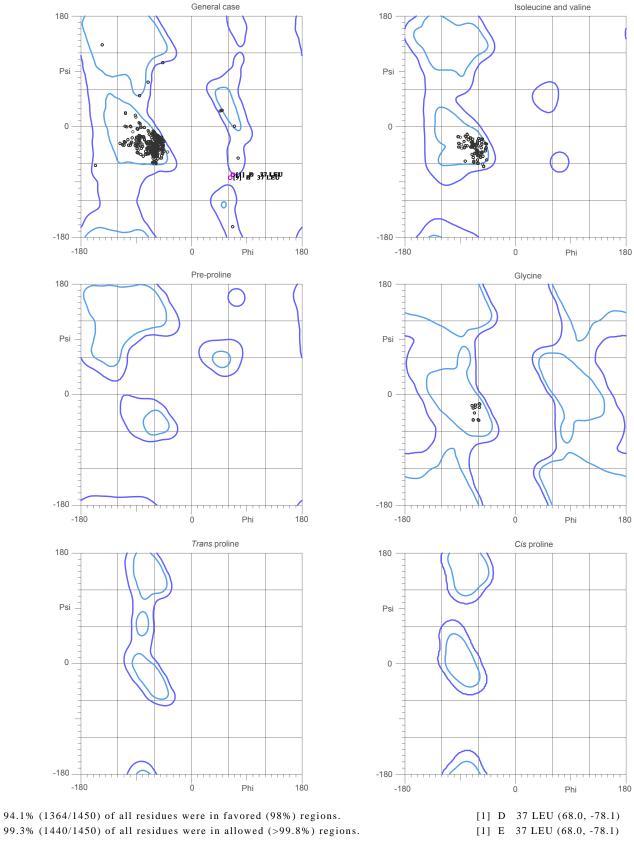
7k3g.H.pdb, all models



99.3% (1440/1450) of all residues were in allowed (>99.8%) regions.

There were 10 outliers (phi, psi):

- [1] A 37 LEU (68.0, -78.1)
- [1] B 37 LEU (68.0, -78.0)
- [1] C 37 LEU (68.1, -78.1)

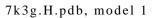
A 37 LEU (63.4, -83.8)

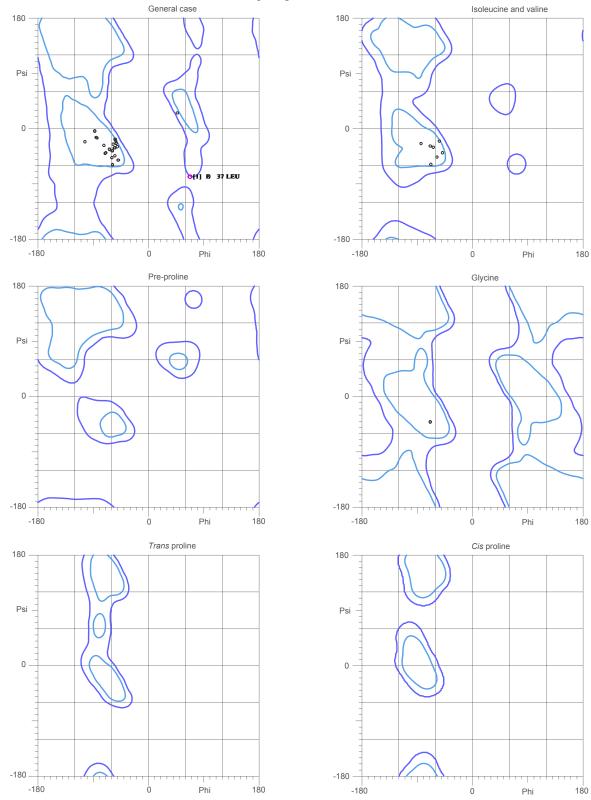
37 LEU (63.4, -83.8)

[9] C 37 LEU (63.5, -83.9)

[9] D 37 LEU (63.3, -83.9)

[9] E 37 LEU (63.3, -83.9)

