A. PmDHFR

B. PoDHFR

Ramachandran Plot
new-entry

E5 (degrees)

-180 -135 -90 -45 0 45 90 135 180

Pll (degrees)

-180 -135 -90 -45 0 45 90 135 180

Plot statistics

Residues in most favored region (Gilk.): 295 0.813
Residues in additionally allowed region (Gilk. & Gyl.): 3 0.008
Residues in generously allowed region (Gilk. & Gyl.): 0 0.000
Residues in disallowed region: G 0.000
Residues in disallowed region: P 0.000
Residues of main chain and non-polar side chains 125 100.000
Number of residues in the class: - 125
Number of residues in the class: 0
Number of residues in the class: 125
Total number of residues: 125

Plot statistics

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Graphs represent 10% of samples of protein with less than 10% of residues in the most favored region.