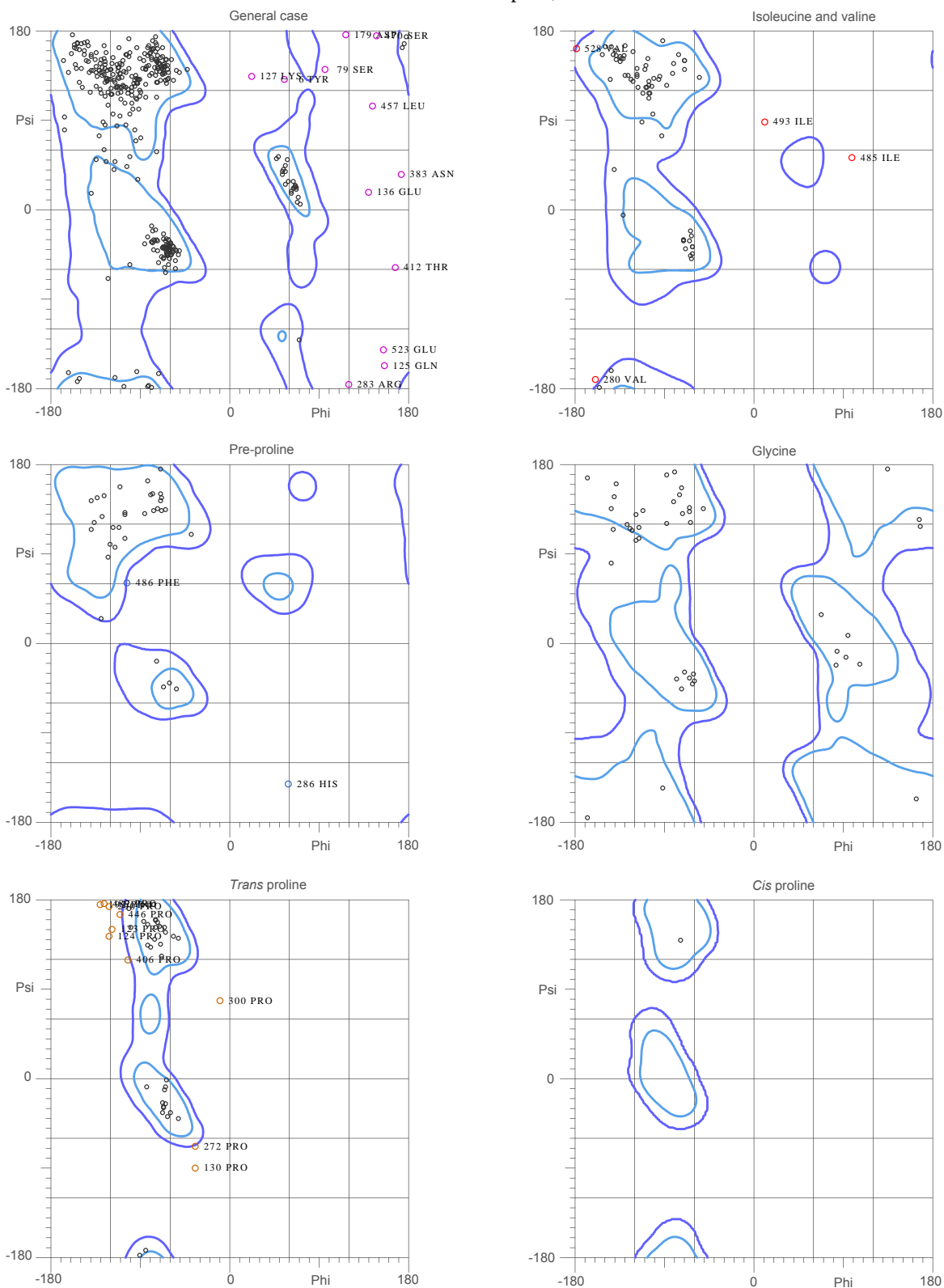


# MolProbity Ramachandran analysis

1aozA.B99990031.pdb, model 1



87.5% (270/309) of all residues were in favored (98%) regions.  
10.0% (32/309) of all residues were in allowed (1.9% (6%)) regions.

There were 26 outliers (8%):

Residue	Phi (deg)	Psi (deg)
179 ASP	179.0	179.0
127 LYS	127.0	127.0
79 SER	79.0	79.0
457 LEU	457.0	457.0
383 ASN	383.0	383.0
136 GLU	136.0	136.0
412 THR	412.0	412.0
523 GLU	523.0	523.0
125 GLN	125.0	125.0
283 ARG	283.0	283.0
486 PHE	486.0	486.0
286 HIS	286.0	286.0
446 PRO	446.0	446.0
124 PRO	124.0	124.0
406 PRO	406.0	406.0
300 PRO	300.0	300.0
272 PRO	272.0	272.0
130 PRO	130.0	130.0