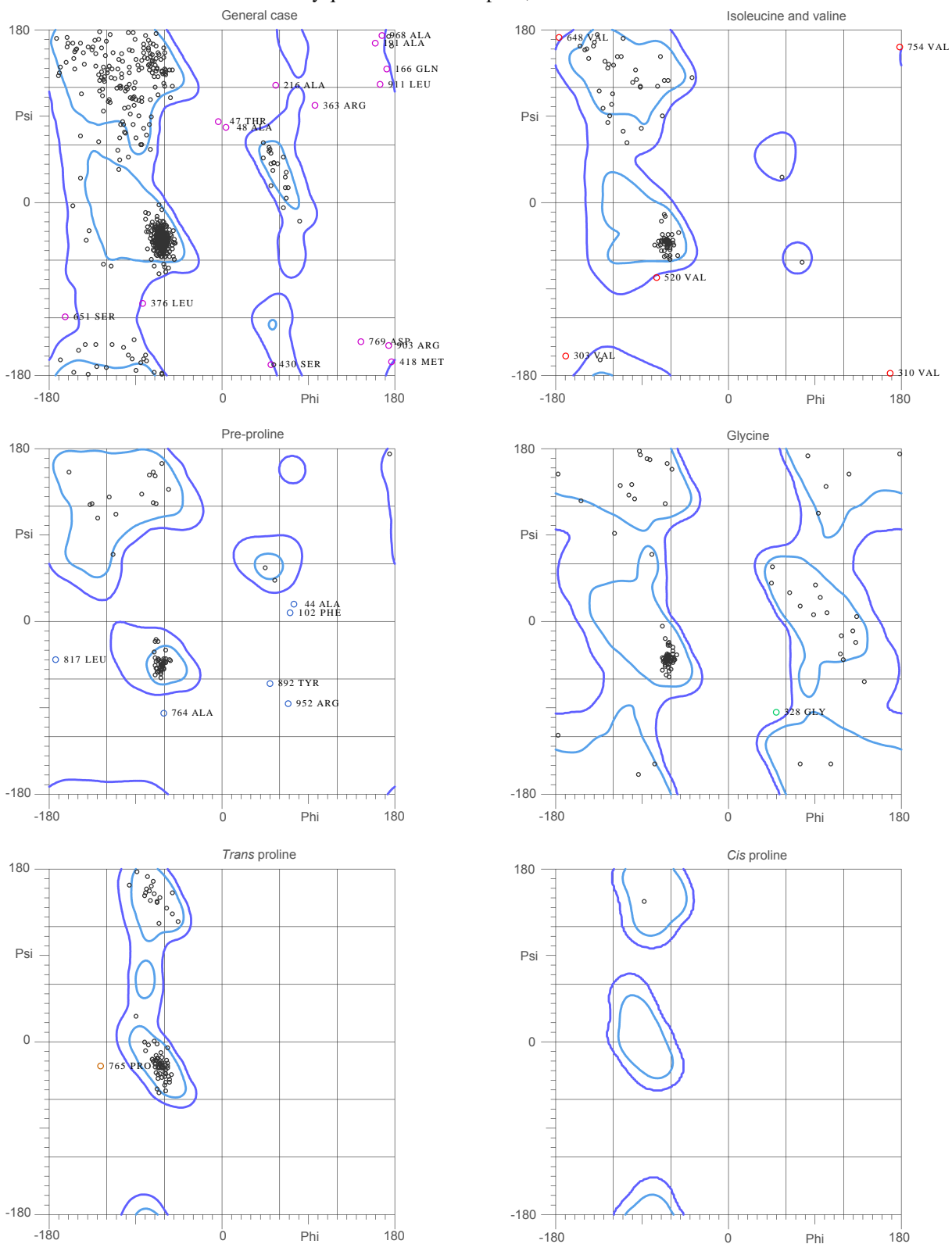


MolProbity Ramachandran analysis

1yq2.B99990004.pdb, model 1



90.7% (319/352) of all residues were in favored (Fav) regions.
97.2% (305/314) of all residues were in allowed (All) regions.

Shown here: 27 outliers (Phi, Psi):

668 ALA (75.9, 18.3)
181 ALA (75.9, 18.3)
216 ALA (75.9, 18.3)
166 GLN (75.9, 18.3)
911 LEU (75.9, 18.3)
47 THR (75.9, 18.3)
48 LEU (75.9, 18.3)
363 ARG (75.9, 18.3)
651 SER (75.9, 18.3)
376 LEU (75.9, 18.3)
430 SER (75.9, 18.3)
769 ARG (75.9, 18.3)
418 MET (75.9, 18.3)
44 ALA (75.9, 18.3)
102 PHE (75.9, 18.3)
817 LEU (75.9, 18.3)
892 TYR (75.9, 18.3)
952 ARG (75.9, 18.3)
764 ALA (75.9, 18.3)
928 GLY (75.9, 18.3)
765 PRO (75.9, 18.3)