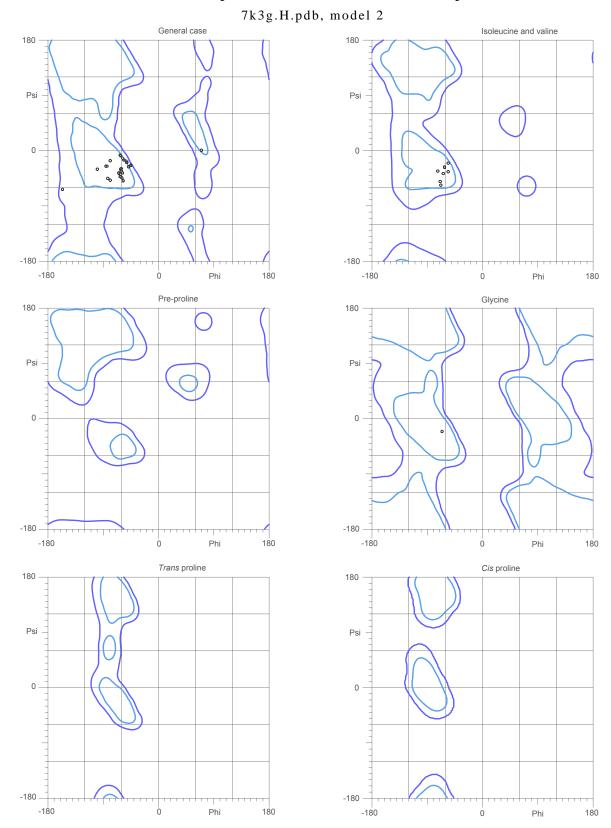




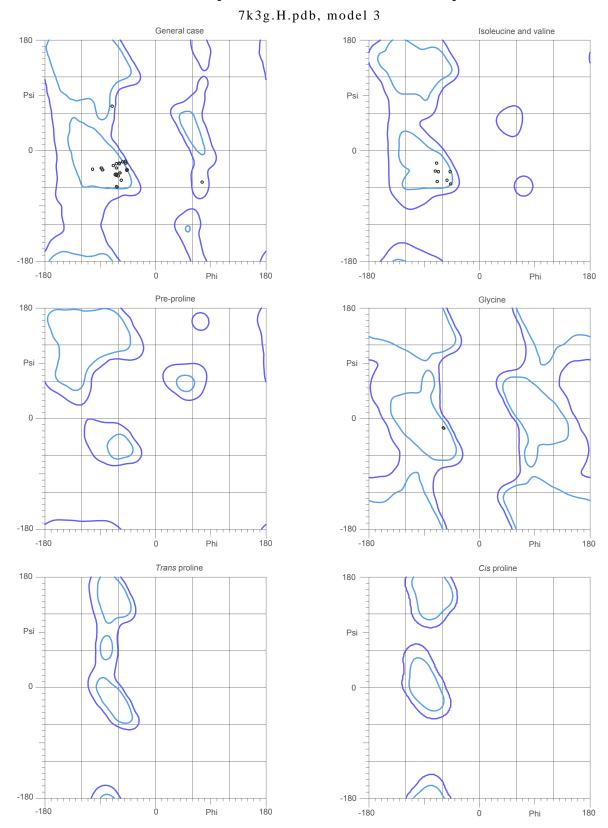
93.1% (135/145) of all residues were in favored (98%) regions. 96.6% (140/145) of all residues were in allowed (>99.8%) regions.

There were 5 outliers (phi, psi):

