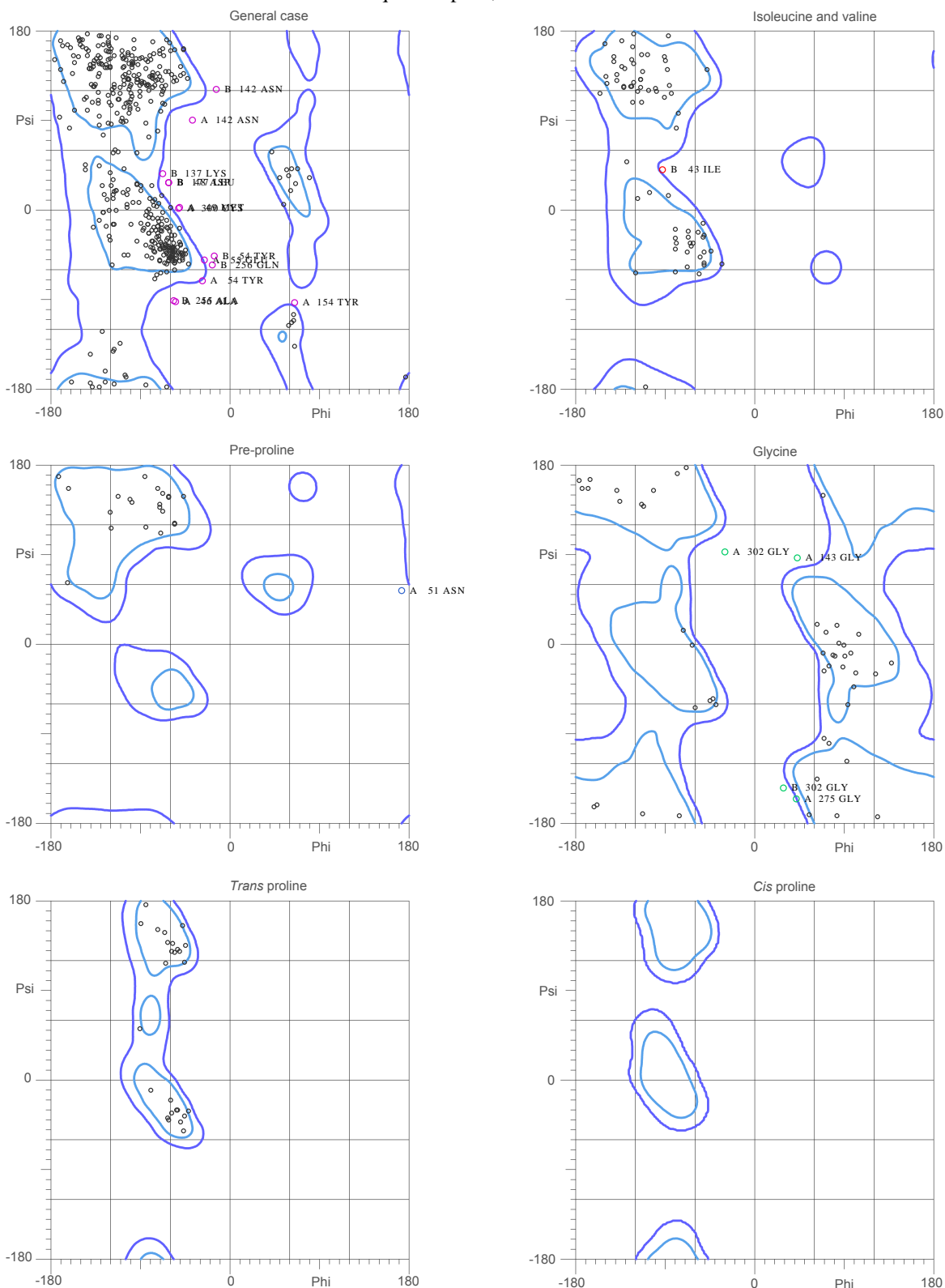


# MolProbity Ramachandran analysis

2qc2.H.pdb, model 1



84.7% (511/600) of all residues were in favored (98%) regions.  
96.7% (586/600) of all residues were in allowed (>95.8%) regions.

There were 20 outliers (Phi, Psi):  
A 46 ALA (-55.2, -92.0)  
A 49 MET (-51.4, 3.3)  
A 51 ASN (171.0, 54.9)  
A 54 TYR (-28.6, -71.3)  
A 55 GLY (-28.6, -10.0)  
A 142 ASN (-38.2, 91.4)  
A 143 GLY (151.7, 87.3)  
A 151 TYR (105.1, -93.2)  
A 270 GLY (42.2, -150.8)  
A 300 CYS (-52.2, 2.3)  
A 302 GLY (-10.8, 93.1)  
B 43 ILE (-63.4, -61.3)  
B 48 ASP (-62.9, 28.4)  
B 54 TYR (-16.2, -46.9)  
B 177 LYS (-48.0, 37.7)  
B 142 ASN (-15.0, 122.8)  
B 177 LEU (-62.7, 28.9)  
B 255 ALA (-57.7, -91.2)  
B 256 GLN (-18.5, -59.3)  
B 302 GLY (29.6, -145.6)