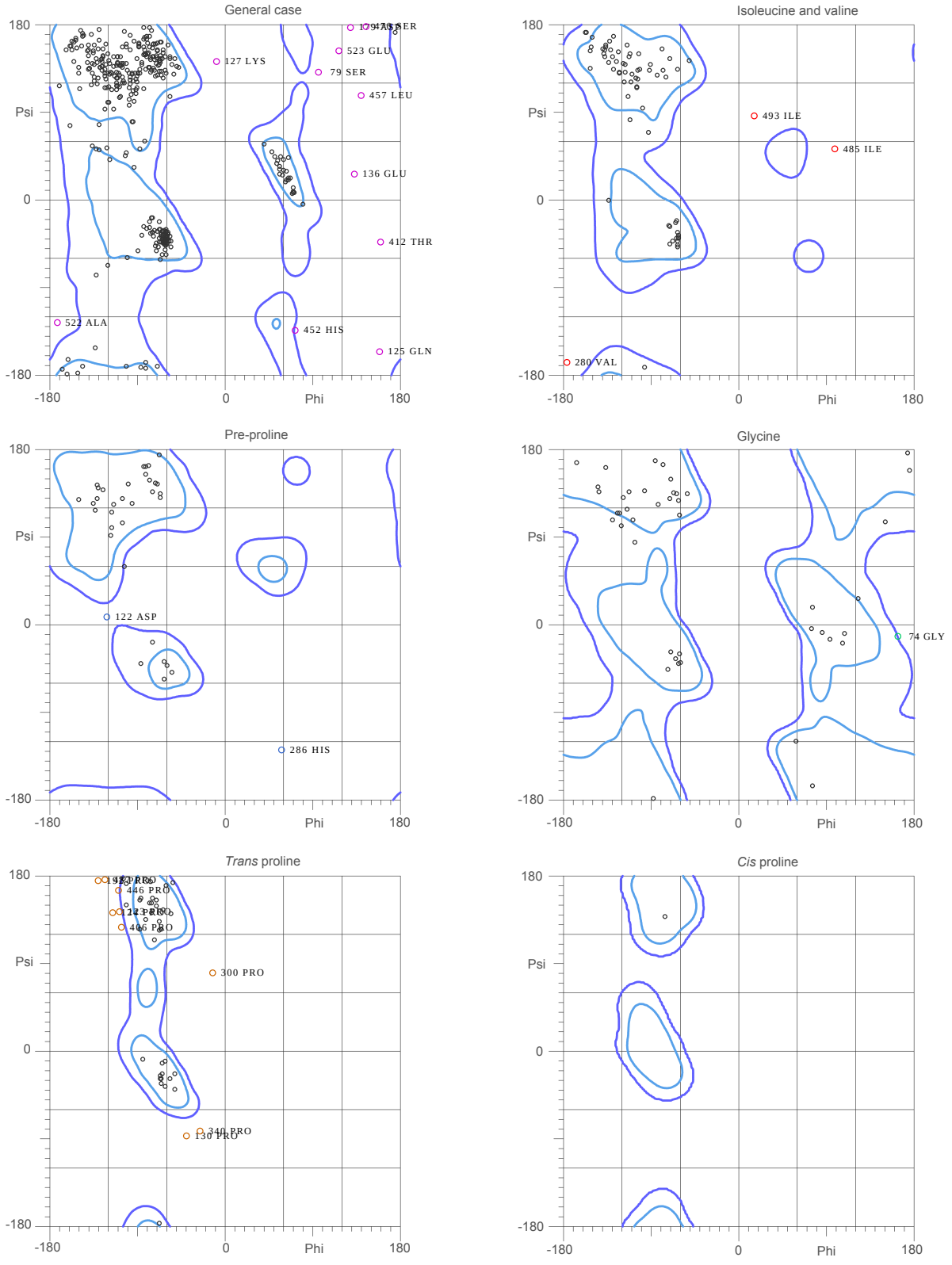
MolProbity Ramachandran analysis

1aozA.B99990007.pdb, model 1



88.0% (484/550) of all residues were in favoured (98%) regions.
95.3% (524/550) of all residues were in allowed (>98.0%) regions.

There were 26 outliers (pht, psi):
74 GLY (164.3, -12.8)
79 SER (96.1, 151.9)
122 ASP (-122.0, 8.9)
123 PRO (-101.9, 144.3)
124 PRO (-116.6, 144.0)
125 GLN (150.2, -156.4)
127 LYS (-9.9, 143.1)
130 PRO (-85.1, -87.9)
136 GLU (133.7, 27.2)
179 ASP (-129.6, 178.5)
192 PRO (-131.3, 176.6)
280 VAL (-177.4, -168.0)
286 HIS (58.3, -125.3)
300 PRO (-51.2, 151.9)
340 PRO (-26.2, -83.0)
406 PRO (-167.6, 139.0)
412 THR (160.9, -43.7)
446 PRO (-115.4, 166.7)
452 HIS (72.9, -134.6)
457 LEU (140.2, 168.3)
479 SER (146.6, 179.5)
485 ILE (99.7, 53.5)
487 PRO (-124.7, 177.1)
493 ILE (66.3, 87.2)
522 ALA (-171.5, -156.5)
523 GLU (117.6, 154.7)

Lovell, Davis, et al. Proteins 50:437 (2003)