP (161.4, 169.9)	1081 LYS (16.4, -18.4)
104 GLN (161.4, 169.9)	1088 THR (-177.3, -159.3)
104 PRO (161.4, 169.9)	1140 THR (-175.1, -127.9)
104 THR (161.4, 169.9)	1141 SER (-150.4, -142.8)
104 VAL (161.4, 169.9)	1142 ILE (176.8, -147.5)
104 MET (161.4, 169.9)	1143 LEU (165.2, -142.4)
104 ASP (161.4, 169.9)	1144 MET (177.4, -147.5)
104 GLN (161.4, 169.9)	1152 VAL (-179.2, -179.9)
104 PRO (161.4, 169.9)	1161 PRO (108.4, 167.9)
104 THR (161.4, 169.9)	1174 GLN (187.4, -169.7)
104 VAL (161.4, 169.9)	1228 ALA (101.8, -159.8)