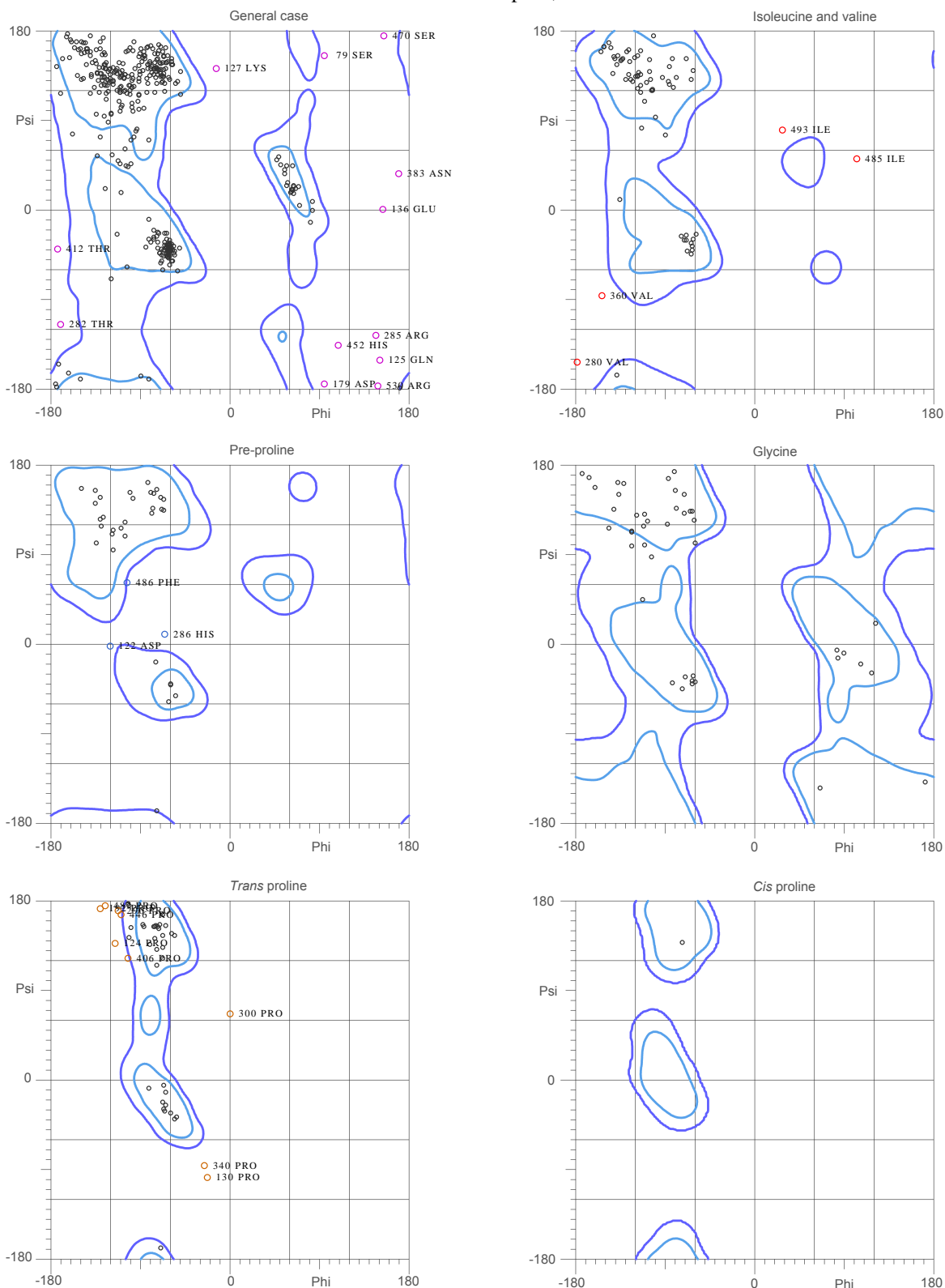


MolProbity Ramachandran analysis

1aozA.B99990035.pdb, model 1



86.9% (278/320) of all residues were in favored (98%) regions.
14.1% (45/320) of all residues were in allowed (1.9%) regions.

There were 28 outliers (phi, psi):

79 SER	(172.0, 178.0)
127 LYS	(172.0, 178.0)
136 GLU	(172.0, 178.0)
142 THR	(172.0, 178.0)
179 ASP	(172.0, 178.0)
282 THR	(172.0, 178.0)
285 ARG	(172.0, 178.0)
340 PRO	(172.0, 178.0)
343 ASN	(172.0, 178.0)
452 HIS	(172.0, 178.0)
485 ILE	(172.0, 178.0)
493 ILE	(172.0, 178.0)
534 ARG	(172.0, 178.0)
536 ARG	(172.0, 178.0)
537 ARG	(172.0, 178.0)
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581 ARG	(172.0, 178.0)
582 ARG	(172.0, 178.0)
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600 ARG	(172.0, 178.0)