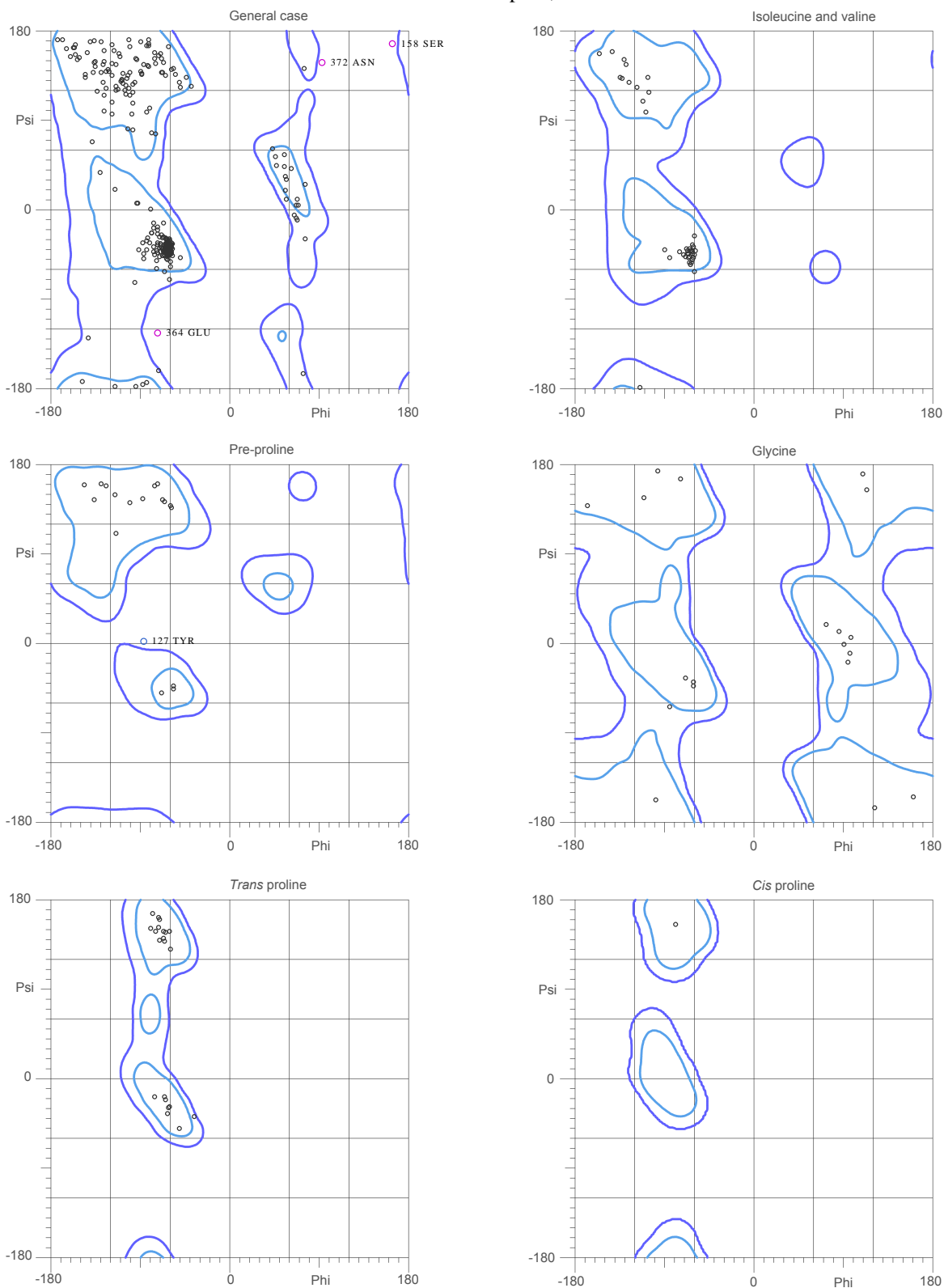


MolProbity Ramachandran analysis

1auiA.B99990017.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 (-87,4, 2,7)
158 SER (164,2, 168,3)
364 GLU (-73,9, -124,3)
372 ASN (93,1, 150,0)