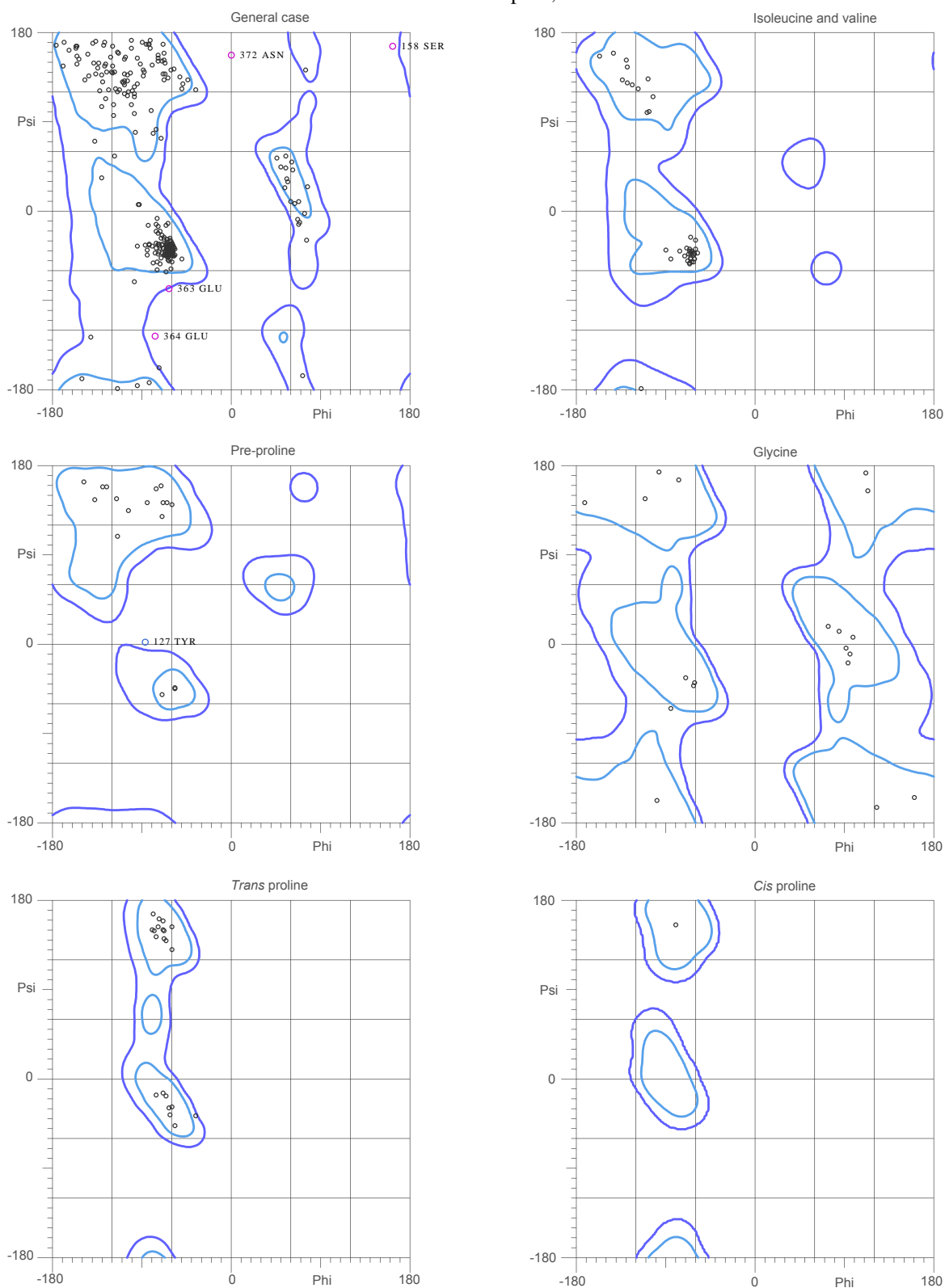


# MolProbity Ramachandran analysis

1auiA.B99990028.pdb, model 1



94,1% (354/376) 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 (-87,3, 2,6)  
158 SER (164,0, 167,8)  
363 GLU (-63,3, -78,5)  
364 GLU (-78,0, -126,4)  
372 ASN (-0,9, 158,6)