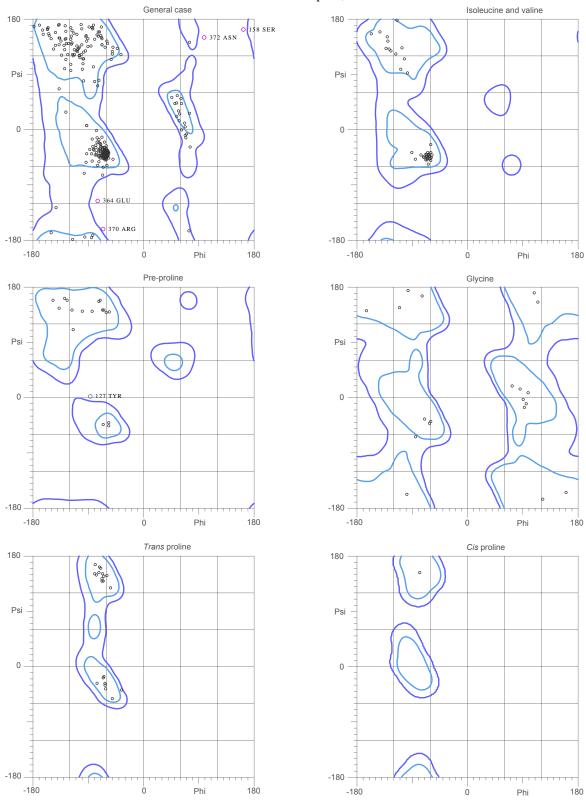
## MolProbity Ramachandran analysis

1auiA.B99990047.pdb, model 1



93,6% (352/376) 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 (-87,4, 2,1)
158 SER (163,7, 164,1)
364 GLU (-76,0, -116,3)
370 ARG (-66,9, -162,9)
372 ASN (99,3, 151,2)