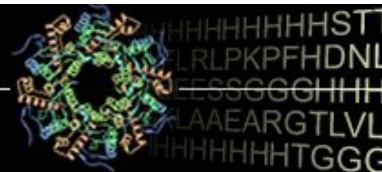
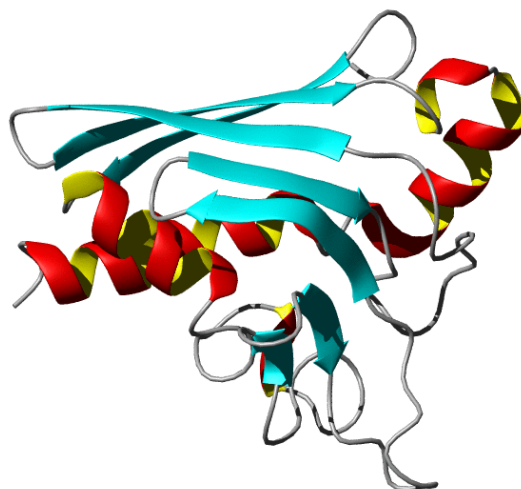


Center for Eukaryotic Structural Genomics

Protein Structure Initiative



Target ID	GO.79751	
Source Organism	<i>Micromonospora echinospora</i>	
Target Name	Q8KNFO	
PDB Entry	1ZXF	Deposition: 08-Jun-2005
Function	putative seed maturation protein-related	
Produced From	<i>E. coli</i>	
Structure by NMR	Restrains/Residue:	Subunits/Molecule: 1
	No. of Residues:	Molecular Weight: 7.5 kDa
	Backbone RMSD(10-64):	All Heavy Atoms RMSD(10-64):
Data Collected At	Nuclear Magnetic Resonance Facility at Madison (NMRFAM)	
Authors	Singh, S., Hager, M.H., Zhang, C., Griffith, B.R., Lee, M.S., Hallenga, K., Markley, J.L., Thorson, J.S.	



Structural Features

NA

Percent Identity with Nearest PDB Structure at Time Solved

Pfam Cluster

Protonet Cluster Size : Structures in PDB

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