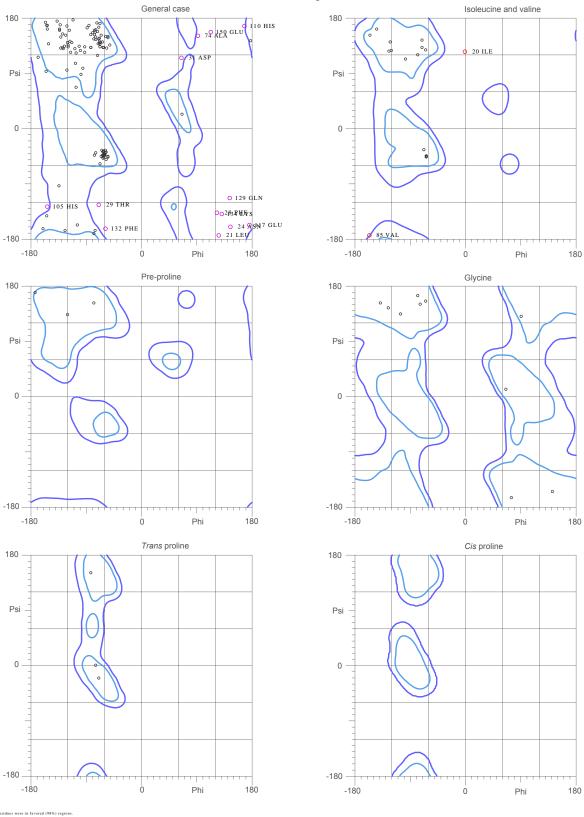
MolProbity Ramachandran analysis

1awcB.B99990012.pdb, model 1



82,8% (125/151) of all residues were in favored (98%) regions. 90,1% (136/151) of all residues were in allowed (>99.8%) region

There were 15 outliers (phi, 20 ILE (-1.0, 126.4)
20 ILE (-1.0, 126.4)
21 LEU (126.1, -174.2)
24 ASN (145.3, -160.9)
28 PHE (123.5, -137.2)
29 THE (-70.5, -124.1)
31 ASP (654, -116.5)
74 ALA (92.4, 153.0)
85 VAL (-157.5, -174.5)
100 RHS (-154.6, -127.7)
110 IHS (168.2, 168.7)
117 GLU (176.5, -157.8)
129 GLN (144.5, -113.3)
129 PHE (-59.9, -163.6)