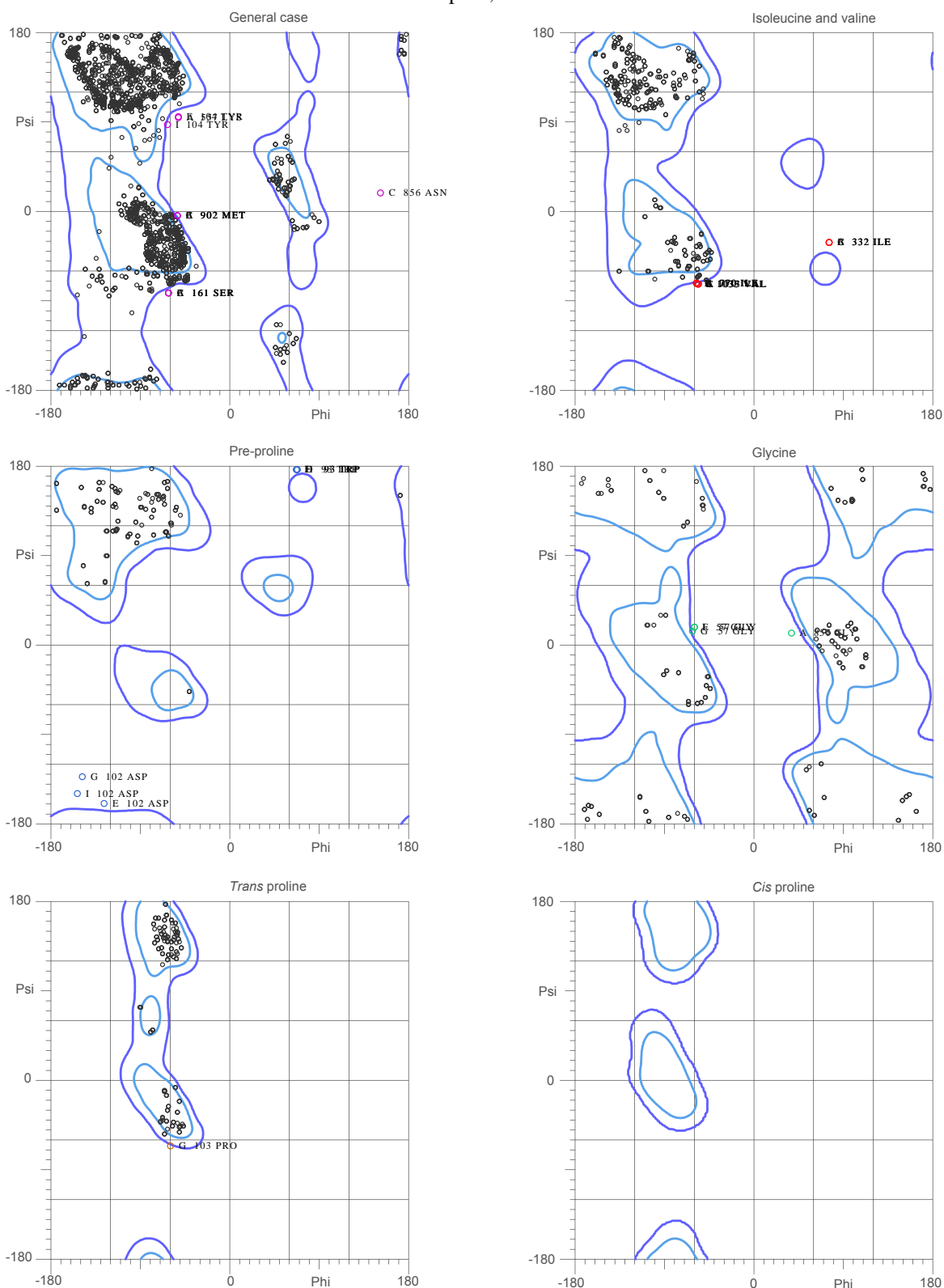


MolProbity Ramachandran analysis

7cak.H.pdb, model 1



40 (4) (100.0%) of all residues were in favored (90%) regions.

40 (4) (100.0%) of all residues were in allowed (100.0%) regions.

There were 50 outliers (40%):

- A 100 ASP (100.0, 100.0)
- A 101 ASP (100.0, 100.0)
- A 102 ASP (100.0, 100.0)
- A 103 ASP (100.0, 100.0)
- A 104 ASP (100.0, 100.0)
- A 105 ASP (100.0, 100.0)
- A 106 ASP (100.0, 100.0)
- A 107 ASP (100.0, 100.0)
- A 108 ASP (100.0, 100.0)
- A 109 ASP (100.0, 100.0)
- A 110 ASP (100.0, 100.0)
- A 111 ASP (100.0, 100.0)
- A 112 ASP (100.0, 100.0)
- A 113 ASP (100.0, 100.0)
- A 114 ASP (100.0, 100.0)
- A 115 ASP (100.0, 100.0)
- A 116 ASP (100.0, 100.0)
- A 117 ASP (100.0, 100.0)
- A 118 ASP (100.0, 100.0)
- A 119 ASP (100.0, 100.0)
- A 120 ASP (100.0, 100.0)
- A 121 ASP (100.0, 100.0)
- A 122 ASP (100.0, 100.0)
- A 123 ASP (100.0, 100.0)
- A 124 ASP (100.0, 100.0)
- A 125 ASP (100.0, 100.0)
- A 126 ASP (100.0, 100.0)
- A 127 ASP (100.0, 100.0)
- A 128 ASP (100.0, 100.0)
- A 129 ASP (100.0, 100.0)
- A 130 ASP (100.0, 100.0)
- A 131 ASP (100.0, 100.0)
- A 132 ASP (100.0, 100.0)
- A 133 ASP (100.0, 100.0)
- A 134 ASP (100.0, 100.0)
- A 135 ASP (100.0, 100.0)
- A 136 ASP (100.0, 100.0)
- A 137 ASP (100.0, 100.0)
- A 138 ASP (100.0, 100.0)
- A 139 ASP (100.0, 100.0)
- A 140 ASP (100.0, 100.0)