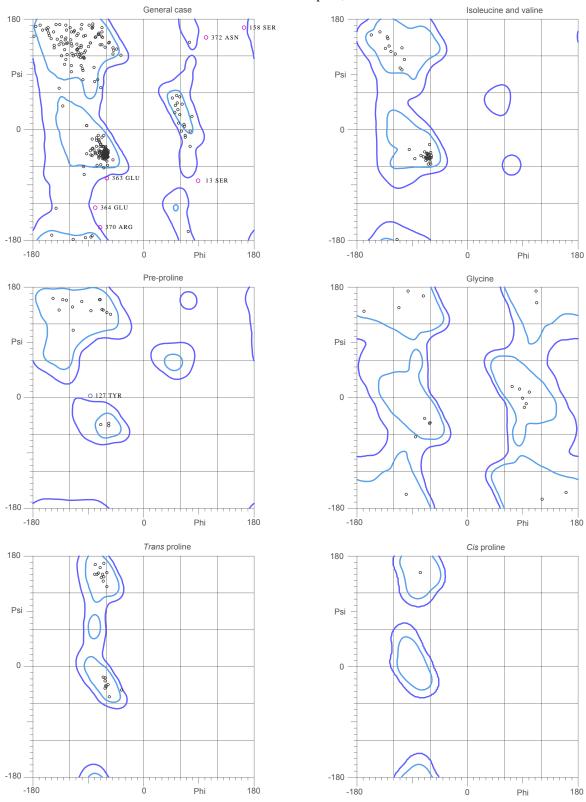
## MolProbity Ramachandran analysis

1auiA.B99990041.pdb, model 1



98,1% (369/376) of all residues were in allowed (>99.8%) regions.

There were 7 outliers (phi, psi): 13 SER (89,1, -83,8) 127 TYR (-87,3, 3,4) 158 SER (164,5, 167,0) 363 GLU (-60,6, -79,5) 364 GLU (-79,1, -127,6) 370 ARG (-71,3, -159,2) 372 ASN (102,1, 151,9)