

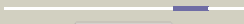
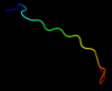

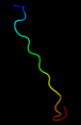










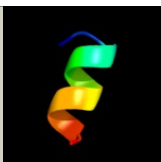
Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">d1n7va_</a>	 Alignment		17.9	24	<b>Fold:</b> Adsorption protein p2 <b>Superfamily:</b> Adsorption protein p2 <b>Family:</b> Adsorption protein p2
2	<a href="#">c3kysC_</a>	 Alignment		10.1	35	<b>PDB header:</b> transcription/protein binding <b>Chain:</b> C: <b>PDB Molecule:</b> transcriptional enhancer factor tef-1; <b>PDBTitle:</b> crystal structure of human yap and tead complex
3	<a href="#">c3l15A_</a>	 Alignment		9.2	25	<b>PDB header:</b> transcription <b>Chain:</b> A: <b>PDB Molecule:</b> transcriptional enhancer factor tef-4; <b>PDBTitle:</b> human tead2 transcriptional factor
4	<a href="#">d2bbga_</a>	 Alignment		7.3	21	<b>Fold:</b> Amb V allergen <b>Superfamily:</b> Amb V allergen <b>Family:</b> Amb V allergen
5	<a href="#">c2cpbA_</a>	 Alignment		7.0	8	<b>PDB header:</b> viral protein <b>Chain:</b> A: <b>PDB Molecule:</b> m13 major coat protein; <b>PDBTitle:</b> solution nmr structures of the major coat protein of2 filamentous bacteriophage m13 solubilized in3 dodecylphosphocholine micelles, 25 lowest energy structures
6	<a href="#">c2kncA_</a>	 Alignment		6.8	17	<b>PDB header:</b> cell adhesion <b>Chain:</b> A: <b>PDB Molecule:</b> integrin alpha-iiB; <b>PDBTitle:</b> platelet integrin alfaIiB-beta3 transmembrane-cytoplasmic2 heterocomplex
7	<a href="#">c3lu2B_</a>	 Alignment		5.7	36	<b>PDB header:</b> hydrolase <b>Chain:</b> B: <b>PDB Molecule:</b> Imo2462 protein; <b>PDBTitle:</b> structure of Imo2462, a listeria monocytogenes amidohydrolase family2 putative dipeptidase

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[d1itua\\_](#)

Alignment



5.2

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**Fold:** TIM beta/alpha-barrel

**Superfamily:** Metallo-dependent hydrolases

**Family:** Renal dipeptidase