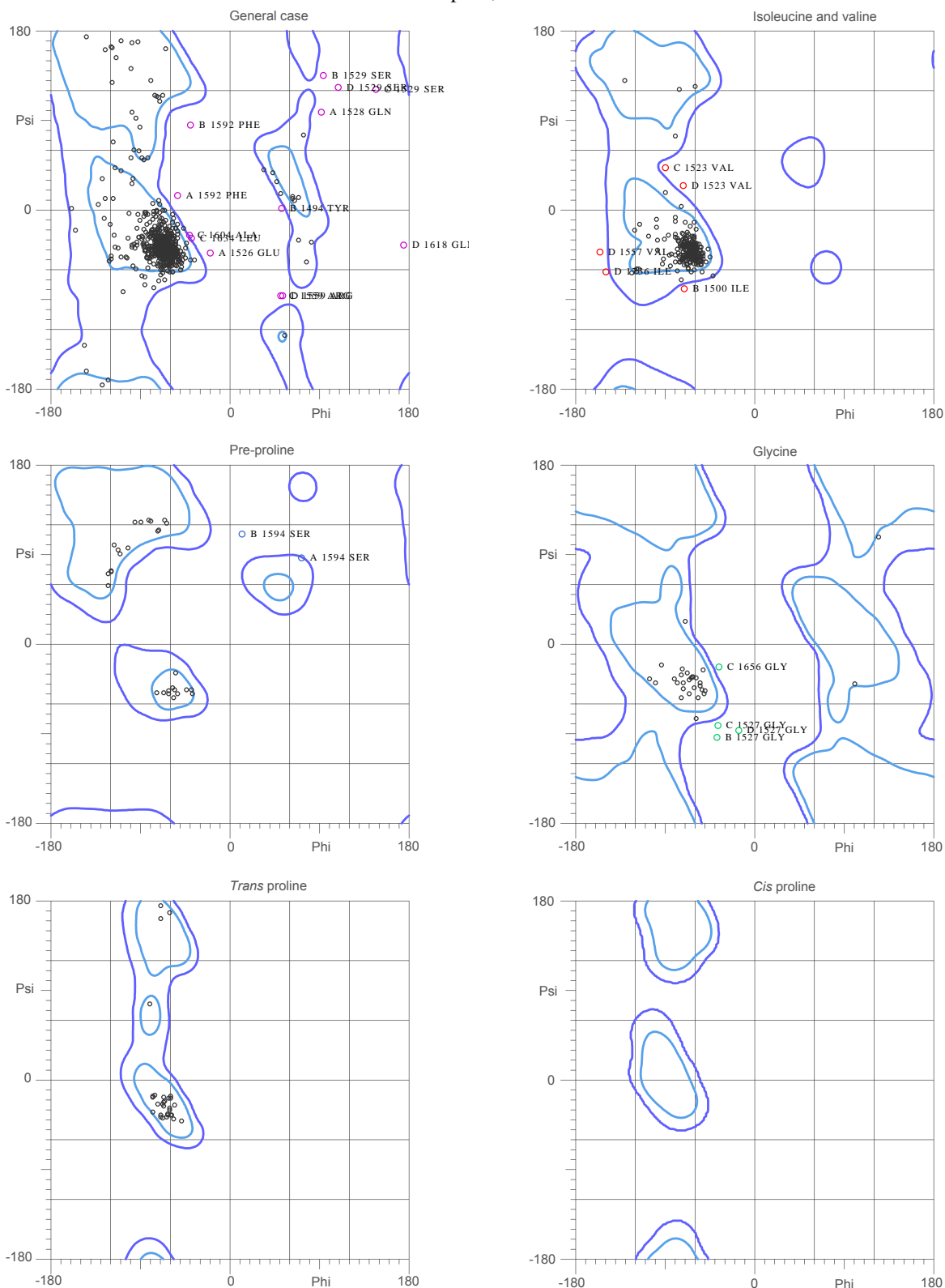


MolProbity Ramachandran analysis

5ek0.H.pdb, model 1



92.6% (105/113) of all residues were in favored (90%) regions.
 7.4% (8/113) of all residues were in allowed (10-90%) regions.

There were 39 outliers (phi, psi):

A 1524 GLU (-28.5, -47.5)
 A 1525 GLU (-33.7, -59.3)
 A 1526 PHE (-53.2, -13.1)
 A 1528 SER (-23.6, -66.6)
 B 1494 TYR (-32.6, 3.7)
 B 1500 SER (-73.1, -79.9)
 B 1517 GLY (-18.6, -66.6)
 B 1518 SER (-6.6, -106.3)
 B 1519 PHE (-49.7, -66.3)
 B 1519 SER (-23.3, -11.9)
 C 1527 VAL (-96.6, -41.3)
 C 1527 GLY (-106.6, -82.6)
 C 1528 SER (-17.7, -123.1)
 C 1529 ARG (-21.6, -86.6)
 C 1530 ALA (-21.0, -23.6)
 C 1531 ALA (-21.0, -23.6)
 C 1532 ALA (-21.0, -23.6)
 C 1533 VAL (-72.3, -24.6)
 D 1527 GLY (-16.7, -67.9)
 D 1528 SER (-109.7, -124.1)
 D 1527 VAL (-156.1, -42.3)
 D 1519 ARG (-53.6, -87.6)
 D 1518 GLU (-107.6, -10.6)
 D 1518 ILE (-139.1, -42.6)