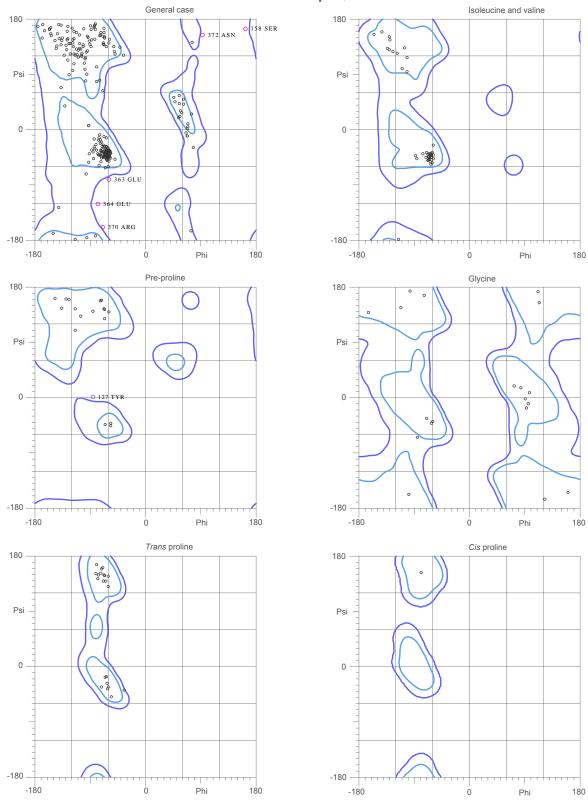
MolProbity Ramachandran analysis

1auiA.B99990019.pdb, model 1



94,1% (354/376) of all residues were in favored (98%) regions. 98,4% (370/376) of all residues were in allowed (>99.8%) regions.

There were 6 outliers (phi, psi):
127 TYR (-86,8, 1,7)
158 SER (163,5, 165,9)
363 GLU (-60,3, -82,0)
364 GLU (-79,0, -121,8)
370 ARG (-70,3, -159,7)
372 ASN (93,9, 155,1)