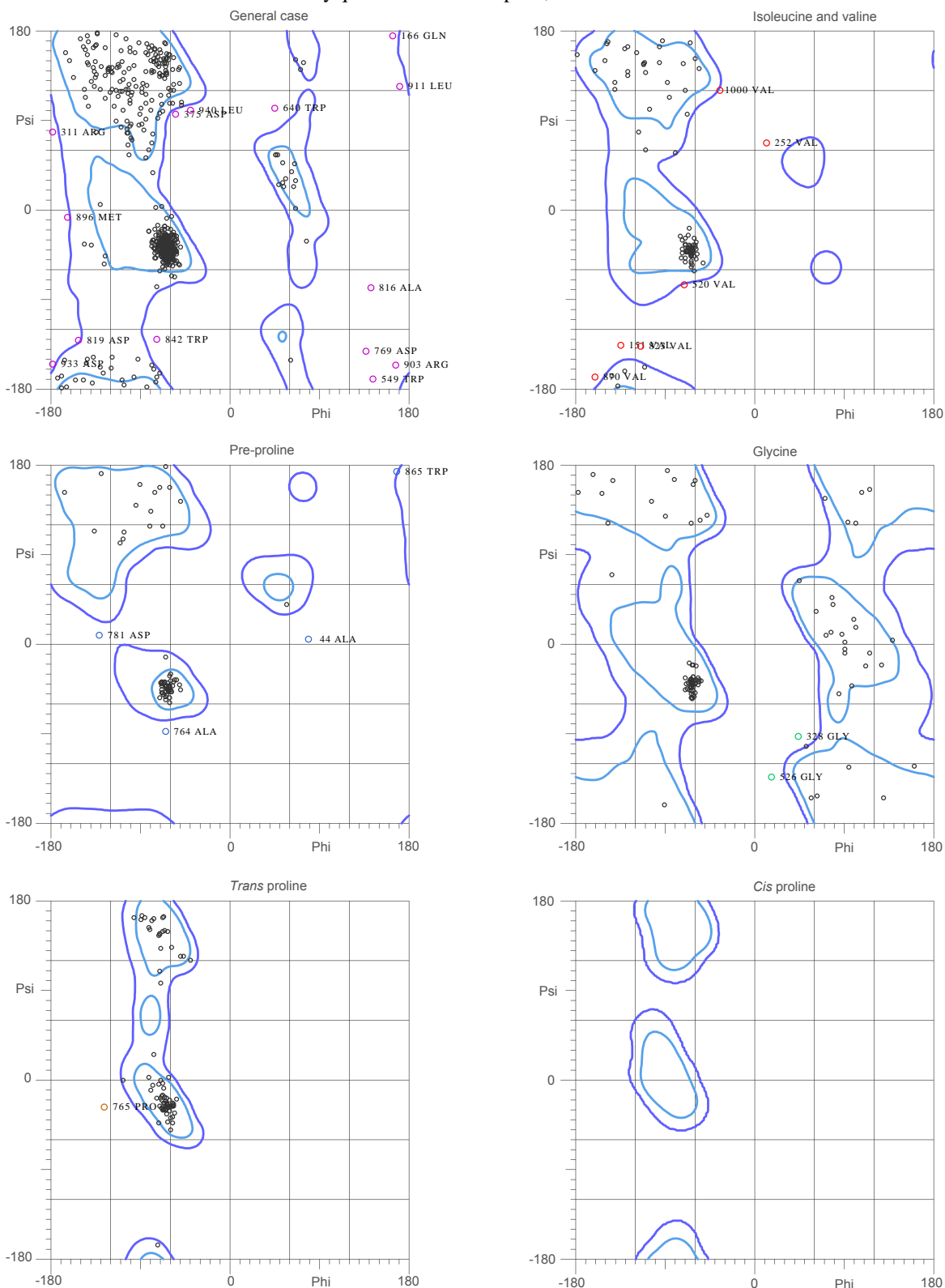


MolProbity Ramachandran analysis

1yq2.B99990032.pdb, model 1



85.7% (313/365) of all residues were in favored (Fav) regions.
10.2% (37/365) of all residues were in allowed (All) regions.

There were 27 outliers (Phi, Psi):

311 ARG (179.5, 1.7)
896 MET (180.0, -180.0)
819 ASP (180.0, -179.5)
842 TRP (180.0, -179.5)
865 TRP (180.0, -179.5)
816 ALA (180.0, -179.5)
769 ASP (180.0, -179.5)
903 ARG (180.0, -179.5)
549 TRP (180.0, -179.5)
781 ASP (179.5, 1.7)
44 ALA (180.0, -180.0)
764 ALA (180.0, -180.0)
328 GLY (180.0, -180.0)
326 GLY (180.0, -180.0)
765 PRO (180.0, -180.0)