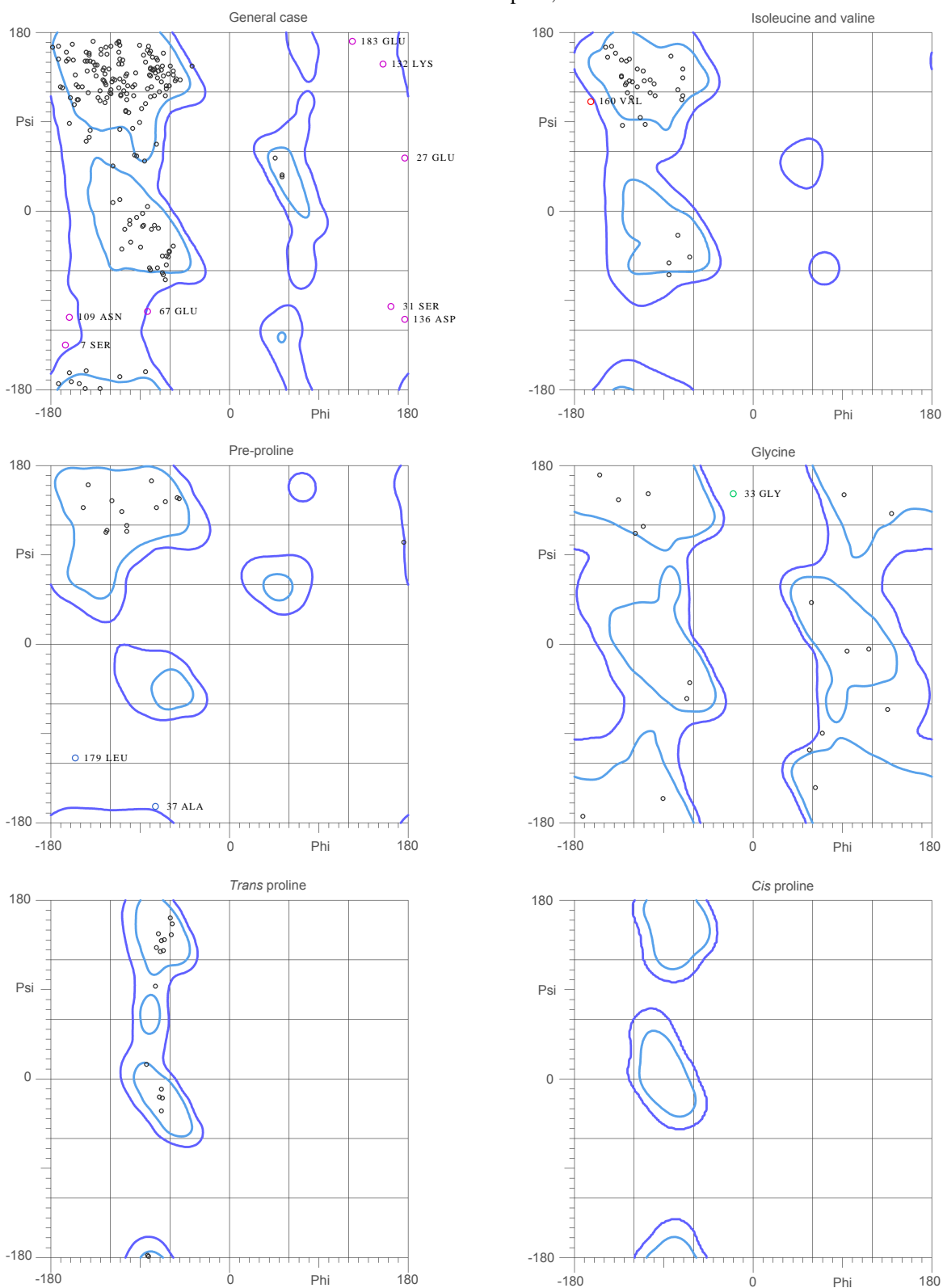


# MolProbity Ramachandran analysis

1a02N.B99990012.pdb, model 1



85.3% (237/278) of all residues were in favored (98%) regions.  
95.7% (266/278) of all residues were in allowed (>99.8%) regions.

There were 12 outliers (phi, psi):

7 SER (-166.1, -135.6)  
27 GLU (177.5, 54.9)  
31 SER (163.6, -96.8)  
33 GLY (-20.6, 152.1)  
37 ALA (-76.0, -164.2)  
67 GLU (-83.4, -101.4)  
109 ASN (-162.8, -107.7)  
132 LYS (155.1, 149.3)  
136 ASP (177.4, -109.9)  
160 VAL (-165.0, 111.5)  
179 LEU (-156.7, -115.9)  
183 GLU (124.4, 172.4)