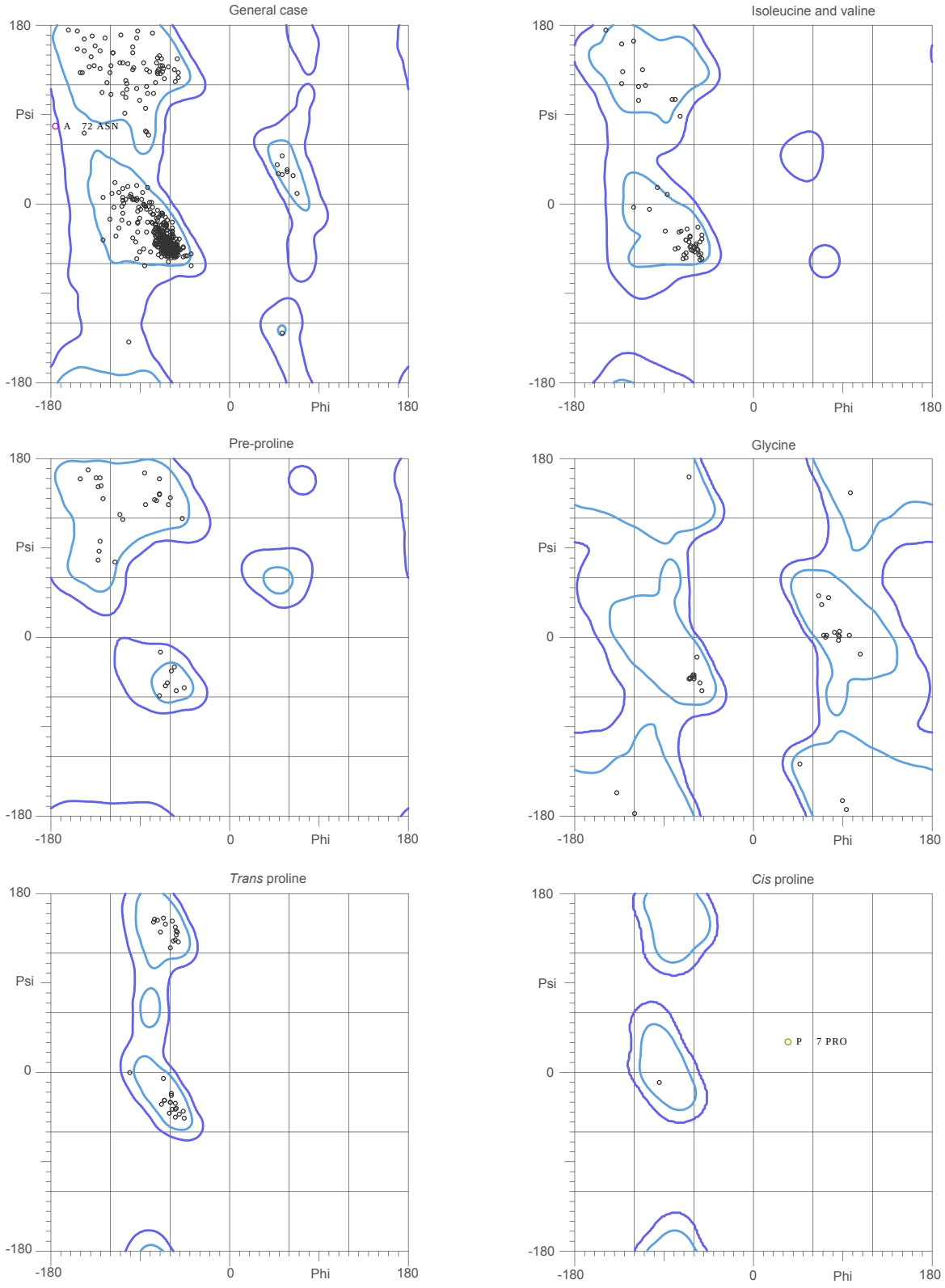
MolProbity Ramachandran analysis

4aph.H.pdb, model 1



98.1% (606/618) of all residues were in favored (98%) regions.
99.7% (616/618) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (phi, psi):

A 72 ASN (-176.8, 79.3)
P 7 PRO (35.3, 31.6)

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