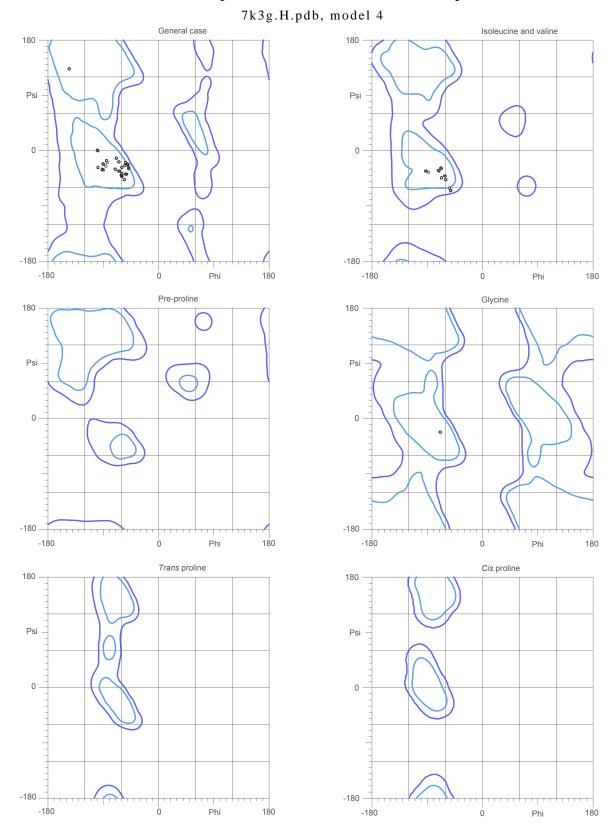
- [1] A 37 LEU (68.0, -78.1)
- [1] B 37 LEU (68.0, -78.0)
- [1] C 37 LEU (68.1, -78.1)
- [1] D 37 LEU (68.0, -78.1)
- [1] E 37 LEU (68.0, -78.1)



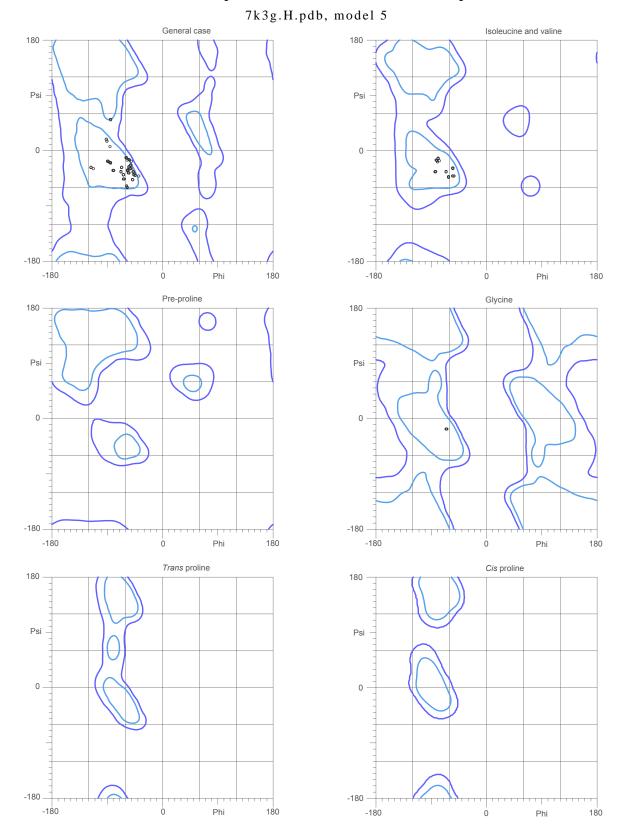
92.4%~(134/145) of all residues were in favored (98%) regions. 100.0%~(145/145) of all residues were in allowed (>99.8%) regions.



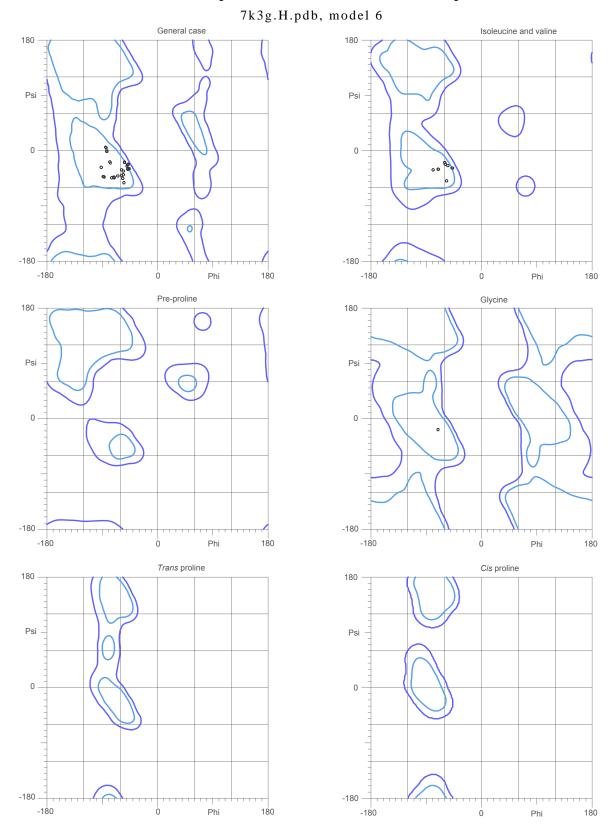
86.2% (125/145) of all residues were in favored (98%) regions. 100.0% (145/145) of all residues were in allowed (>99.8%) regions.



