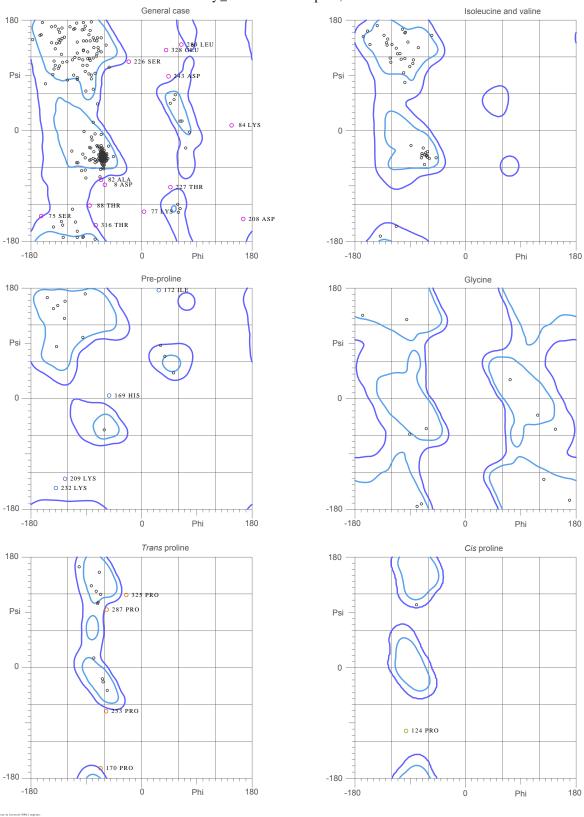
MolProbity Ramachandran analysis

1a8y_.B99990025.pdb, model 1



1,0% (279/336) of all residues were in favored (99%) regions.

93,5% (314/336) of all residue There were 22 outliers (phi, po 8 ASP (-60,4, -88,9) 75 SER (-164,1, -129,3) 77 LYS (4,9, -132,5)

75 SER (164,1, -197,1) 77 LYS (49, -132,5) 82 ALA (-64, -89,4) 84 LYS (147,9, 9,7) 82 PRO (-97,9, -104,0) 169 BIS (-53,5,5,4) 172 ILE (28,9, 177,6) 208 ASF (167,0, -144,5) 920 LYS (-125,5, -131,7,2) 216 SER (-21,9, 131,4)

172 LEC (23.9, 177.6)
208 ASP (167.0, -144.4)
209 LYS (-125.5, -131.7)
226 SER (-21.9, 131.4)
226 SER (-21.9, 131.4)
226 SER (-21.9, 131.4)
227 THE (67.2, -92.0)
232 LYS (-140.7, -146.5)
233 PEO (-58.6, -71.6)
234 PEO (-58.6, -71.6)
237 PEO (-58.0, 95.2)
237 SEP (-58.0, 95.2)
237 PEO (-58.0, 95.2)
237 PEO (-58.0, 95.2)