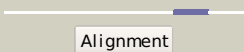

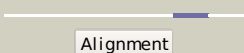

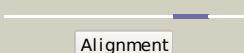

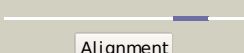

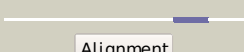

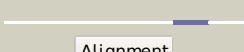

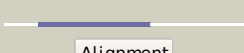

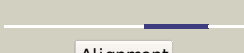





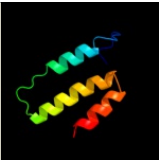


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">d1uptd_</a>	 Alignment		19.7	22	<b>Fold:</b> GRIP domain <b>Superfamily:</b> GRIP domain <b>Family:</b> GRIP domain
2	<a href="#">c2j5dA_</a>	 Alignment		18.8	22	<b>PDB header:</b> membrane protein <b>Chain:</b> A: <b>PDB Molecule:</b> bcl2/adenovirus e1b 19 kda protein-interacting <b>PDBTitle:</b> nmr structure of bnip3 transmembrane domain in lipid2 bicelles
3	<a href="#">c2ka2B_</a>	 Alignment		18.0	23	<b>PDB header:</b> membrane protein <b>Chain:</b> B: <b>PDB Molecule:</b> bcl2/adenovirus e1b 19 kda protein-interacting <b>PDBTitle:</b> solution nmr structure of bnip3 transmembrane peptide dimer2 in detergent micelles with his173-ser172 intermonomer3 hydrogen bond restraints
4	<a href="#">c2ka1A_</a>	 Alignment		18.0	23	<b>PDB header:</b> membrane protein <b>Chain:</b> A: <b>PDB Molecule:</b> bcl2/adenovirus e1b 19 kda protein-interacting <b>PDBTitle:</b> solution nmr structure of bnip3 transmembrane peptide dimer2 in detergent micelles
5	<a href="#">c2ka2A_</a>	 Alignment		18.0	23	<b>PDB header:</b> membrane protein <b>Chain:</b> A: <b>PDB Molecule:</b> bcl2/adenovirus e1b 19 kda protein-interacting <b>PDBTitle:</b> solution nmr structure of bnip3 transmembrane peptide dimer2 in detergent micelles with his173-ser172 intermonomer3 hydrogen bond restraints
6	<a href="#">c2ka1B_</a>	 Alignment		18.0	23	<b>PDB header:</b> membrane protein <b>Chain:</b> B: <b>PDB Molecule:</b> bcl2/adenovirus e1b 19 kda protein-interacting <b>PDBTitle:</b> solution nmr structure of bnip3 transmembrane peptide dimer2 in detergent micelles
7	<a href="#">c3rkoK_</a>	 Alignment		16.0	9	<b>PDB header:</b> oxidoreductase <b>Chain:</b> K: <b>PDB Molecule:</b> nadh-quinone oxidoreductase subunit k; <b>PDBTitle:</b> crystal structure of the membrane domain of respiratory complex i from2 e. coli at 3.0 angstrom resolution
8	<a href="#">c3fewX_</a>	 Alignment		6.6	19	<b>PDB header:</b> immune system <b>Chain:</b> X: <b>PDB Molecule:</b> colicin s4; <b>PDBTitle:</b> structure and function of colicin s4, a colicin with a2 duplicated receptor binding domain
9	<a href="#">c3hfxA_</a>	 Alignment		6.4	6	<b>PDB header:</b> transport protein <b>Chain:</b> A: <b>PDB Molecule:</b> l-carnitine/gamma-butyrobetaine antiporter; <b>PDBTitle:</b> crystal structure of carnitine transporter
10	<a href="#">d1a87a_</a>	 Alignment		6.1	15	<b>Fold:</b> Toxins' membrane translocation domains <b>Superfamily:</b> Colicin <b>Family:</b> Colicin

11 [c1a87A](#)

Alignment



6.1

15

**PDB header:**bacteriocin  
**Chain:** A: **PDB Molecule:**colicin n;  
**PDBTitle:** colicin n