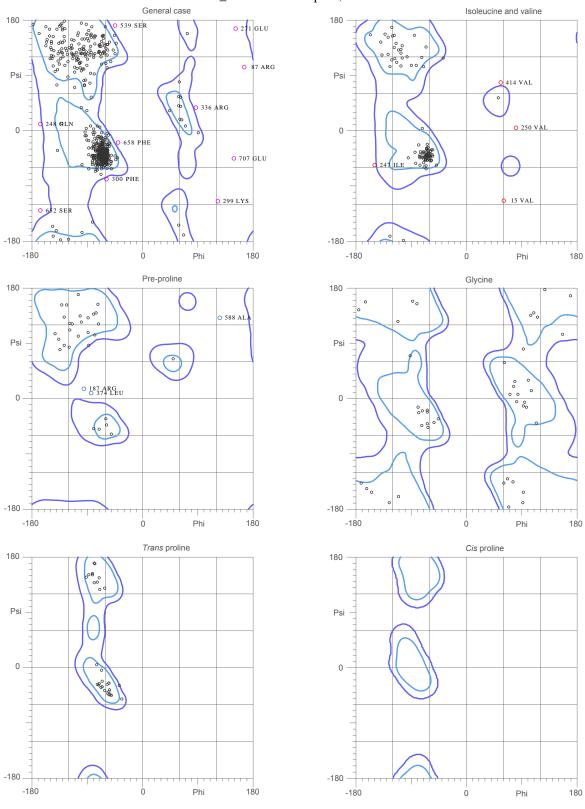
MolProbity Ramachandran analysis

1a8i_.B99990002.pdb, model 1



92,1% (746/810) of all residues were in favored (98%) regions. 97,9% (793/810) of all residues were in allowed (>99.8%) regions

here were 17 outliers (ph.; p. 15 VAL (62.0, -114.5)
15 VAL (62.0, -114.5)
77 ABG (16.6, 104.6)
187 ABG (96.3, 16.8)
187 ABG (96.3, 16.8)
247 ILE (-150.7, -56.2)
248 GIN (-167.8, 11.1)
250 VAL (81.5, 5.6)
271 GIU (152.8, 167.3)
270 JUX (123.7, -115.6)
300 PHE (-59.7, -79.9)
374 LEU (-44.3, 5.5)
374 LEU (-44.1, 56.3, 79.9)
539 SER (45.4, 79.9)
539 SER (45.4, 12.8)
588 ALA (12.6, 132.3)
588 ALA (12.6, 132.3)
588 ALA (12.6, 132.3)