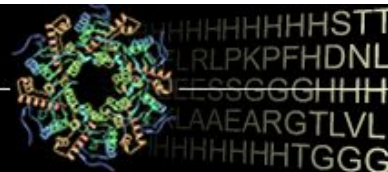
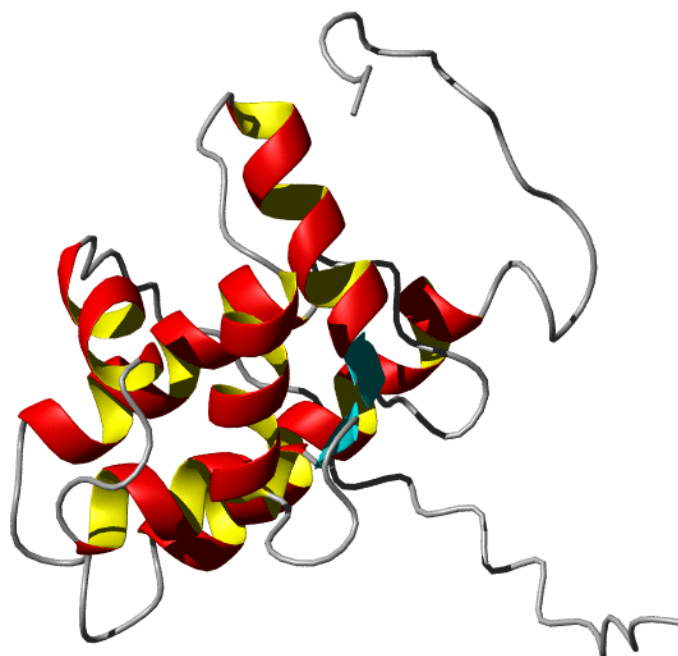


Center for Eukaryotic Structural Genomics

Protein Structure Initiative



Target ID	GO.33780	
Source Organism	<i>Homo sapiens</i>	
Target Name	BC009474	
PDB Entry	2G2B	Deposition: 15-Feb-2006
BMRB Entry	6980	Deposition: 28-Feb-2006
Function	human allograft inflammatory factor 1	
Produced From	Cell-free	
Structure by NMR	Restraints/Residue: 16.0	Subunits/Molecule: 1
	No. of Residues: 150	Molecular Weight: 17.0 kDa
	Backbone RMSD(16-35,38-40, 46-59, 70-80, 86-96,106-113, 121-125): 1.28 Å	All Heavy Atoms RMSD(16-35,38-40, 46-59, 70-80, 86-96,106-113, 121-125): 1.96 Å
Data Collected At	Nuclear Magnetic Resonance Facility at Madison (NMRFAM)	
Authors	Song, J., Tyler, R.C., Newman, C.L., Vinarov, D., Markley, J.L.	



Structural Features

NA

Percent Identity with Nearest PDB Structure at Time Solved 20% coverage (1CLM)

Pfam Cluster EFHAND,Tup
N,9644,1769,76953

Sequence Family Size 1010

Center for Eukaryotic Structural Genomics (CESG), University of Wisconsin-Madison Biochemistry Department, 433 Babcock Drive, Madison, WI 53706-1549; phone: 608.263.2183; fax: 608.890.1942; email: cesginfo@biochem.wisc.edu; website: <http://www.uwstructuralgenomics.org>. This research funded by NIH / NIGMS Protein Structure Initiative grants U54 GM074901 and P50 GM064598.