MolProbity Ramachandran analysis

1avgI.B99990001.pdb, model 1
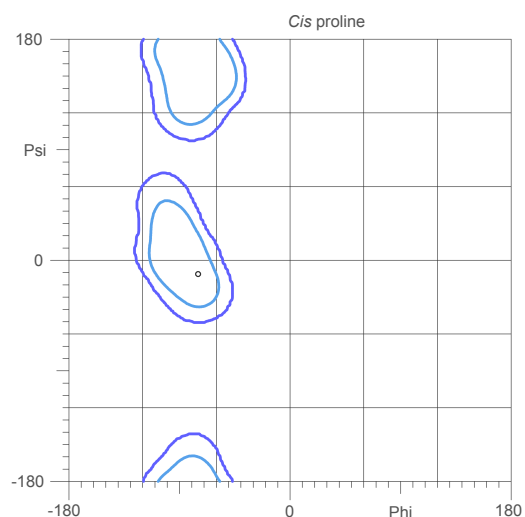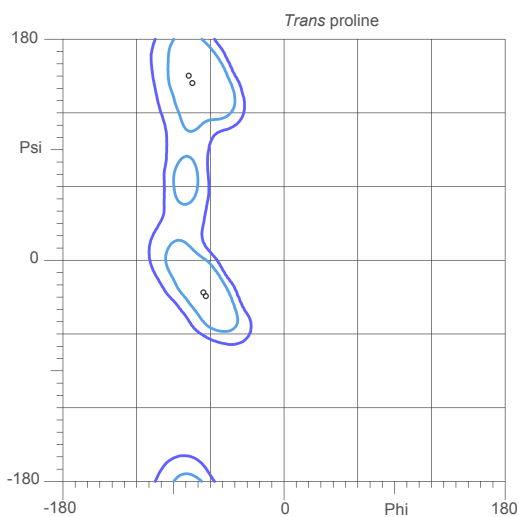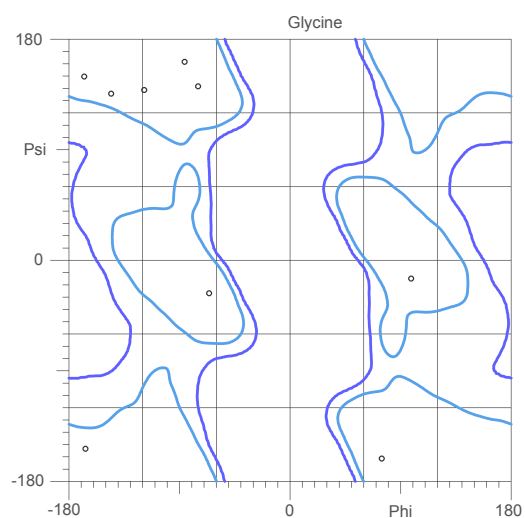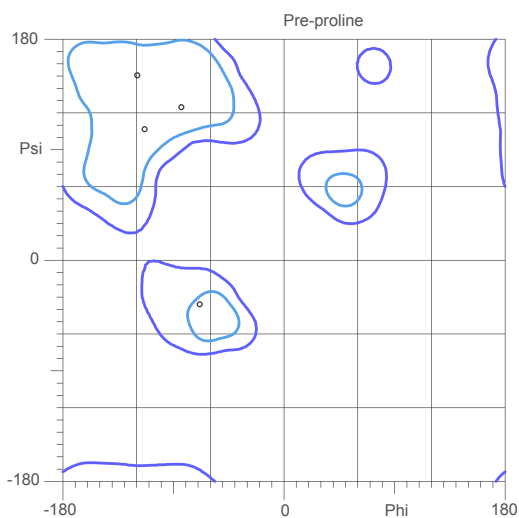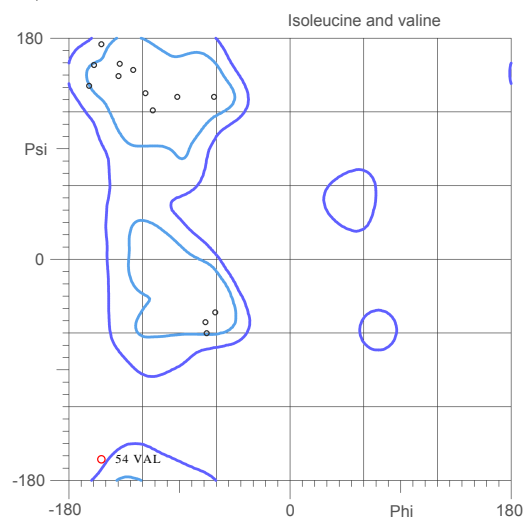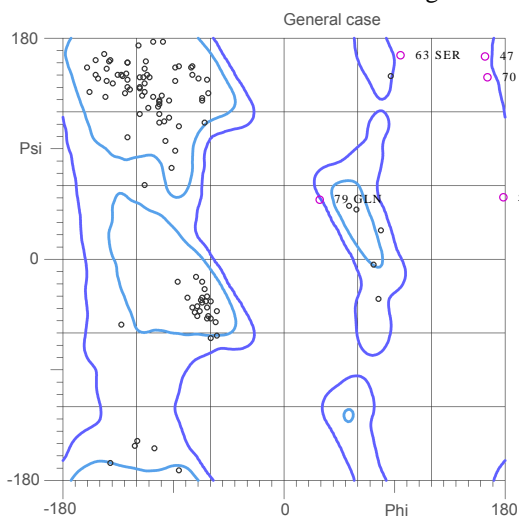

86,4% (121/140) of all residues were in favored (98%) regions.
95,7% (134/140) of all residues were in allowed (>99.8%) regions.

There were 6 outliers (phi, psi):
    47 SER (164,3, 166,2)
    51 THR (179,1, 51,8)
    54 VAL (-155,0, -163,1)
    63 SER (95,3, 167,3)
    70 ASN (166,3, 149,8)
    79 GLN (29,4, 49,1)

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Lovell, Davis, et al. Proteins 50:437 (2003)