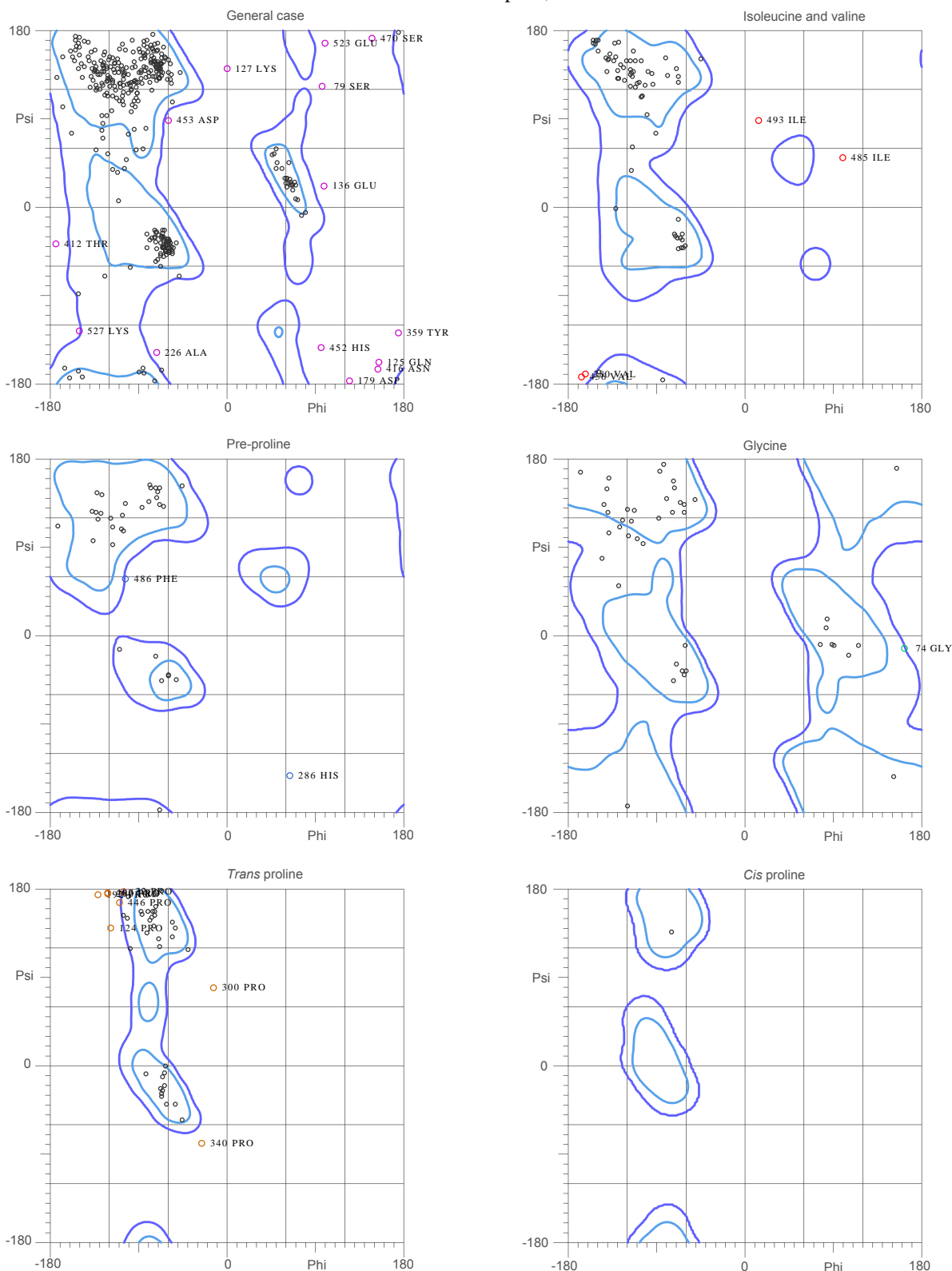


# MolProbity Ramachandran analysis

1aozA.B99990008.pdb, model 1



40.4% (271/669) of all residues were in favored (Fav) regions.  
96.7% (632/669) of all residues were in allowed (All) regions.

There were 30 outliers (3.0%):

10 PRO (1.0%), 1.0%  
10 SER (1.0%), 1.0%  
10 ASP (1.0%), 1.0%  
10 THR (1.0%), 1.0%  
10 LYS (1.0%), 1.0%  
10 GLU (1.0%), 1.0%  
10 HIS (1.0%), 1.0%  
10 TYR (1.0%), 1.0%  
10 ILE (1.0%), 1.0%  
10 VAL (1.0%), 1.0%  
10 PHE (1.0%), 1.0%  
10 LEU (1.0%), 1.0%  
10 ARG (1.0%), 1.0%  
10 CYS (1.0%), 1.0%  
10 MET (1.0%), 1.0%  
10 GLY (1.0%), 1.0%  
10 ALA (1.0%), 1.0%  
10 ASP (1.0%), 1.0%  
10 THR (1.0%), 1.0%  
10 LYS (1.0%), 1.0%  
10 GLU (1.0%), 1.0%  
10 HIS (1.0%), 1.0%  
10 TYR (1.0%), 1.0%  
10 ILE (1.0%), 1.0%  
10 VAL (1.0%), 1.0%  
10 PHE (1.0%), 1.0%  
10 LEU (1.0%), 1.0%  
10 ARG (1.0%), 1.0%  
10 CYS (1.0%), 1.0%  
10 MET (1.0%), 1.0%  
10 GLY (1.0%), 1.0%  
10 ALA (1.0%), 1.0%