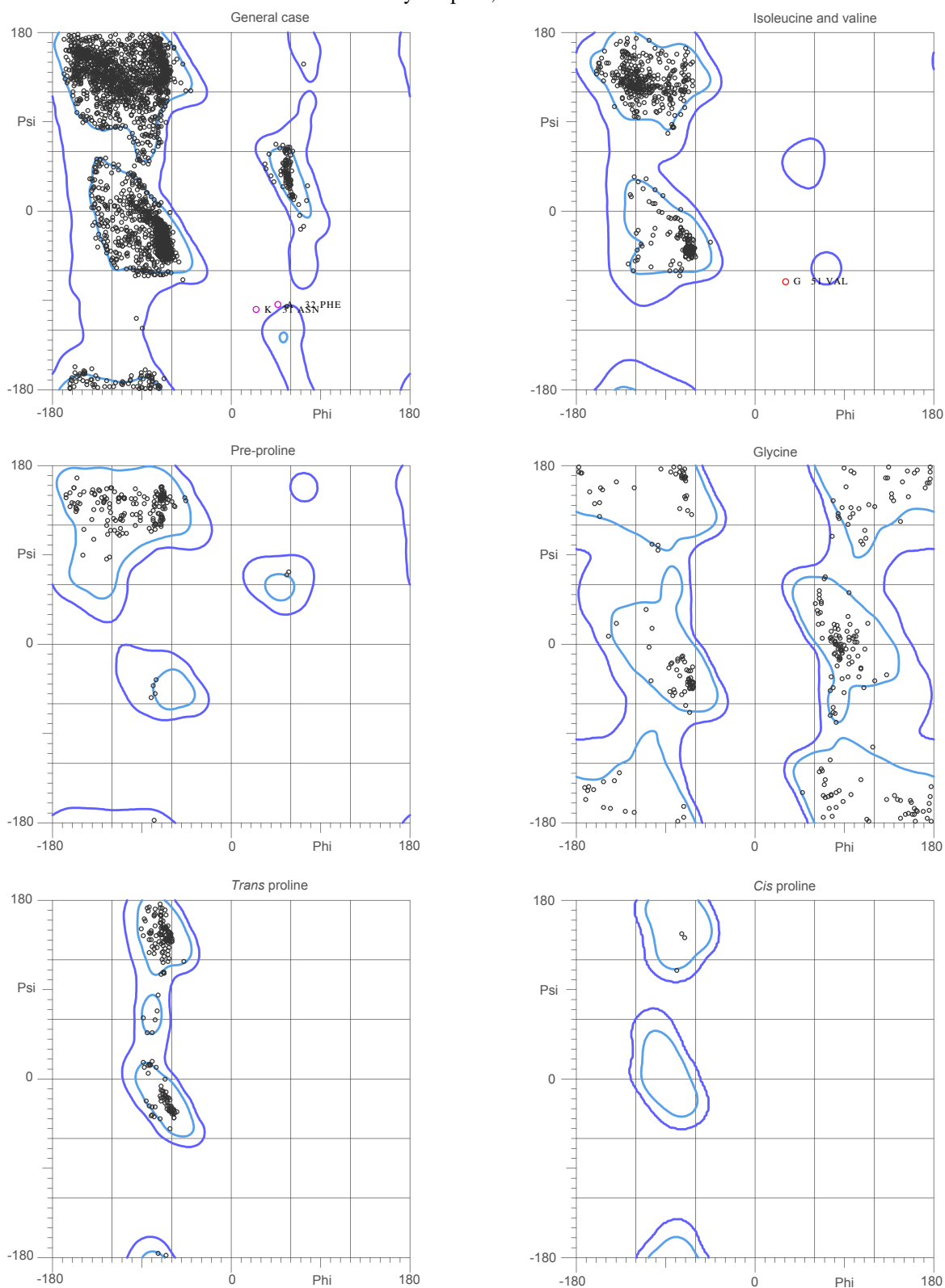


# MolProbity Ramachandran analysis

6xy.H.pdb, model 1



94.8% (3541/3734) of all residues were in favored (98%) regions.  
99.9% (3731/3734) of all residues were in allowed (>99.8%) regions.

There were 3 outliers (phi, psi):

- A 32 PHE (47.7, -94.6)
- G 51 VAL (31.1, -71.7)
- K 31 ASN (25.5, -99.1)