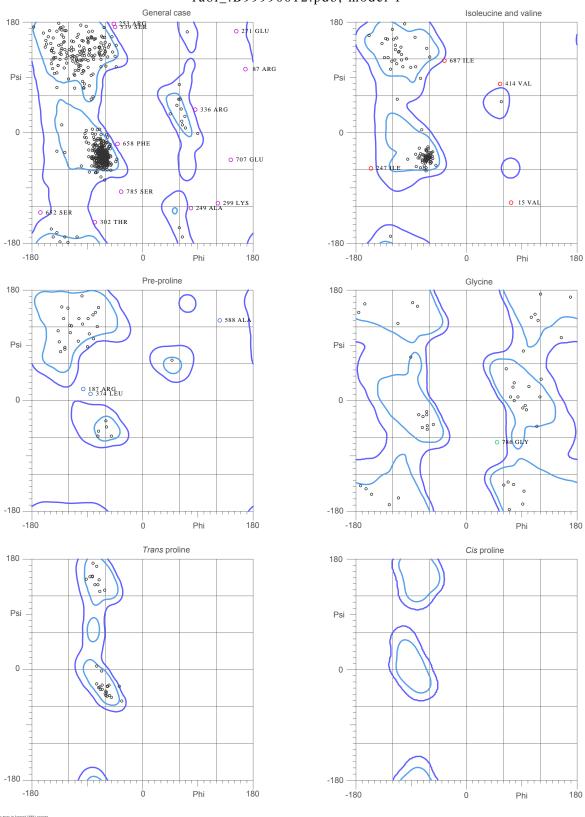
MolProbity Ramachandran analysis

1a8i_.B99990012.pdb, model 1



92.3% (748/810) of all residues were in favored (98%) regions.

There were 20 outliers (phi., 15 VAL (73.1, -114.8) 87 ARG (168.5, 104.7) 187 ARG (-97.3, 19.6) 247 ILE (-156., -58.4) 240 ALA (79.9, -122.0) 253 ARG (-47.9, 178.4) 271 GLU (153.6, 166.7) 299 LYS (123.1, -115.1)

240 ALA (79,6, -213,0) 253 ARG (-47,0, 178,4) 271 GLU (153,6, 166,7) 299 LYS (123,1, -115,1) 302 THE (-78,0, -146,3) 316 ARG (86,9, 38,0) 374 LEU (-58,4, 11,1) 414 VAL (55,1, 80,6) 579 SEE (-54,4, 173,1) 588 ALA (126,3, 121,6) 652 SEE (-56,5, -130,6) 687 PHE (-41,1,18,1) 687 LEE (-56,6, 118,5)