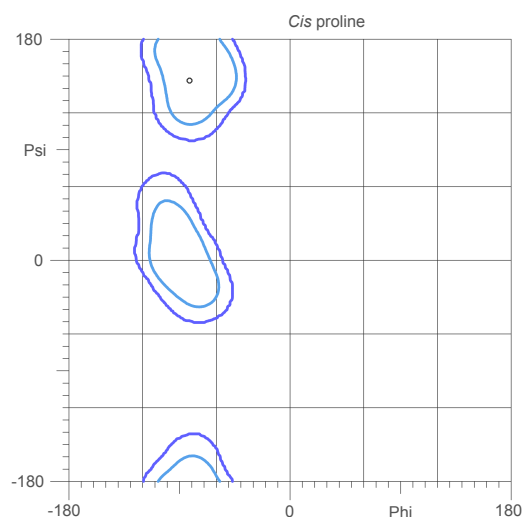
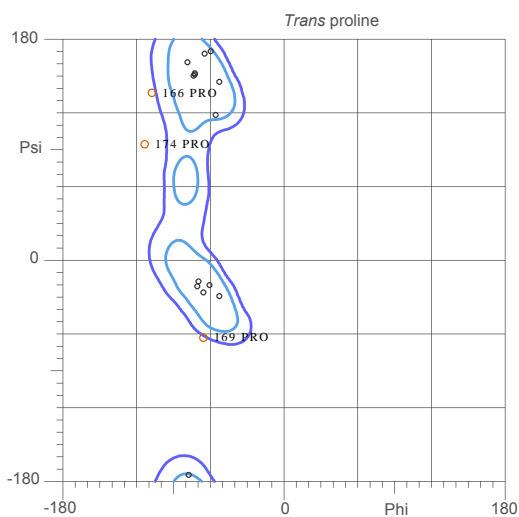
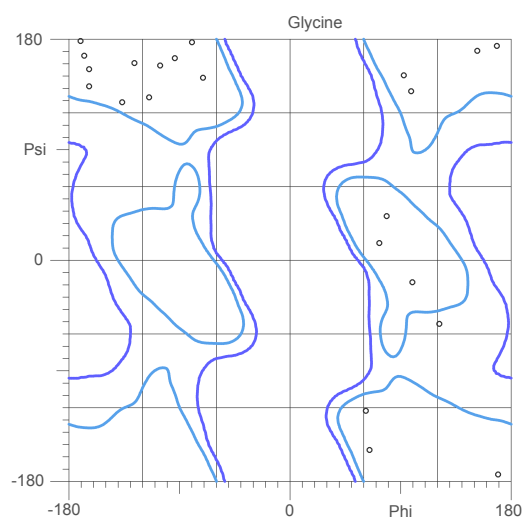
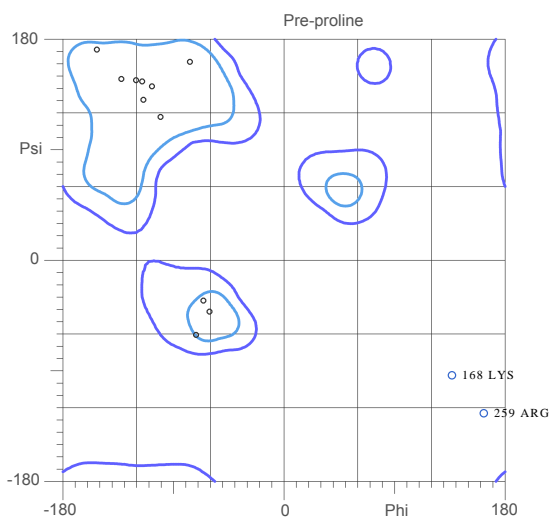
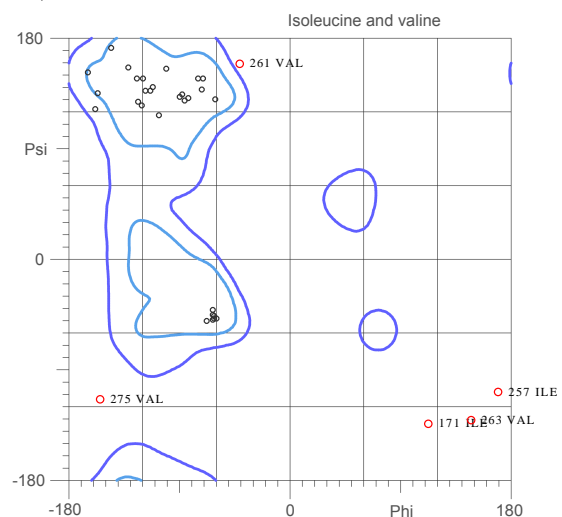
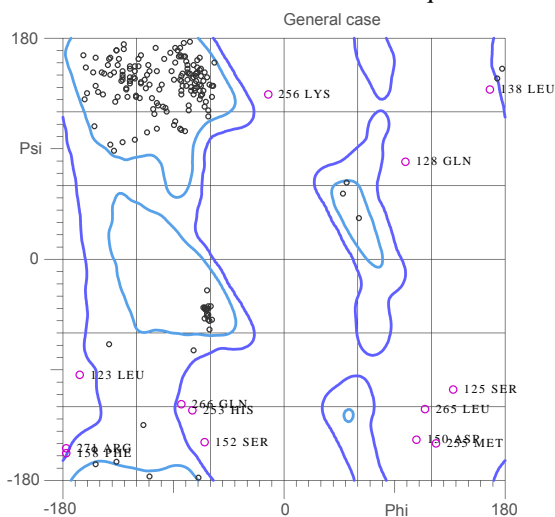


MolProbity Ramachandran analysis

1a3qA.B99990032.pdb, model 1



88.0% (249/283) of all residues were in Allowed (99%) regions.
91.0% (260/283) of all residues were in allowed (>99.0%) regions.

There were 25 outliers (outliers):

123 LEU (147.4, -94.9)
125 SER (178.8, -96.9)
126 GLN (99.1, 80.0)
128 GLN (106.5, -109.0)
138 LEU (108.7, -147.9)
152 SER (108.7, -147.9)
155 HIS (108.7, -147.9)
156 MET (178.8, -118.2)
158 ARG (108.7, -117.0)
168 LYS (117.7, -94.9)
169 PRO (108.7, -94.9)
174 PRO (117.7, -94.9)
175 ILE (111.1, -114.2)
176 ILE (111.1, -114.2)
177 ILE (111.1, -114.2)
178 ILE (111.1, -114.2)
179 ILE (111.1, -114.2)
180 ILE (111.1, -114.2)
256 LYS (111.1, -114.2)
257 ILE (111.1, -114.2)
259 ARG (108.7, -117.0)
261 VAL (108.7, -117.0)
275 VAL (108.7, -117.0)
257 ILE (111.1, -114.2)
259 ARG (108.7, -117.0)
261 VAL (108.7, -117.0)
275 VAL (108.7, -117.0)