

MolProbity Ramachandran analysis

3d0h.H.pdb, model 1



Source: 3d0h.H.pdb, model 1
 Residue: A 289 PRO
 Chain: A
 Conformation: trans
 MolProbity score: 0.00
 Ramachandran plot: 0.00
 Clash score: 0.00
 Rotamer score: 0.00
 Overall score: 0.00