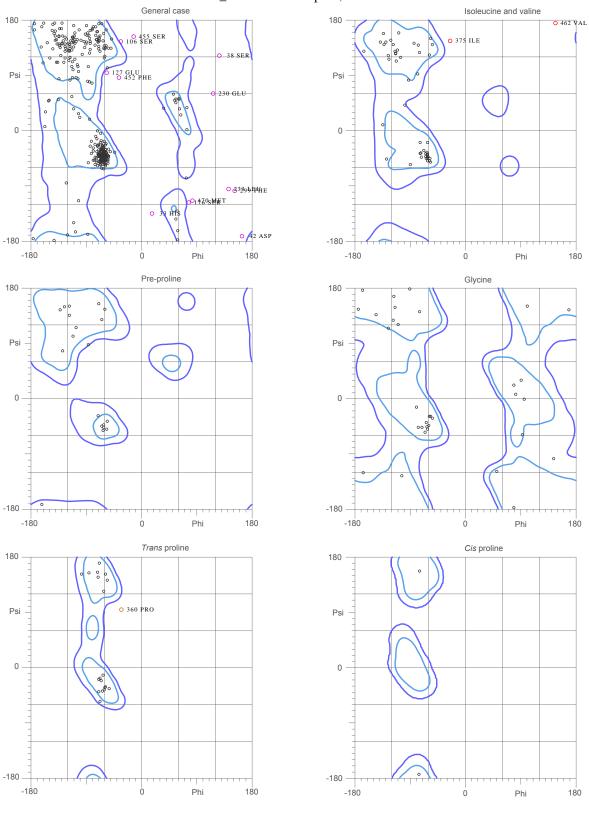
## MolProbity Ramachandran analysis

1ac5\_.B99990045.pdb, model 1



88,8% (427/481) of all residues were in favored (98%) regions. 96,9% (466/481) of all residues were in allowed (>99.8%) region

There were 15 outliers (phi, 33 HIS (17.6, -135.7) 33 KIR (127.6, -135.7) 42 ASP (164.2, -172.7) 42 ASP (164.2, -172.7) 42 ASP (164.2, -172.7) 106 SER (-34, -195.4) 127 GLIU (-57.4, -195.4) 230 GLU (117.2, -61.0) 233 LEU (142.4, -96.0) 360 PRO (-33.5, -95.6) 360 PRO (-33.5, -95.6) 375 LLE (-52.1, 147.1) 452 PHE (-37.7, -87.8)