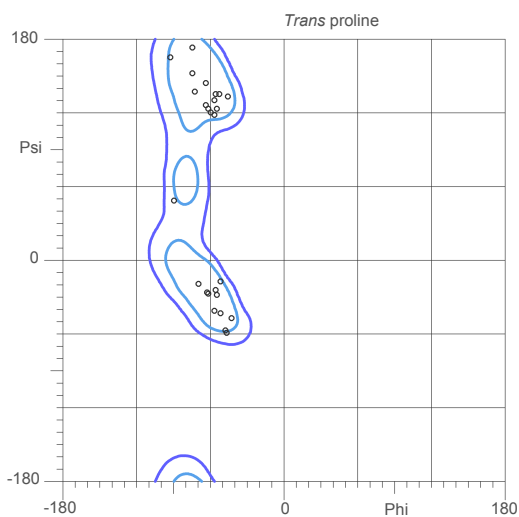
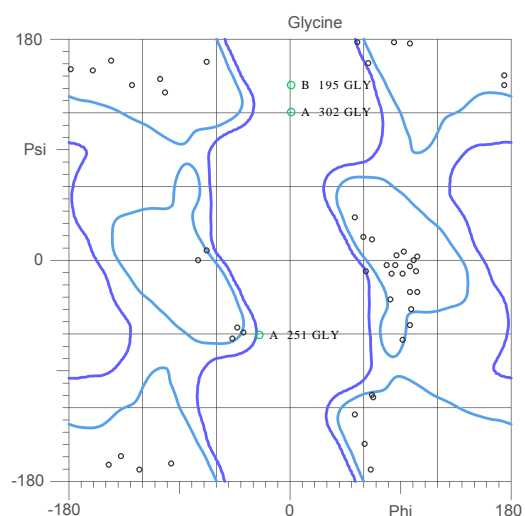
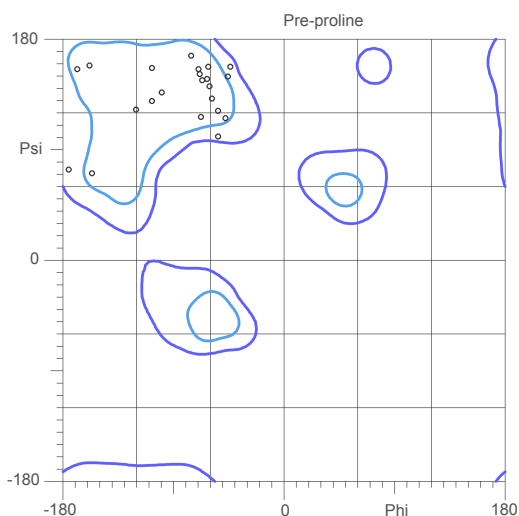
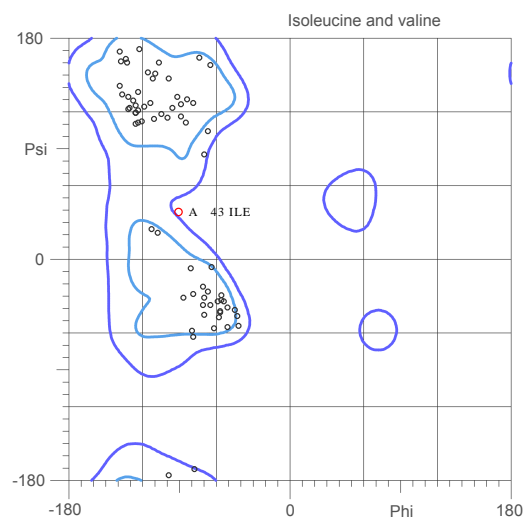
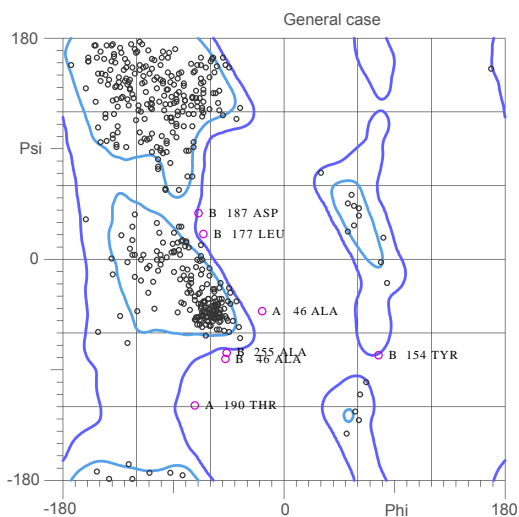


MolProbity Ramachandran analysis

2d2d.H.pdb, model 1



88.8% (532/599) of all residues were in favored (98%) regions.
98.2% (588/599) of all residues were in allowed (>99.8%) regions.

There were 11 outliers (phi, psi):

A 43 ILE (-91.9, 39.0)
A 46 ALA (-18.2, -42.1)
A 190 THR (-73.6, -120.0)
A 251 GLY (-25.9, -61.6)
A 302 GLY (1.8, 121.6)
B 46 ALA (-48.2, -82.0)
B 154 TYR (77.3, -79.0)
B 177 LEU (-66.7, 21.3)
B 187 ASP (-70.6, 38.5)
B 195 GLY (1.6, 143.1)
B 255 ALA (-47.8, -76.5)