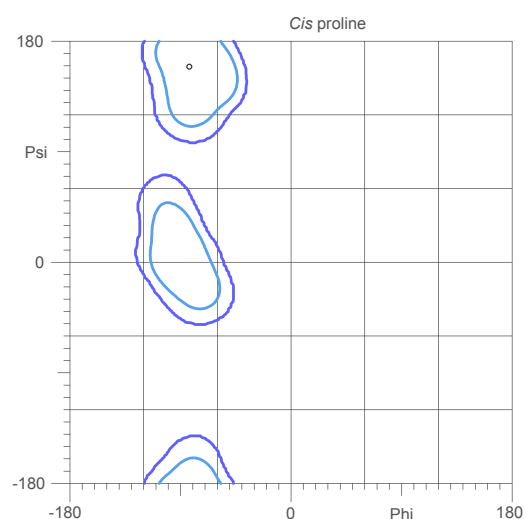
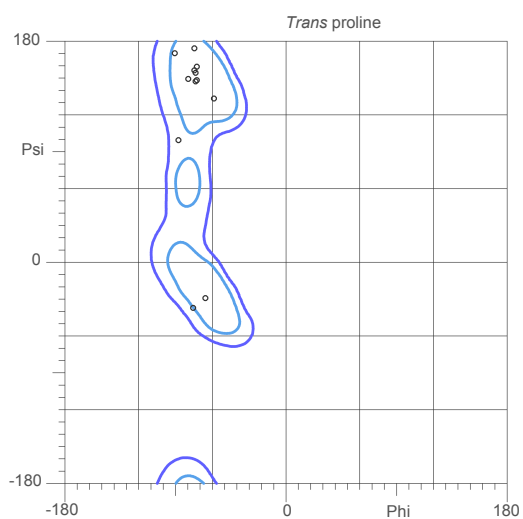
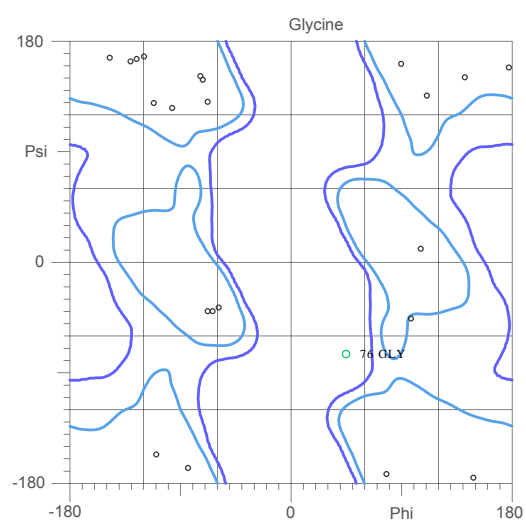
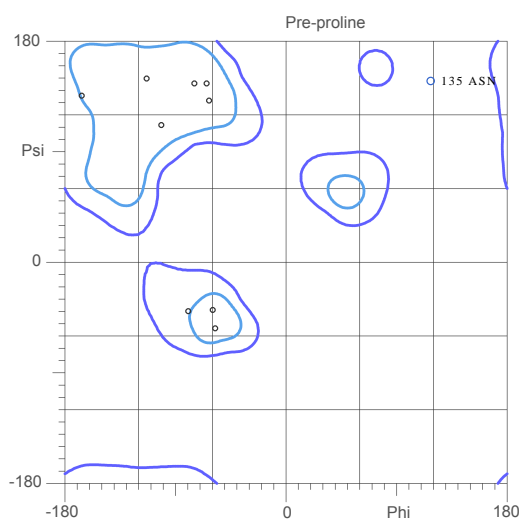
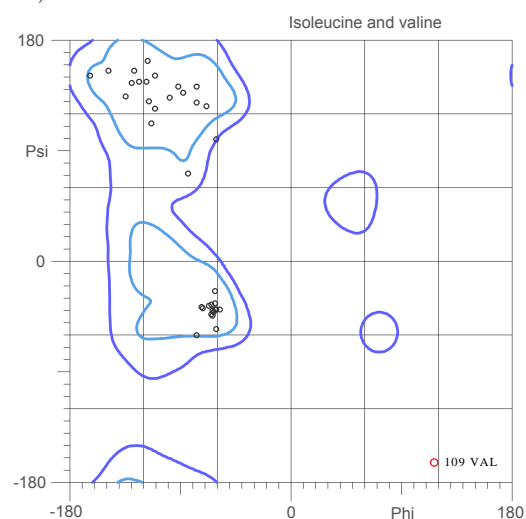
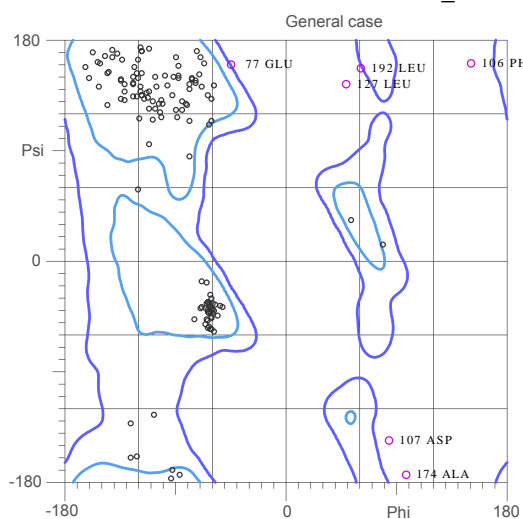


MolProbity Ramachandran analysis

1ad2_.B99990008.pdb, model 1



90.1% (200/222) of all residues were in favored (98%) regions.
95.9% (213/222) of all residues were in allowed (>99.8%) regions.

There were 9 outliers (phi, psi):

76 GLY (45.6, -75.7)
77 GLU (-45.4, 161.2)
106 PHE (151.3, 162.1)
107 ASP (84.2, -146.4)
109 VAL (117.6, -164.4)
127 LEU (49.9, 145.1)
135 ASN (118.1, 148.5)
174 ALA (98.1, -174.6)
192 LEU (61.1, 158.7)