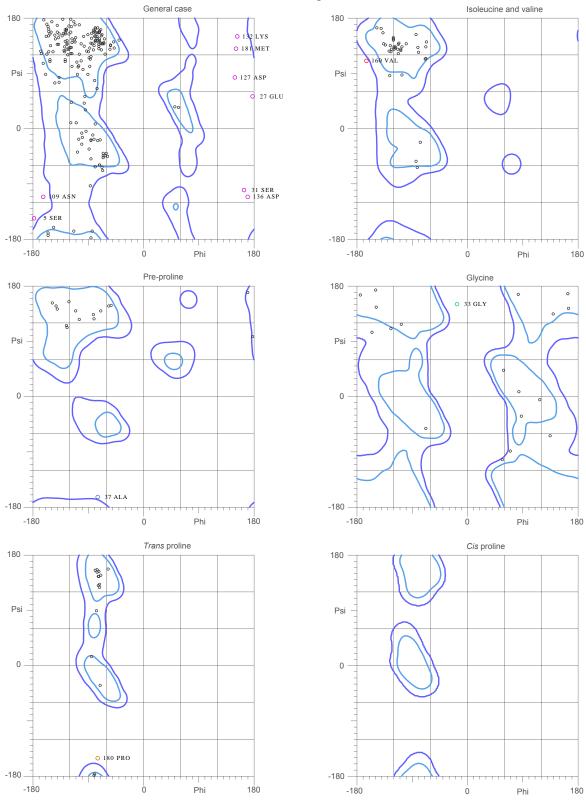
## MolProbity Ramachandran analysis

1a02N.B99990015.pdb, model 1



85,6% (238/278) of all residues were in favored (98%) regions. 95,7% (266/278) of all residues were in allowed (>99.8%) regions

7.9 (200.78) of all residues we see were 12 outliers (phi, psi): 5 SEE (-179.2, -146.7) 27 GLU (178.4, 53.8) 31 SEE (164.5, 100.8) 33 GLY (-17.2, 151.6) 37 ALA (-73.3, 164.4) 109 ASN (-164.7, -111.1) 122 LYS (153.6, 151.0) 136 ASP (170.5, 111.4) 160 VAL (-165.5, 111.5) 180 PRO (-75.6, -151.2) 181 MET (151.3, 131.2)