








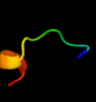

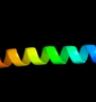





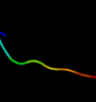

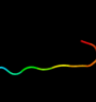


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	dlvkoa1	 Alignment		28.0	54	Fold: NAD(P)-binding Rossmann-fold domains Superfamily: NAD(P)-binding Rossmann-fold domains Family: Glyceraldehyde-3-phosphate dehydrogenase-like, N-terminal domain
2	d1p1ja1	 Alignment		20.3	54	Fold: NAD(P)-binding Rossmann-fold domains Superfamily: NAD(P)-binding Rossmann-fold domains Family: Glyceraldehyde-3-phosphate dehydrogenase-like, N-terminal domain
3	c1vkoaA	 Alignment		18.4	54	PDB header: isomerase Chain: A: PDB Molecule: inositol-3-phosphate synthase; PDBTitle: crystal structure of inositol-3-phosphate synthase (ce21227) from2 caenorhabditis elegans at 2.30 a resolution
4	d1pv0a	 Alignment		12.8	30	Fold: Long alpha-hairpin Superfamily: Sporulation inhibitor Sda Family: Sporulation inhibitor Sda
5	c1p1hD	 Alignment		11.8	54	PDB header: isomerase Chain: D: PDB Molecule: inositol-3-phosphate synthase; PDBTitle: crystal structure of the 1l-myo-inositol/nad+ complex
6	c3bj4B	 Alignment		11.6	37	PDB header: signaling protein Chain: B: PDB Molecule: potassium voltage-gated channel subfamily kqt PDBTitle: the kcnq1 (kv7.1) c-terminus, a multi-tiered scaffold for2 subunit assembly and protein interaction
7	d2j44a2	 Alignment		11.2	30	Fold: Prealbumin-like Superfamily: Starch-binding domain-like Family: PUD-like
8	c2lfeA	 Alignment		11.2	34	PDB header: ligase Chain: A: PDB Molecule: e3 ubiquitin-protein ligase hecw2; PDBTitle: solution nmr structure of n-terminal domain of human e3 ubiquitin-2 protein ligase hecw2, northeast structural genomics consortium (nesg)3 target ht6306a
9	c2le2B	 Alignment		9.0	78	PDB header: hydrolase inhibitor Chain: B: PDB Molecule: p56; PDBTitle: novel dimeric structure of phage phi29-encoded protein p56: insights2 into uracil-dna glycosylase inhibition
10	d2j43a2	 Alignment		7.4	27	Fold: Prealbumin-like Superfamily: Starch-binding domain-like Family: PUD-like

11 [c2qbxB](#)

Alignment



6.6

37

PDB header:signaling protein
Chain: B: **PDB Molecule:**ephrin type-b receptor 2;
PDBTitle: ephb2/snew antagonistic peptide complex