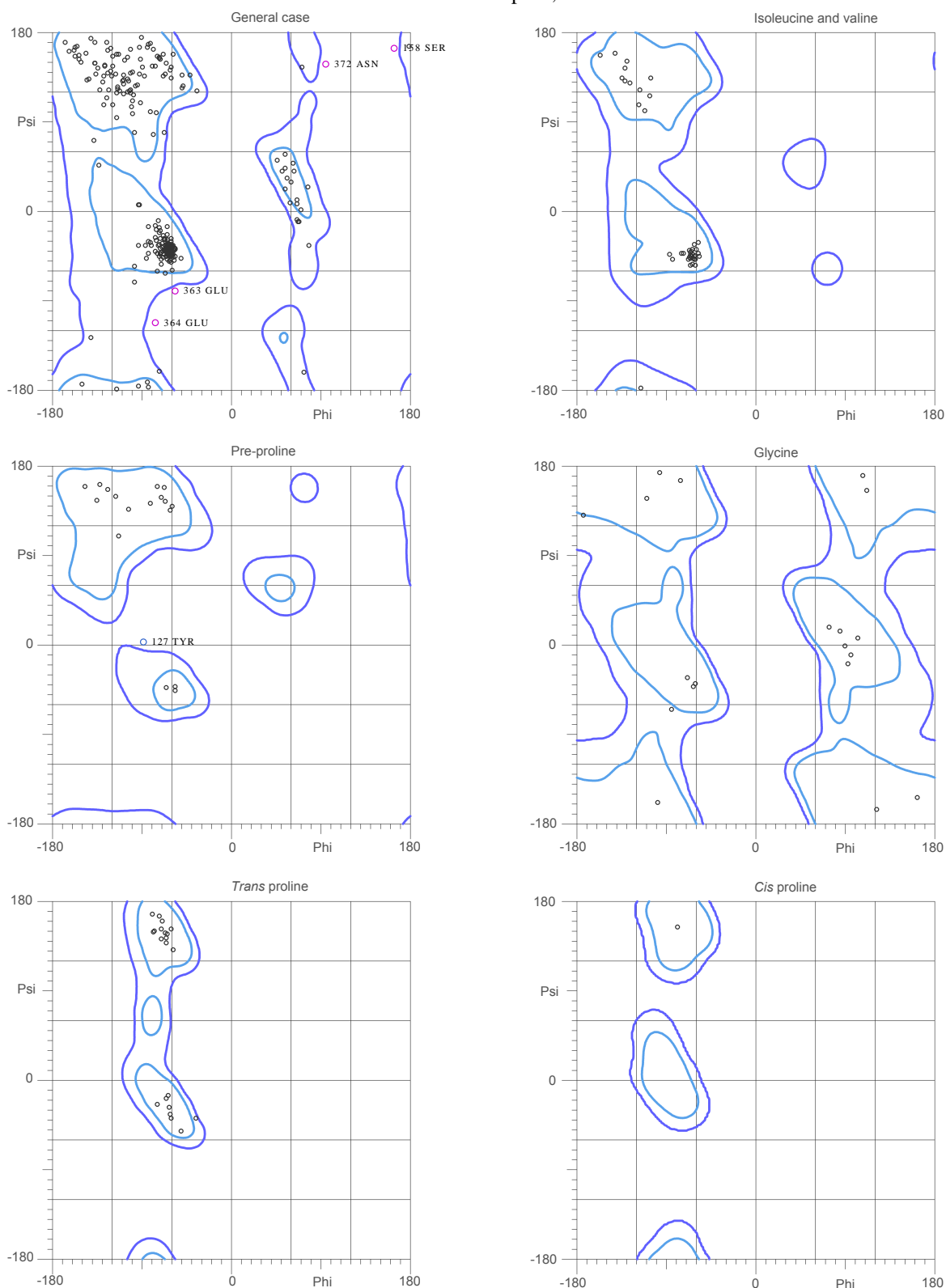


MolProbity Ramachandran analysis

1auiA.B99990035.pdb, model 1



93.4% (351/376) of all residues were in favored (98%) regions.
98.7% (371/376) of all residues were in allowed (>99.8%) regions.

There were 5 outliers (phi, psi):

127 TYR (-89.0, 3.1)
158 SER (164.6, 166.0)
363 GLU (-57.6, -80.0)
364 GLU (-77.1, -112.1)
372 ASN (95.8, 149.8)