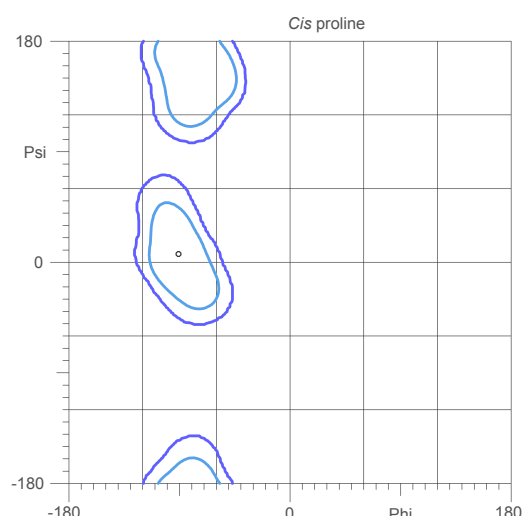
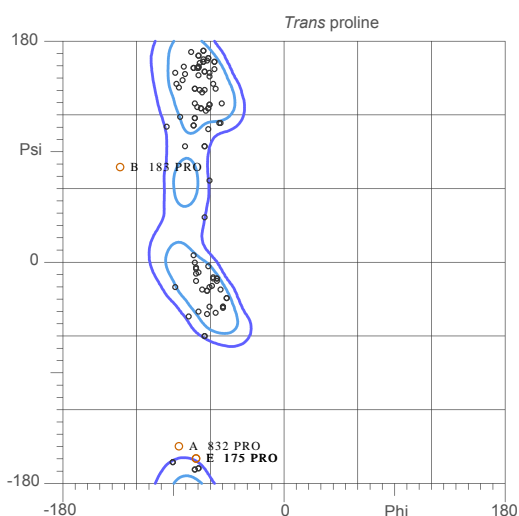
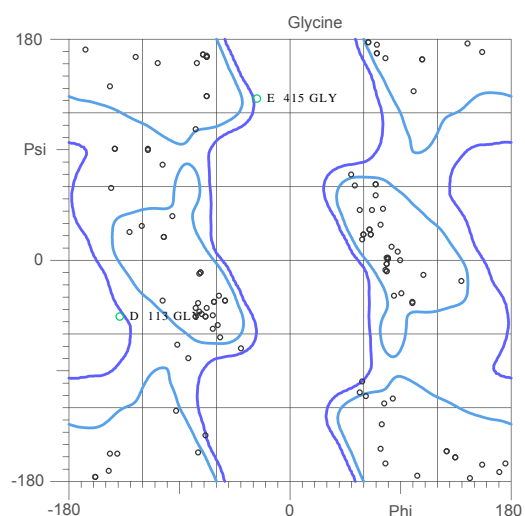
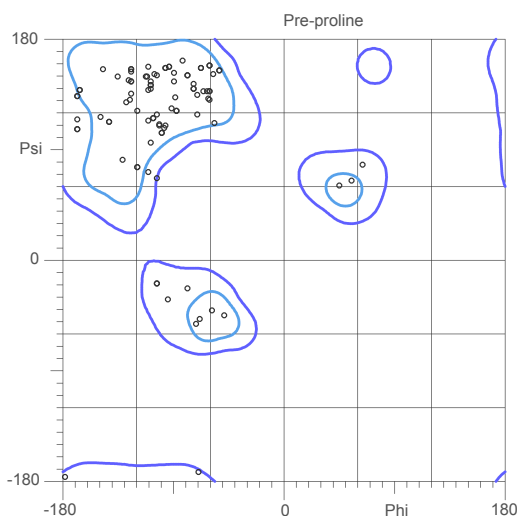
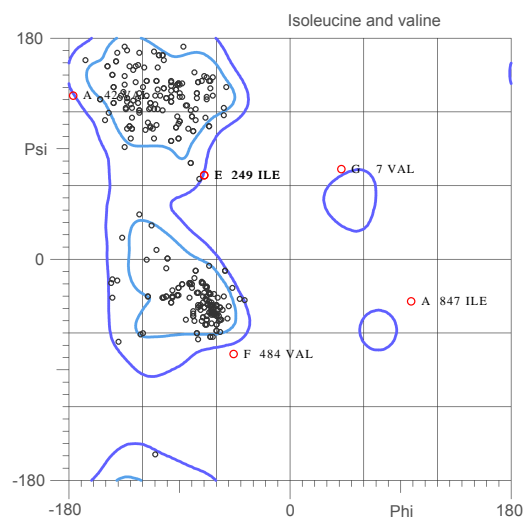
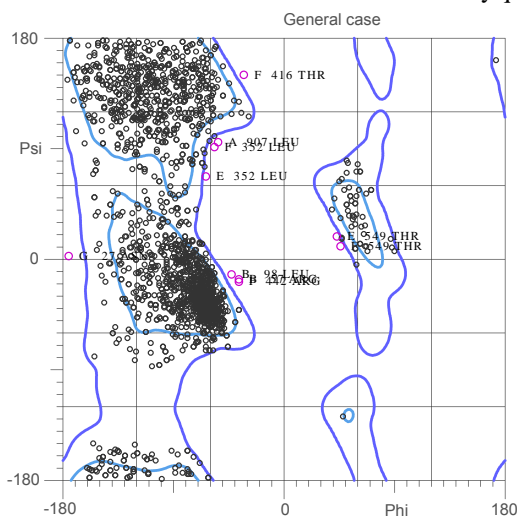


MolProbity Ramachandran analysis

7cyq.H.pdb, model 1



87.7% (2115/2409) of all residues were in favored (90%) regions.
99.2% (2417/2439) of all residues were in allowed (≥90%) regions.

There were 22 outliers (Phi, Psi):

A 42 VAL (-177.4, 134.6)
A 832 PRO (-86.4, -151.0)
A 867 LEU (-99.2, -18.9)
A 907 LEU (-134.4, 96.9)
B 37 ARG (-17.4, -16.9)
B 98 LEU (-13.1, -12.2)
B 183 PRO (-134.6, 76.0)
D 113 GLY (-139.4, -47.0)
D 175 PRO (-72.1, -101.0)
F 249 ILE (-70.6, 49.0)
F 352 LEU (-17.7, 92.1)
F 414 THR (-33.4, 131.5)
F 482 ARG (-17.4, -18.9)
F 484 VAL (-46.7, -77.7)
F 549 THR (-66.7, 11.1)
E 175 PRO (-72.4, -160.9)
E 249 ILE (-206.6, 60.0)
E 352 LEU (-44.0, 68.2)
E 415 GLY (-27.4, 132.2)
E 549 THR (-43.7, 18.1)
G 7 VAL (-63.6, 74.7)
G 27 ASN (-176.2, 3.2)