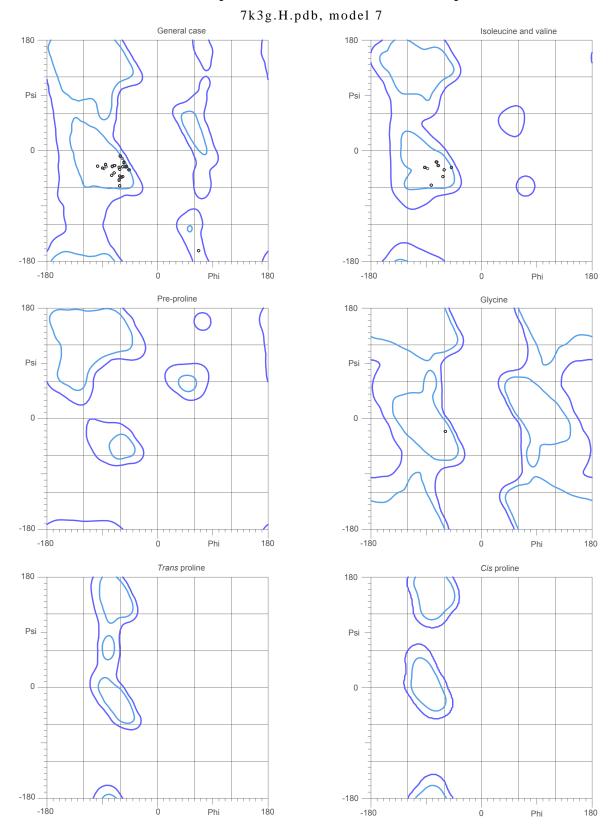
96.6% (140/145) of all residues were in favored (98%) regions. 100.0% (145/145) of all residues were in allowed (>99.8%) regions.



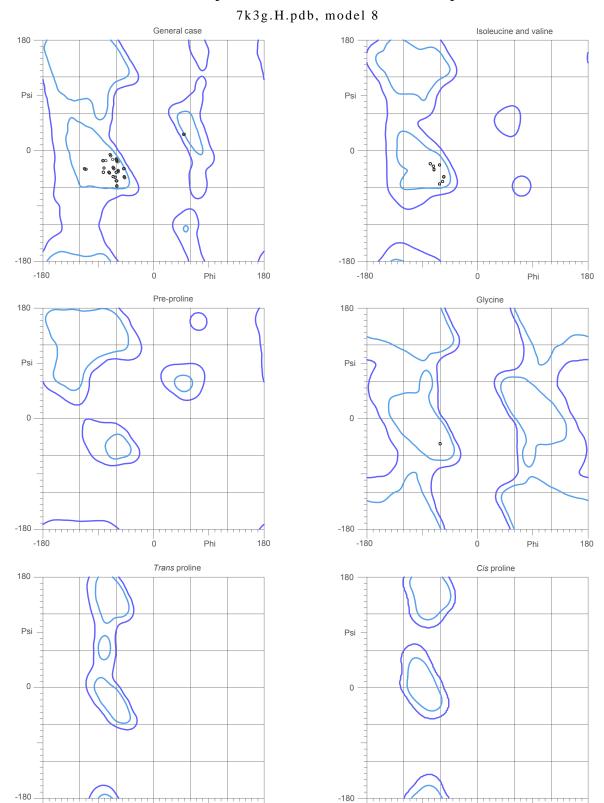
97.2%~(141/145) of all residues were in favored (98%) regions. 100.0%~(145/145) of all residues were in allowed (>99.8%) regions.



89.7%~(130/145) of all residues were in favored (98%) regions. 100.0%~(145/145) of all residues were in allowed (>99.8%) regions.



