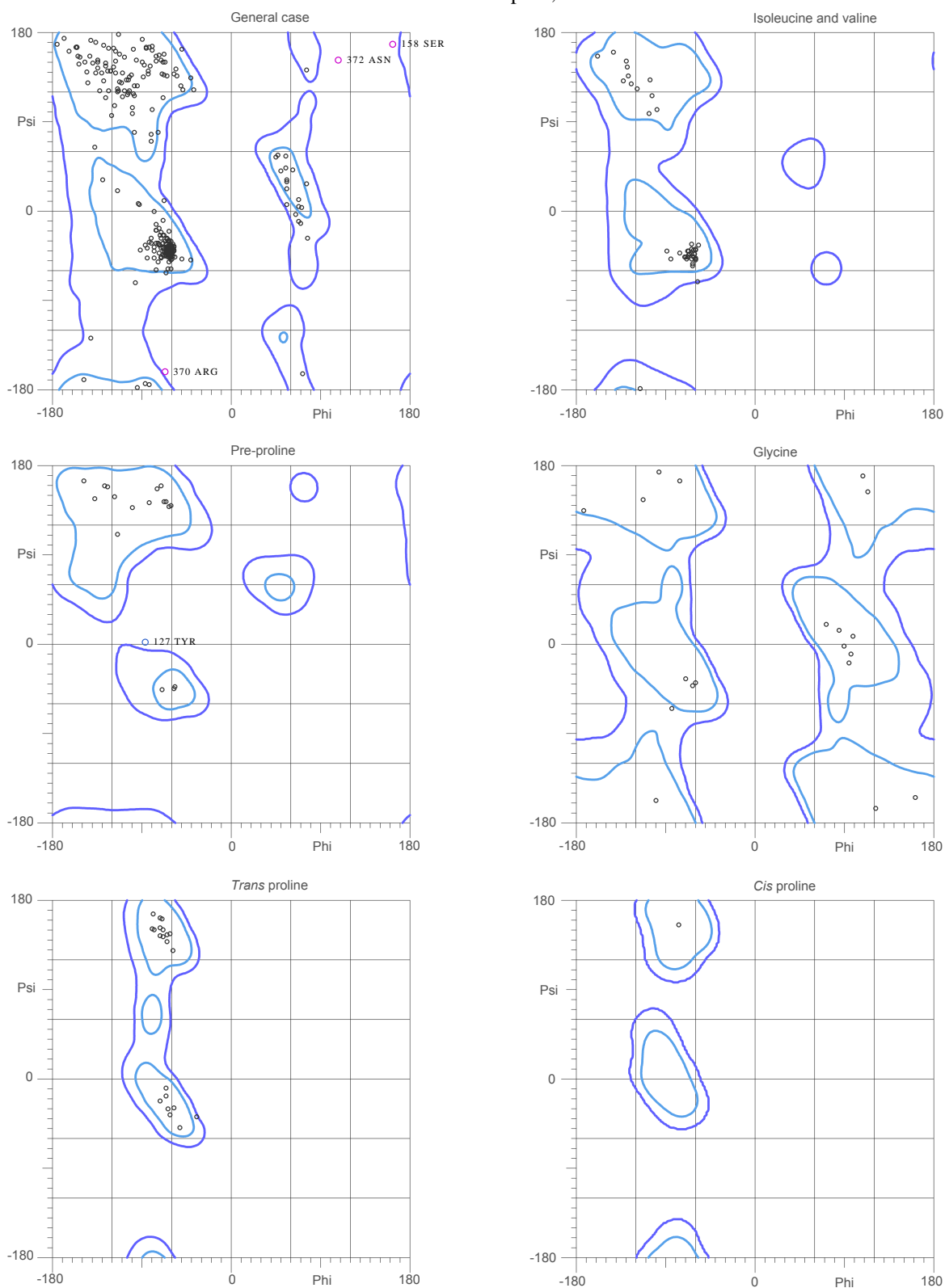


# MolProbity Ramachandran analysis

1auiA.B99990049.pdb, model 1



93,6% (352/376) of all residues were in favored (98%) regions.  
98,9% (372/376) of all residues were in allowed (>99.8%) regions.

There were 4 outliers (phi, psi):

127 TYR (-87,6, 2,8)  
158 SER (163,7, 169,0)  
370 ARG (-67,9, -162,7)  
372 ASN (108,5, 153,7)