

# MolProbity Ramachandran analysis

3d0h.H.pdb, model 1



Source: 3d0h.H.pdb, model 1  
 Residue: 322, Chain: B, Type: ASN  
 Residue: 614, Chain: A, Type: ALA  
 Residue: 387, Chain: , Type: PHE  
 Residue: 471, Chain: E, Type: ALA  
 Residue: 416, Chain: E, Type: PHE  
 Residue: 416, Chain: F, Type: PHE  
 Residue: 370, Chain: E, Type: SER  
 Residue: 330, Chain: E, Type: ASN  
 Residue: 254, Chain: A, Type: SER  
 Residue: 409, Chain: A, Type: SER  
 Residue: 401, Chain: A, Type: TRP  
 Residue: 91, Chain: A, Type: LEU  
 Residue: 163, Chain: A, Type: TRP  
 Residue: 338, Chain: B, Type: ASN  
 Residue: 20, Chain: A, Type: THR  
 Residue: 338, Chain: B, Type: ASN  
 Residue: 338, Chain: A, Type: ASN