

MolProbity Ramachandran analysis

3d0h.H.pdb, model 1



Source: 3D0H.H.pdb, model 1
 Residue: A 289 PRO
 Psi: -170.0
 Phi: -150.0
 B-factor: 100.0
 Occupancy: 1.00
 Displacement: 0.00
 ... (repeated for all residues) ...