96.6% (140/145) of all residues were in favored (98%) regions. 100.0% (145/145) of all residues were in allowed (>99.8%) regions.



100.0% (145/145) of all residues were in favored (98%) regions. 100.0% (145/145) of all residues were in allowed (>99.8%) regions.

Phi

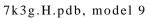
180

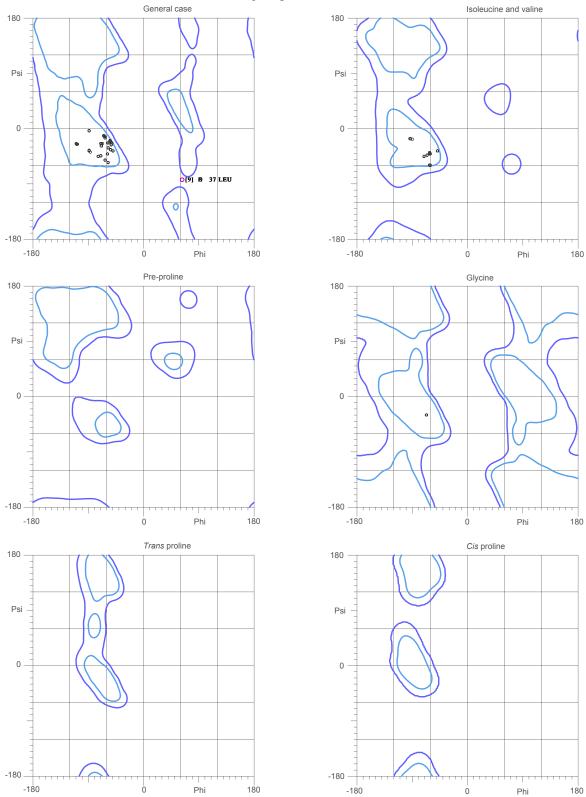
-180

There were no outliers.

-180

Phi





96.6% (140/145) of all residues were in favored (98%) regions. 96.6% (140/145) of all residues were in allowed (>99.8%) regions.

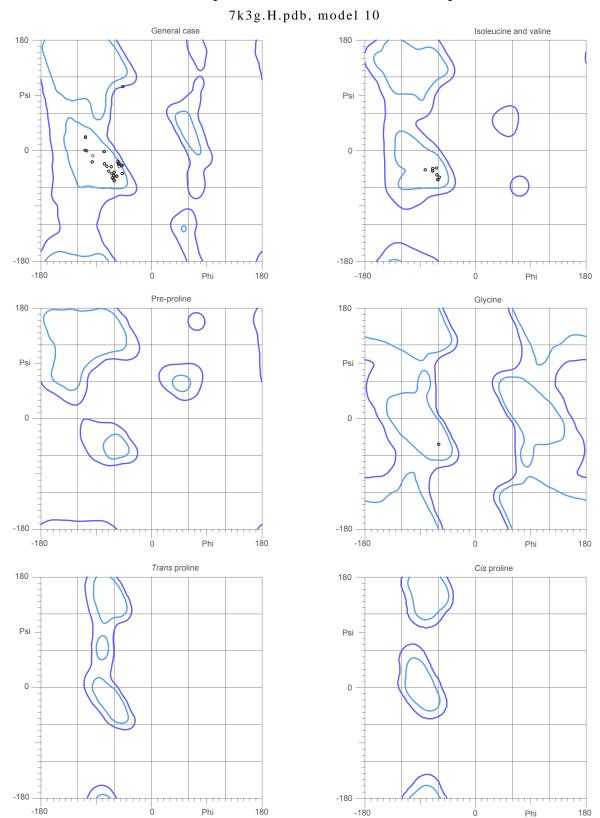
There were 5 outliers (phi, psi):

[9] A 37 LEU (63.4, -83.8)

[9] B 37 LEU (63.4, -83.8)

[9] C 37 LEU (63.5, -83.9) [9] D 37 LEU (63.3, -83.9)

[9] E 37 LEU (63.3, -83.9)



92.4% (134/145) of all residues were in favored (98%) regions. 100.0% (145/145) of all residues were in allowed (>99.8%) regions.