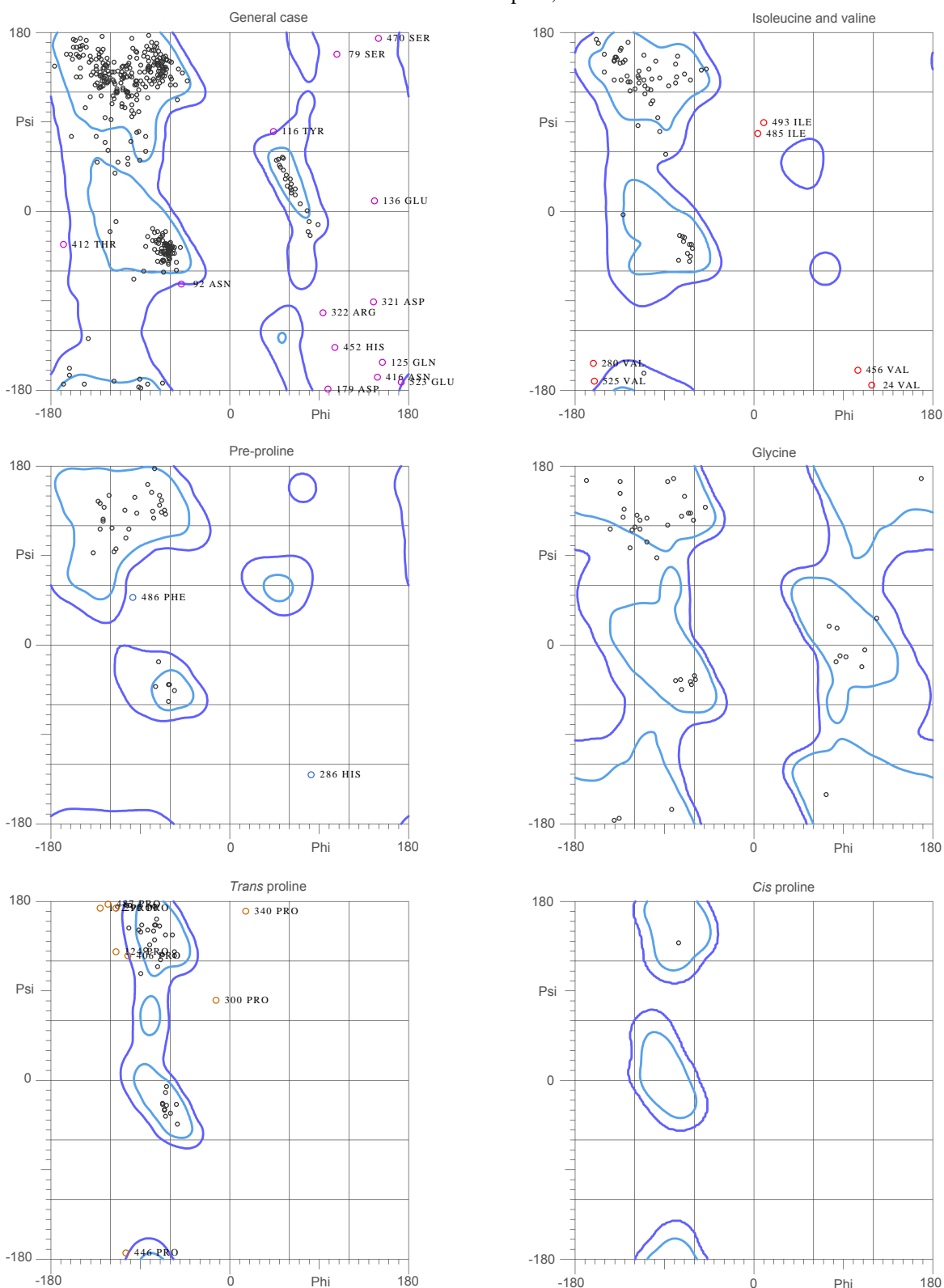


MolProbity Ramachandran analysis

1aozA.B99990034.pdb, model 1



48.0% (240/500) of all residues were in favored (98%) regions.
56.7% (283/500) of all residues were in allowed (1.0%) regions.

There were 39 outliers (8%):

39 VAL (116.1, 17.0)
79 SER (108.7, 10.0)
92 ASN (107.1, 10.0)
116 TYR (108.0, 10.0)
136 GLU (116.0, 10.0)
179 ASP (107.0, 10.0)
280 VAL (116.0, 10.0)
300 PRO (116.0, 10.0)
321 ASP (116.0, 10.0)
322 ARG (116.0, 10.0)
323 GLU (116.0, 10.0)
412 THR (116.0, 10.0)
416 ASN (116.0, 10.0)
452 HIS (116.0, 10.0)
485 ILE (116.0, 10.0)
486 PHE (116.0, 10.0)
493 ILE (116.0, 10.0)
525 VAL (116.0, 10.0)
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