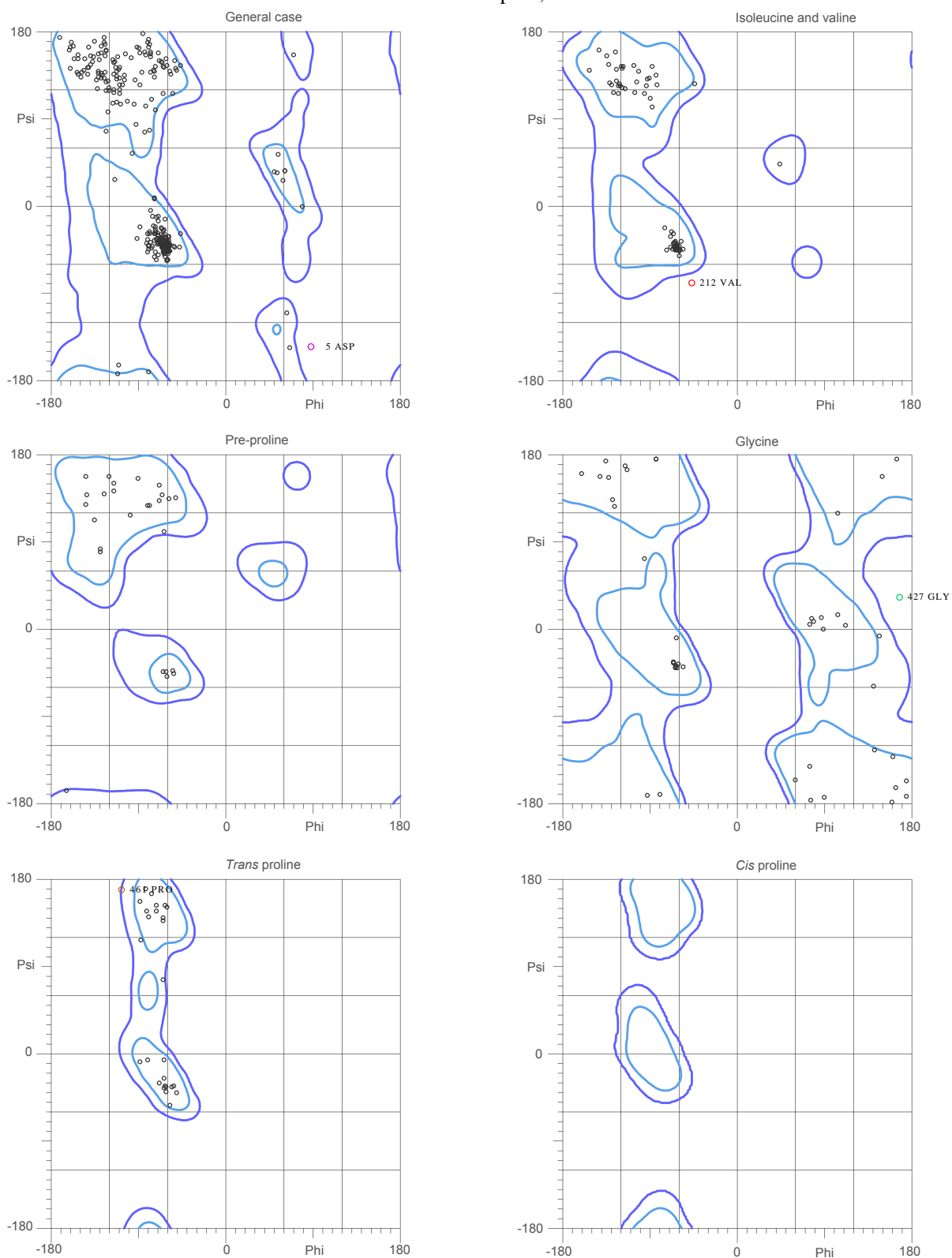# MolProbity Ramachandran analysis

1a4sA.B99990002.pdb, model 1

96,0% (481/501) of all residues were in favored (98%) regions.
99,2% (497/501) of all residues were in allowed (>99.8%) regions.

There were 4 outliers (phi, psi):
    5 ASP (88,3, -145,2)
    212 VAL (-47,6, -79,5)
    427 GLY (168,2, 33,9)
    461 PRO (-108,0, 170,7)

Lovell, Davis, et al. Proteins 50:437 (2003)