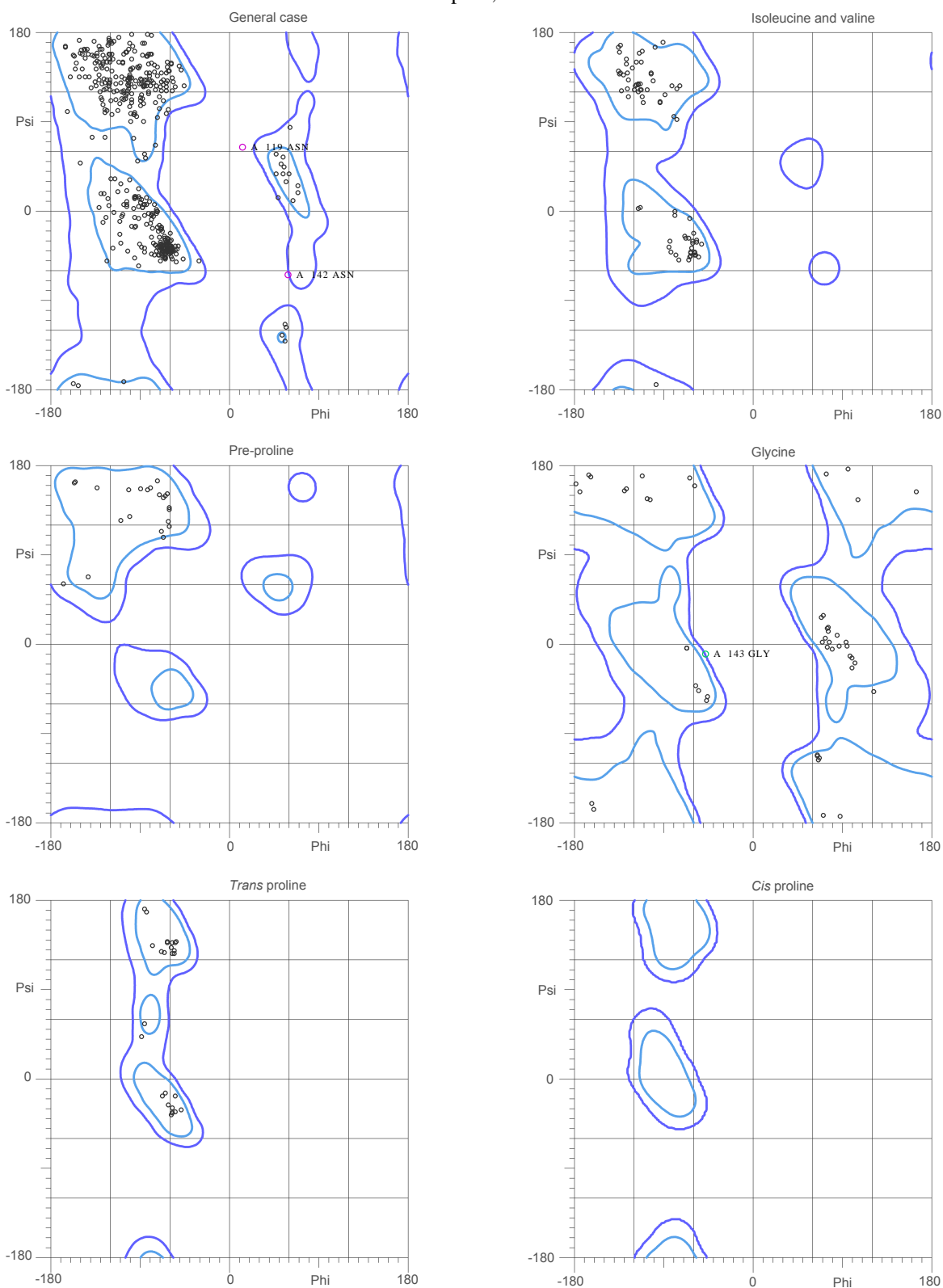


MolProbity Ramachandran analysis

7kw5.H.pdb, model 1



96.2% (579/602) of all residues were in favored (98%) regions.
99.5% (599/602) of all residues were in allowed (>99.8%) regions.

There were 3 outliers (ϕ , ψ):

- A 119 ASN (13.5, 65.6)
- A 142 ASN (59.6, -64.8)
- A 143 GLY (-48.1, -10.4)