

# MolProbity Ramachandran analysis

3d0h.H.pdb, model 1



Source: 3D0H.H.pdb, model 1  
 Residue: A 20 THR  
 Residue: B 338 THR  
 Residue: A 163 TRP  
 Residue: A 91 LEU  
 Residue: A 401 ASP  
 Residue: A 409 SER  
 Residue: A 254 SER  
 Residue: E 330 ASN  
 Residue: E 370 SEI  
 Residue: 387 PHE  
 Residue: A 614 ALA  
 Residue: B 322 ASN  
 Residue: E 416 PHE  
 Residue: F 416 PHE  
 Residue: E 471 ALA  
 Residue: B 463 VAL  
 Residue: B 212 VAL  
 Residue: B 126 ILE  
 Residue: A 339 VAL  
 Residue: B 339 VAL  
 Residue: E 369 VAL  
 Residue: E 465 LYS  
 Residue: F 465 LYS  
 Residue: F 368 GLY  
 Residue: F 369 GLY  
 Residue: A 466 GLY  
 Residue: A 289 PRO  
 Residue: E 470 PRO  
 Residue: F 470 PRO