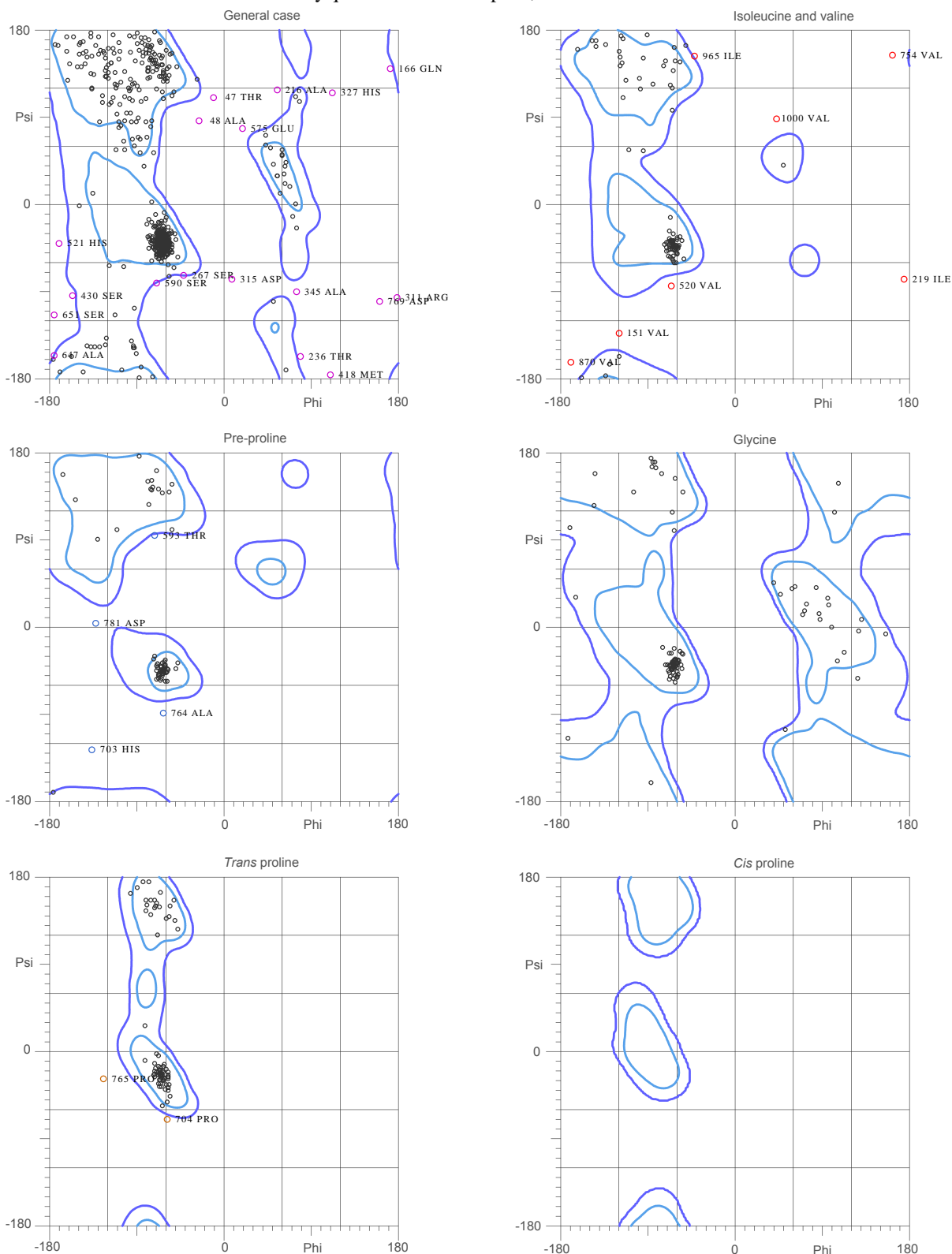


MolProbity Ramachandran analysis

1yq2.B99990028.pdb, model 1



96.0% (252/262) of all residues were in favored (Fav) regions.
0.0% (0/262) of all residues were in allowed (All) regions.

Residue (chain ID, residue number)

166 GLN (A, 166)
47 THR (A, 47)
216 ALA (A, 216)
327 HIS (A, 327)
48 ALA (A, 48)
576 GLU (A, 576)
521 HIS (A, 521)
430 SER (A, 430)
651 SER (A, 651)
647 ALA (A, 647)
267 SER (A, 267)
315 ASP (A, 315)
345 ALA (A, 345)
765 ASP (A, 765)
236 THR (A, 236)
418 MET (A, 418)
965 ILE (A, 965)
754 VAL (A, 754)
1000 VAL (A, 1000)
520 VAL (A, 520)
219 ILE (A, 219)
151 VAL (A, 151)
870 VAL (A, 870)
593 THR (A, 593)
781 ASP (A, 781)
764 ALA (A, 764)
703 HIS (A, 703)
765 PRO (A, 765)
704 PRO (A, 704)