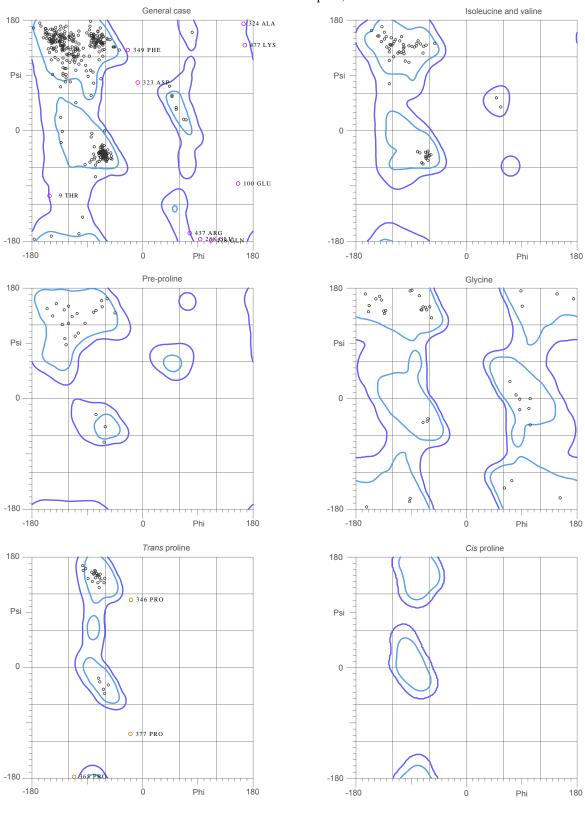
MolProbity Ramachandran analysis

1a6cA.B99990014.pdb, model 1



93,2% (476/511) of all residues were in favored (98%) regions. 97,7% (499/511) of all residues were in allowed (>99.8%) regions

7,7% (4995;11) of all residues we here were 12 outliers (phi, psi): 9 THR (-1523, -106,3) 100 GLU (156,3 -86,9) 100 GLU (156,3 -86,9) 268 GLU (94,9-107,9) 323 ASP (-8.8, 80,0) 324 ALA (652, 175,6) 346 PHG (-94,8, 132,1) 365 PHG (-12,3, -178,0) 377 PHG (-92,4-108,0) 437 ARG (77,9, -167,5) 477 LYS (167,9, 140,1) 478 GLN (112,4, -179,3)