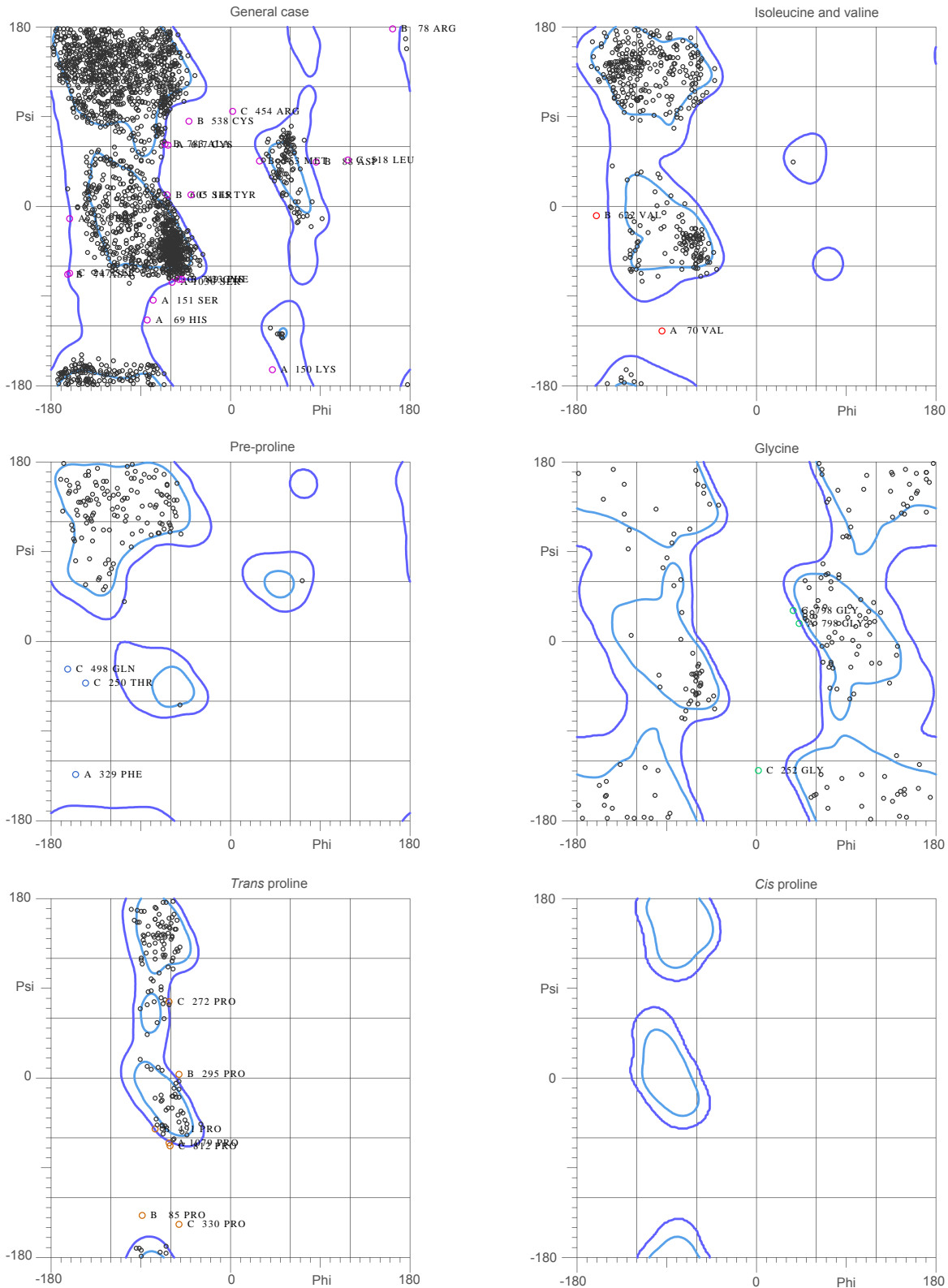


# MolProbity Ramachandran analysis

7cn9.H.pdb, model 1



75 (100%) of all residues were in favored (blue) regions.  
0 (0%) of all residues were in allowed (cyan) regions.  
None were in outlier (red) regions.

Residues in favored regions (100%):

- A 1033 SER
- A 150 LYS
- A 151 SER
- A 69 HIS
- B 518 LEU
- B 538 CYS
- B 605 SER
- B 608 TYR
- B 767 ALA
- B 78 ARG
- C 454 ARG
- C 98 GLY
- C 230 THR
- C 251 GLY
- C 272 PRO
- C 330 PRO
- C 798 GLY
- K 947 ASN

Residues in allowed regions (0%):

- A 329 PHE
- B 85 PRO
- B 295 PRO
- C 1479 PRO

Residues in outlier regions (0%):

- A 70 VAL
- B 622 VAL