

MolProbity Ramachandran analysis

6m17.H.pdb, model 1



89.7% (2746/3060) of all residues were in favored (90%) regions.
99.4% (3012/3060) of all residues were in allowed (99.9%) regions.

There were 48 outliers (phi, psi):

A 146 GLU (-45.5, -95.7)	E 483 VAL (60.9, -42.2)
A 147 PRO (-51.5, 88.8)	E 485 ILE (-59.5, -91.8)
C 108 HIS (71.5, 88.8)	F 420 PRO (-45.3, -10.0)
C 147 PRO (-51.5, 87.8)	F 421 GLY (151.1, 87.7)
D 394 GLU (-18.1, -105.8)	F 422 GLY (-44.0, -10.3)
D 397 GLY (-48.8, 31.5)	F 423 THR (161.9, 135.9)
D 401 VAL (-18.1, 140.0)	F 424 CYS (160.3, 108.6)
D 402 VAL (-28.8, 110.0)	F 483 VAL (60.9, -42.2)
D 403 LEU (-108.1, 8.0)	F 485 ILE (-59.5, -91.8)
D 404 LYS (164.9, 9.7)	
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