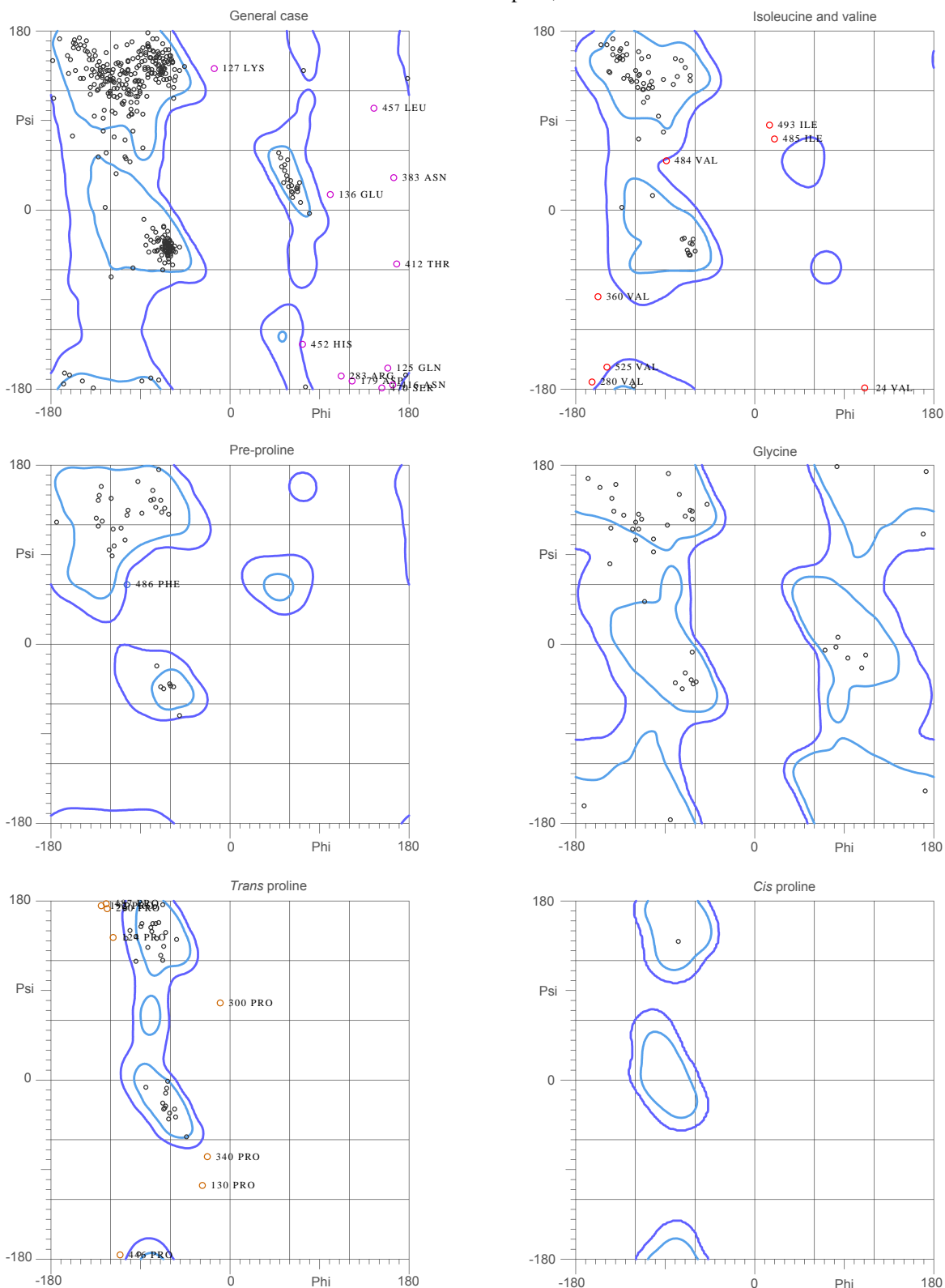


# MolProbity Ramachandran analysis

1aozA.B99990032.pdb, model 1



86.9% (178/203) of all residues were in favored (Fav) regions.  
10.2% (20/203) of all residues were in allowed (All) regions.

Below are 27 outliers (Phi, Psi):  
 127 LYS (113.1, 179.2)  
 457 LEU (108.6, 164.4)  
 136 GLU (108.6, 164.4)  
 383 ASN (108.6, 164.4)  
 412 THR (108.6, 164.4)  
 452 HIS (108.6, 164.4)  
 283 ARG (108.6, 164.4)  
 125 GLN (108.6, 164.4)  
 179 SER (108.6, 164.4)  
 493 ILE (108.6, 164.4)  
 485 ILE (108.6, 164.4)  
 484 VAL (108.6, 164.4)  
 360 VAL (108.6, 164.4)  
 525 VAL (108.6, 164.4)  
 580 VAL (108.6, 164.4)  
 24 VAL (108.6, 164.4)