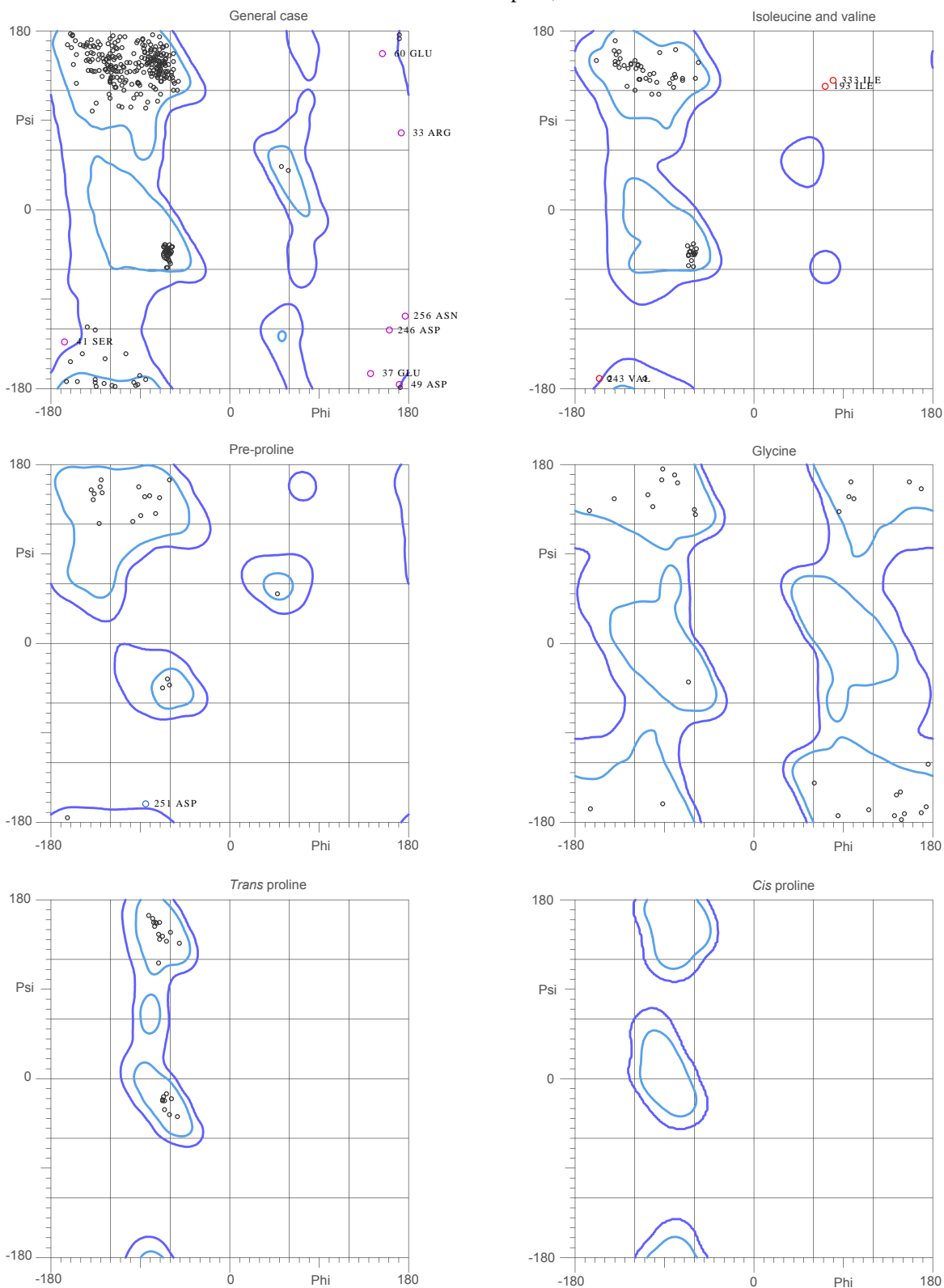# MolProbity Ramachandran analysis

1ad3A.B99990048.pdb, model 1

94.4% (419/444) of all residues were in favored (98%) regions.
97.5% (433/444) of all residues were in allowed (>99.8%) regions.

There were 11 outliers (phi, psi):
    33 ARG (173.5, 78.8)
    37 GLU (142.4, -165.7)
    41 SER (-167.7, -133.0)
    49 ASP (171.5, -176.8)
    60 GLU (154.1, 158.4)
    193 ILE (72.5, 125.7)
    243 VAL (-156.2, -170.8)
    246 ASP (161.7, -121.2)
    251 ASP (-85.7, -162.5)
    256 ASN (177.9, -107.9)
    333 ILE (80.1, 131.6)