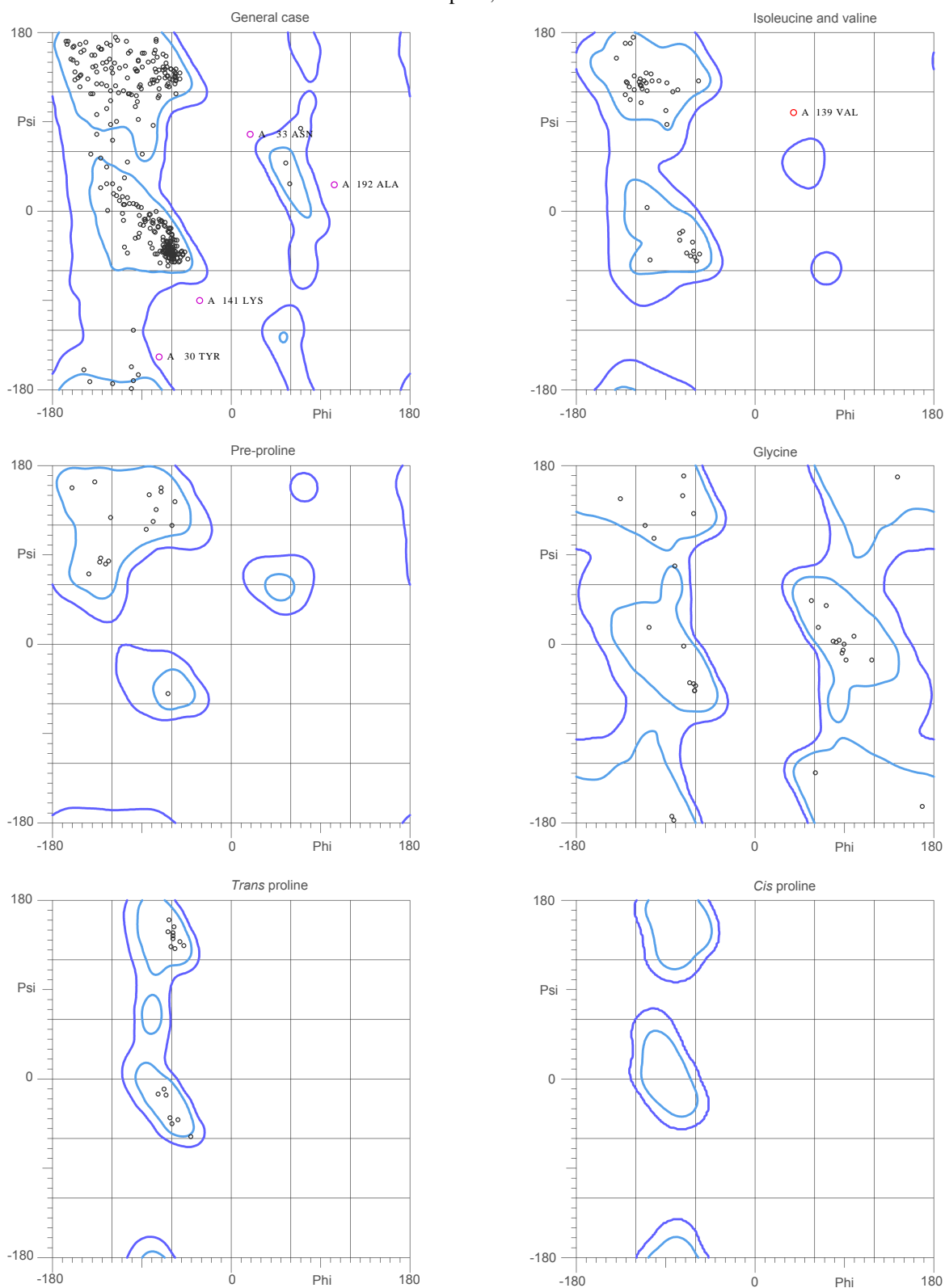


# MolProbity Ramachandran analysis

3r24.H.pdb, model 1



95.8% (392/409) of all residues were in favored (98%) regions.  
98.8% (404/409) of all residues were in allowed (>99.8%) regions.

There were 5 outliers (phi, psi):

- A 30 TYR (-73.5, -147.2)
- A 33 ASN (19.8, 78.1)
- A 139 VAL (39.8, 100.9)
- A 141 LYS (-32.6, -90.7)
- A 192 ALA (104.1, 27.6)