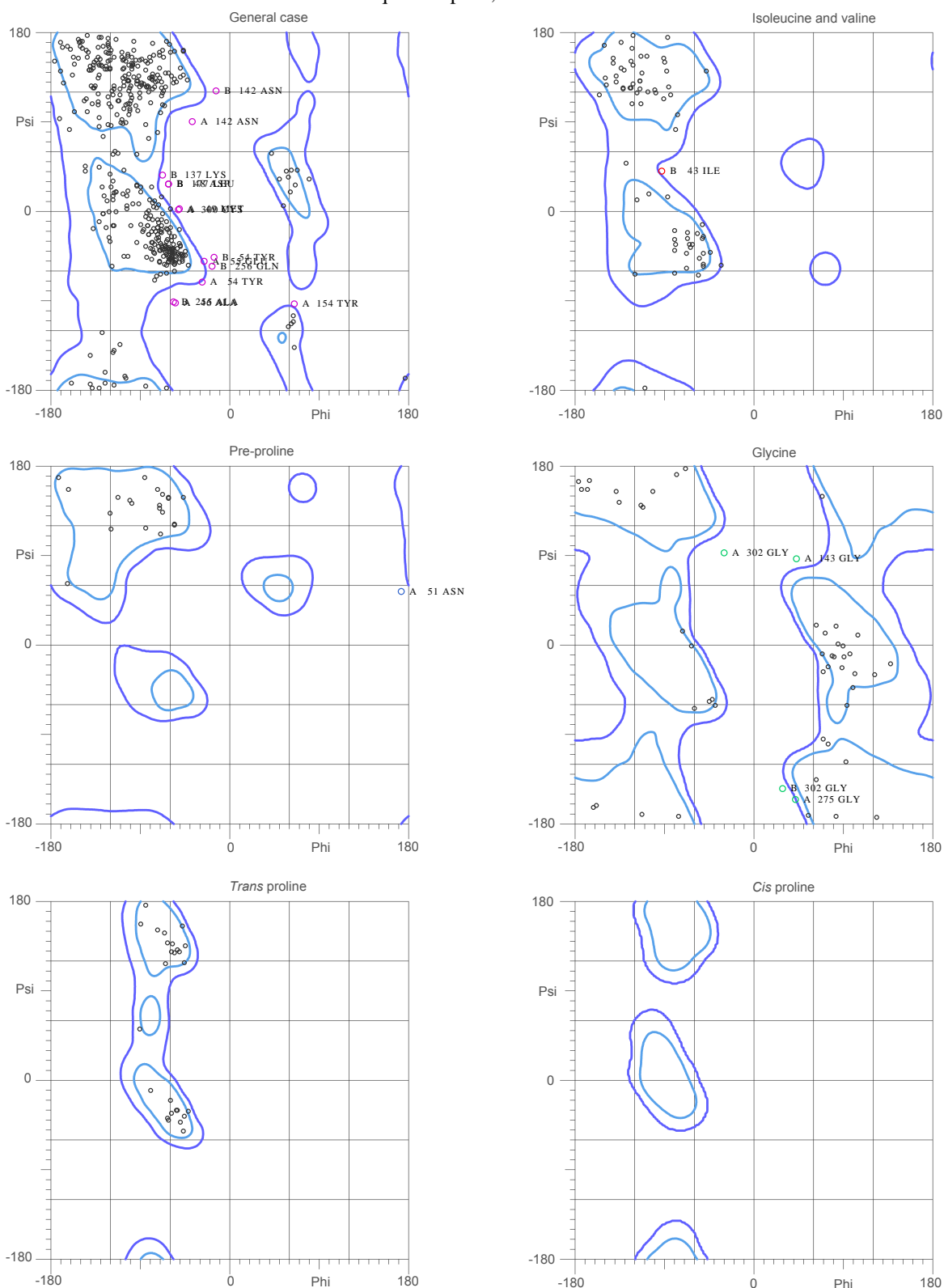


MolProbity Ramachandran analysis

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 (173.0, 54.9)
- A 54 TYR (-28.6, -71.3)
- A 55 GLY (-28.6, -55.0)
- A 142 ASN (-38.2, 91.4)
- A 143 GLY (81.3, 87.3)
- A 151 TYR (85.1, -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 (-68.0, 37.7)
- B 142 ASN (-15.0, 122.8)
- B 177 LEU (-62.3, 28.9)
- B 255 ALA (-57.7, -91.2)
- B 256 GLN (-18.5, -59.3)
- B 302 GLY (29.0, -145.6)