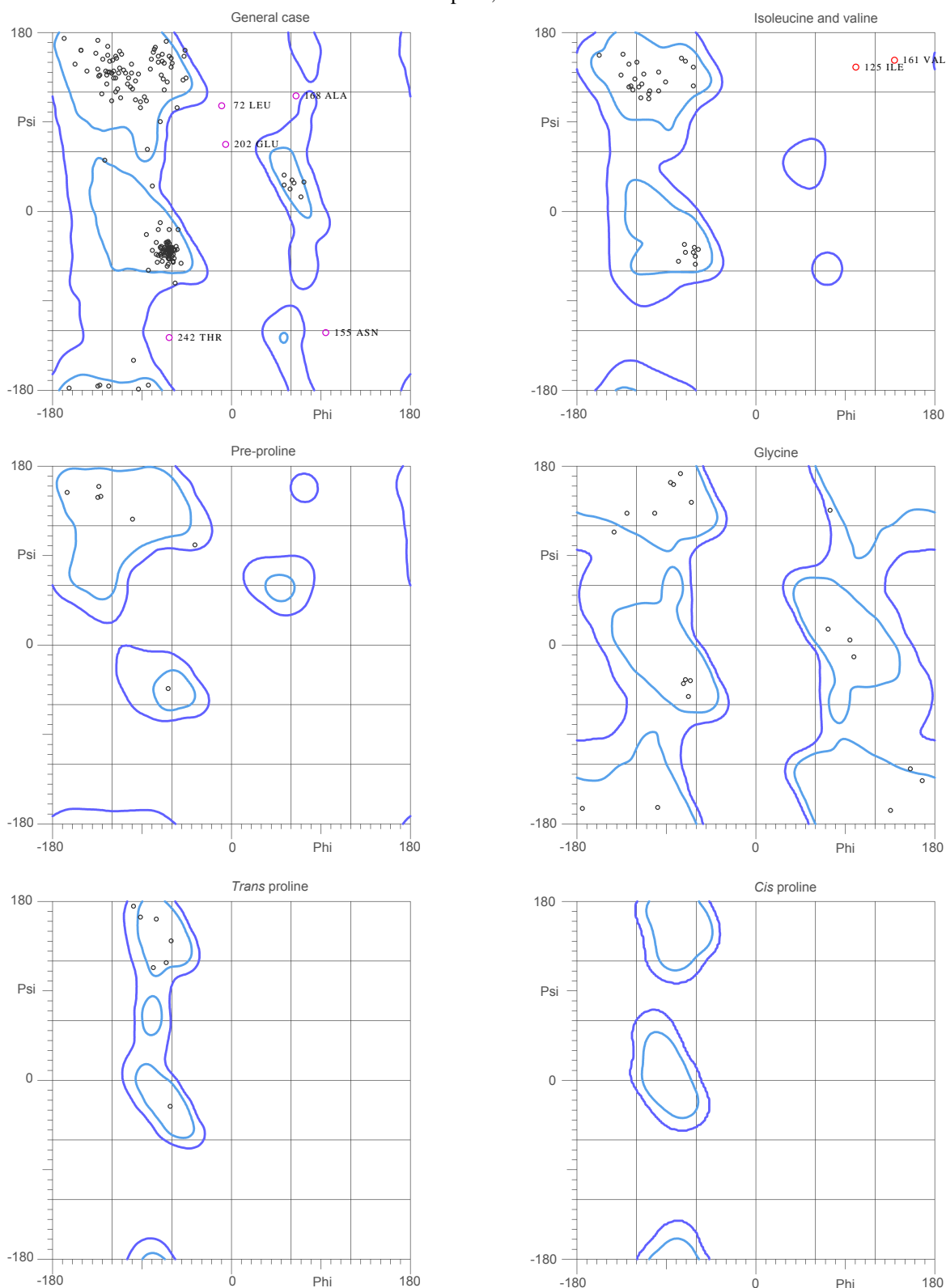


MolProbity Ramachandran analysis

CD168.pdb, model 1



91.4% (224/245) of all residues were in favored (98%) regions.
97.1% (238/245) of all residues were in allowed (>99.8%) regions.

There were 7 outliers (phi, psi):

72 LEU (-10.3, 107.4)
125 ILE (101.5, 146.3)
155 ASN (95.6, -122.0)
161 VAL (140.7, 153.7)
168 ALA (65.2, 117.7)
202 GLU (-6.5, 68.9)
242 THR (-63.4, -127.4)