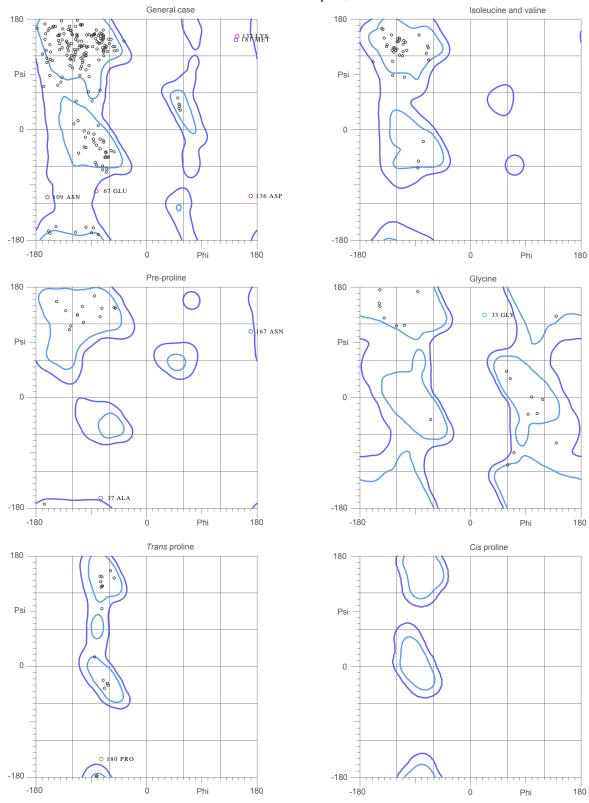
## MolProbity Ramachandran analysis

1a02N.B99990032.pdb, model 1



 $86,7\%\ (241/278)$  of all residues were in favored (98%) regions

96,8% (269/278) of all residues were in allowed (>99.8%) regions

There were 9 outliers (phi, psi): 33 GLY (23.5, 135.7) 37 ALA (-75.1, -164.8) 67 GLU (-82.7, -100.3) 109 ASN (-162.8, -110.3) 12 LYS (149.0, 153.7) 136 ASP (170.8, -108.3) 167 ASN (170.6, 108.7) 180 PRO (-75.0, -151.4) 181 MET (146.6, 147.2)