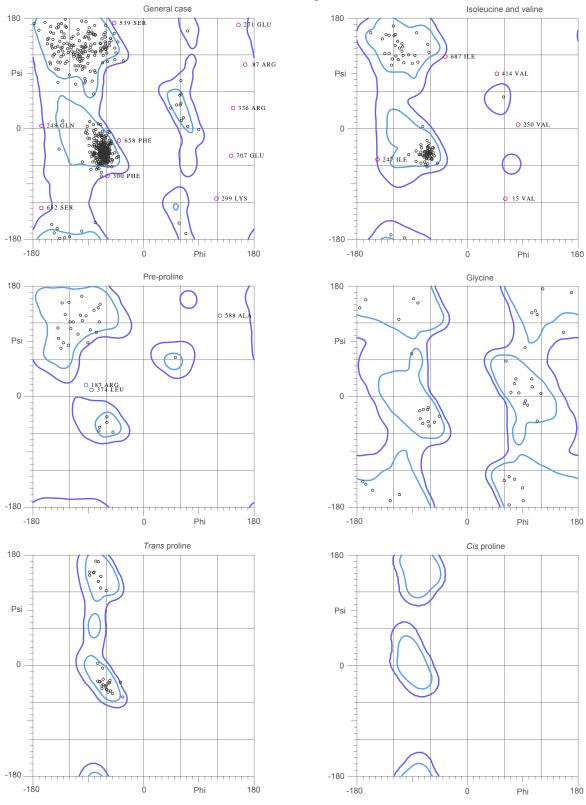
MolProbity Ramachandran analysis

1a8i_.B99990011.pdb, model 1



92,8% (752/810) of all residues were in favored (98%) regions. 97,8% (792/810) of all residues were in allowed (>99.8%) regions

here were 18 outliers (phi.; 15 VAL (e2.6, -114.2) 15 VAL (e2.6, -114.2) 87 ABG (16-6, 108.2) 187 ABG (-94.2, 19.7) 187 ABG (-94.2, 19.7) 247 ILE (-147.8, -0.6) 248 GLN (-168.0, 6.0) 250 VAL (83, 7.4) 271 GLU (155.1, 170.1) 270 GLU (155.1, 170.1) 300 PHE (-60.0, -77.9) 310 ABG (146.9, 34.2) 3174 LEU (-85.5, 11.5) 318 ABG (146.9, 34.2) 318 ALA (125.8, 10.9) 622 SER (-48.3, 173.3) 623 SER (-167.8, -129.4) 638 PHE (-46.5, 11.8)