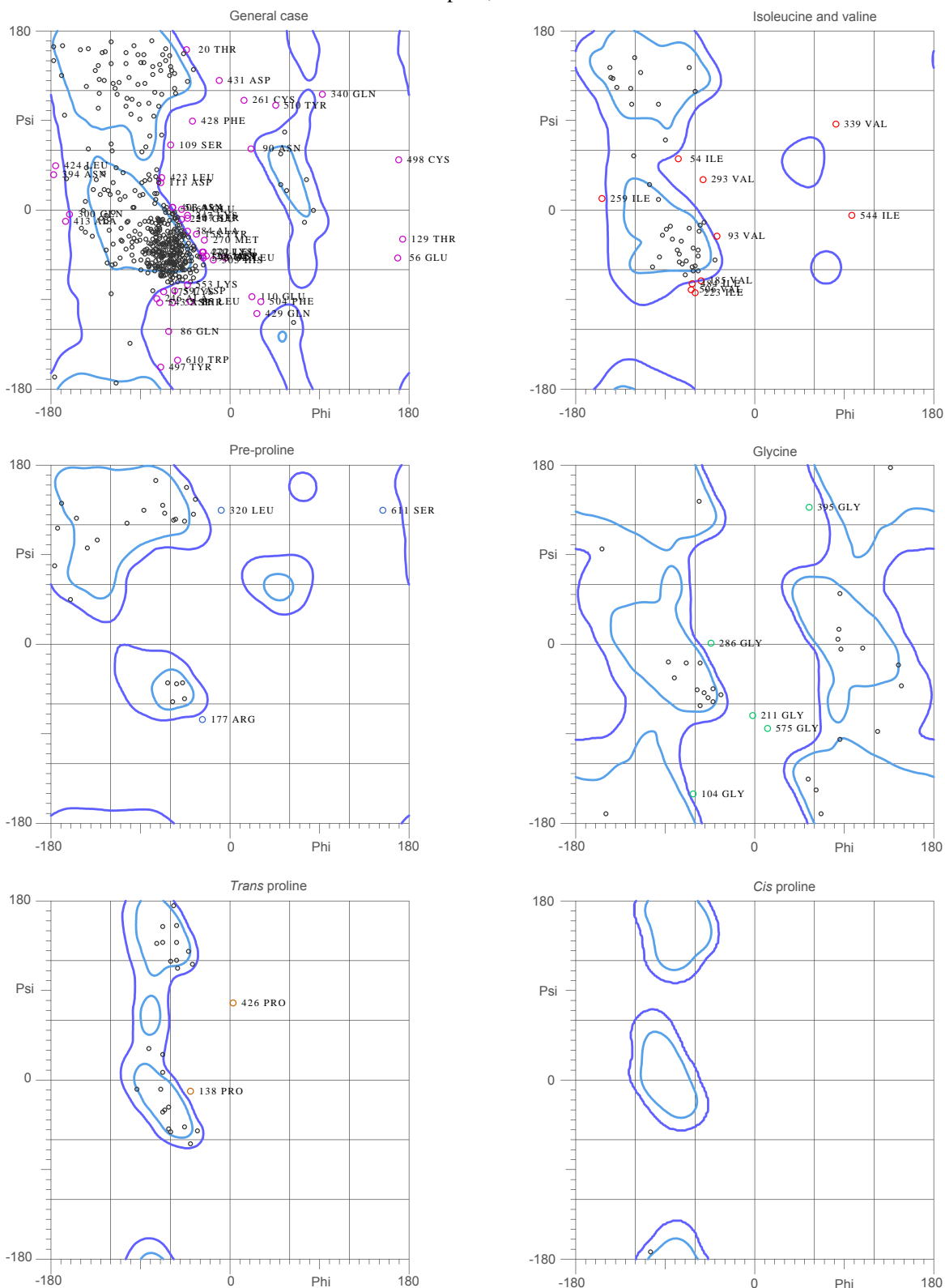


MolProbity Ramachandran analysis

1r4l.H.pdb, model 1



47.0% (100/210) of all residues were in favored (99%) regions.
50.0% (105/210) of all residues were in allowed (100%) regions.
There were 0 outliers (0% out).

50 THR, 48.0, 10.0, 10.0
51 ASP, 10.0, 10.0, 10.0
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53 THR, 10.0, 10.0, 10.0
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