MolProbity Ramachandran analysis

3sna.H.pdb, model 1

General case

A 80 HIS
A 277 ASN
A 41 HIS
A 211 ALA
A 50 LEU
A 49 MET
A 154 TYR
A 33 ASP
H 2 SER

Isoleucine and valine

Pre-proline

Glycine

A 278 GLY

Trans proline

*Cis* proline
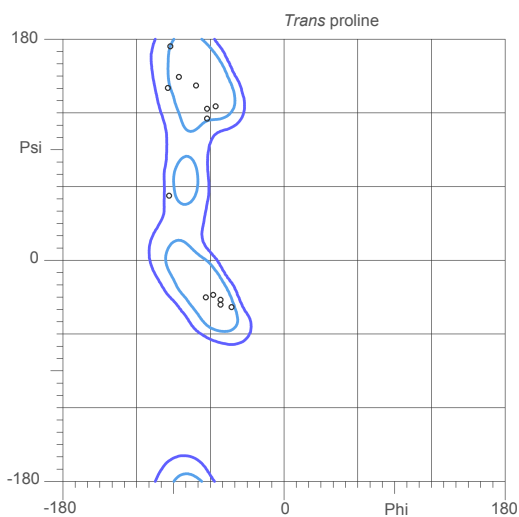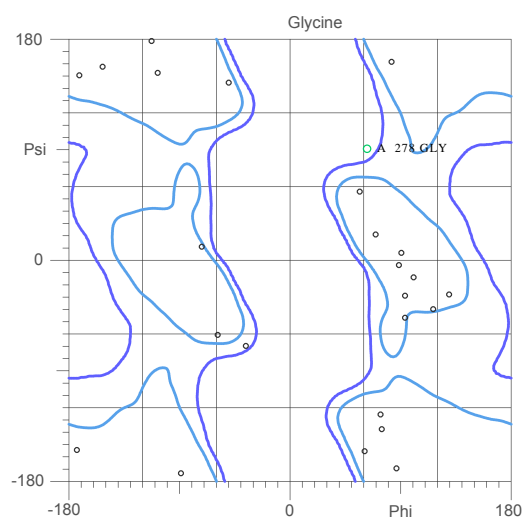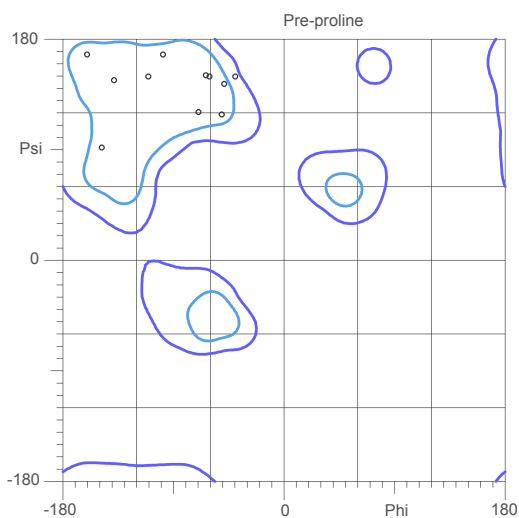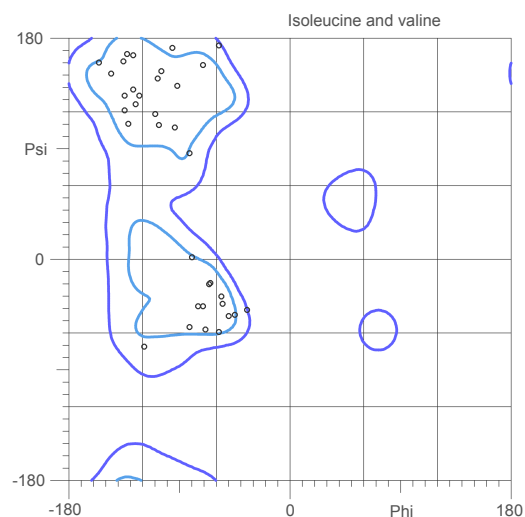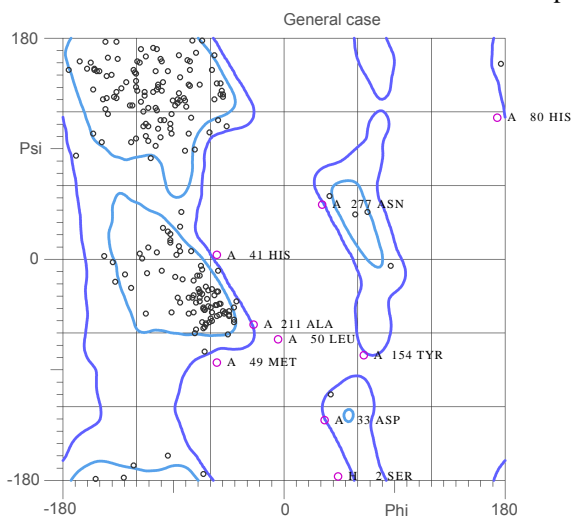
84.5% (256/303) of all residues were in favored (98%) regions.
96.7% (293/303) of all residues were in allowed (>99.8%) regions.

There were 10 outliers (phi, psi):
    A   33 ASP (33.2, -131.2)
    A   41 HIS (-56.0, 4.7)
    A   49 MET (-55.2, -84.4)
    A   50 LEU (-5.7, -65.5)
    A   80 HIS (174.6, 117.0)
    A 154 TYR (66.0, -78.8)
    A 211 ALA (-25.6, -53.6)
    A 277 ASN (31.4, 45.1)
    A 278 GLY (63.0, 91.6)
    H   2 SER (44.0, -177.2)