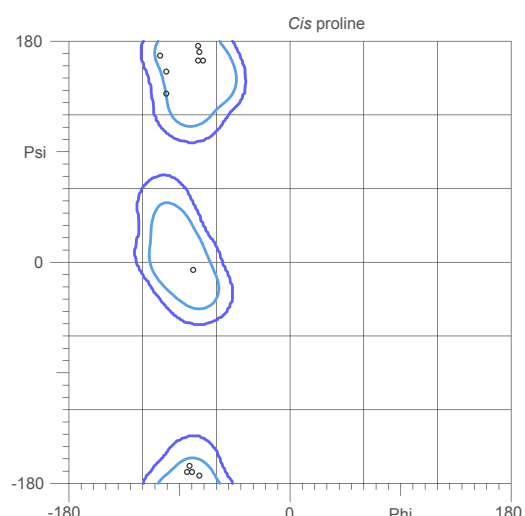
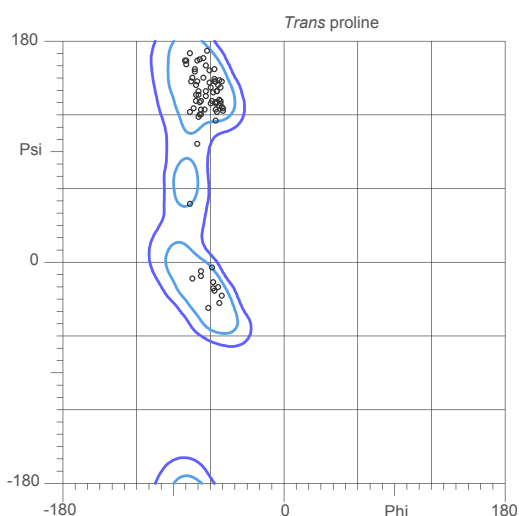
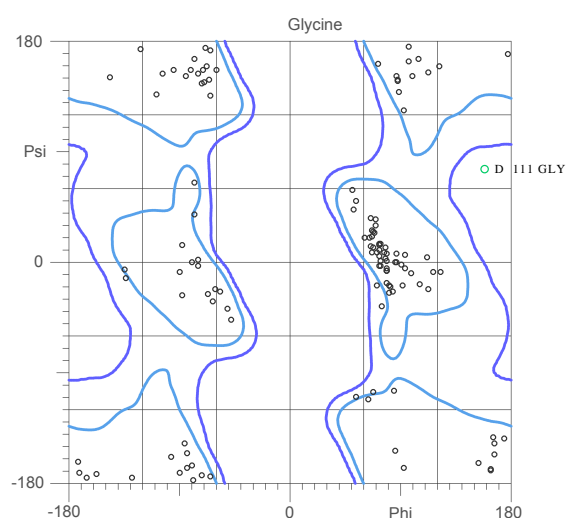
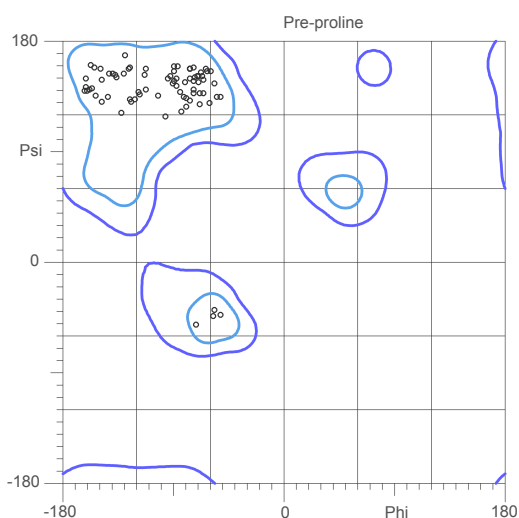
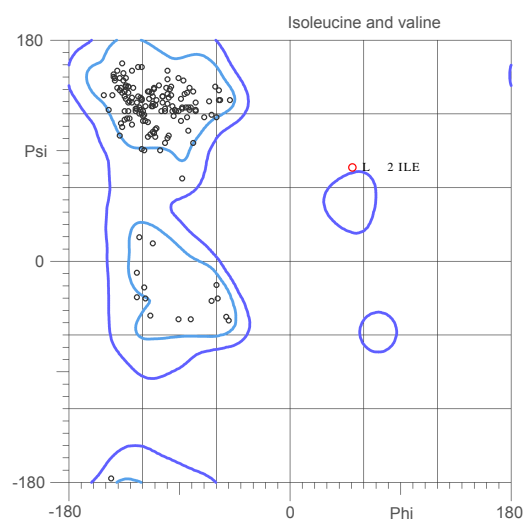
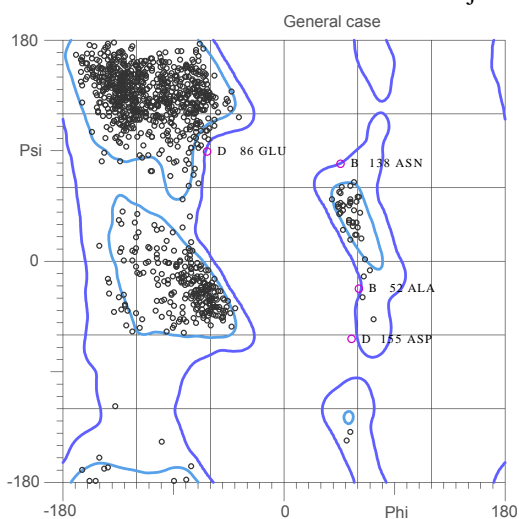


# MolProbity Ramachandran analysis

7jx3.H.pdb, model 1



95.3% (1426/1496) of all residues were in favored (98%) regions.  
99.6% (1490/1496) of all residues were in allowed (>99.8%) regions.

There were 6 outliers (phi, psi):

B 52 ALA (61.6, -22.7)  
B 138 ASN (46.3, 80.6)  
D 86 GLU (-63.3, 90.5)  
D 111 GLY (159.9, 76.9)  
D 155 ASP (56.0, -64.0)  
L 2 ILE (51.7, 77.8)