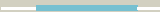




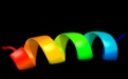









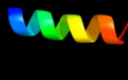

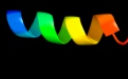




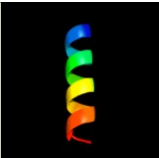
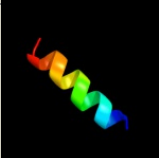
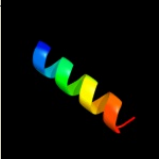

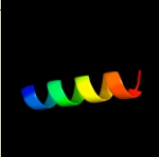
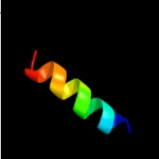


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c1dfwA_	 Alignment		37.7	58	PDB header: immune system Chain: A: PDB Molecule: lung surfactant protein b; PDBTitle: conformational mapping of the n-terminal segment of 2 surfactant protein b in lipid using 13c-enhanced fourier3 transform infrared spectroscopy (ftir)
2	c1sszA_	 Alignment		22.8	58	PDB header: surface active protein Chain: A: PDB Molecule: pulmonary surfactant-associated protein b; PDBTitle: conformational mapping of mini-b: an n-terminal/c-terminal2 construct of surfactant protein b using 13c-enhanced3 fourier transform infrared (ftir) spectroscopy
3	c3lhwC_	 Alignment		20.6	56	PDB header: transport protein Chain: C: PDB Molecule: m2 protein; PDBTitle: high resolution crystal structure of transmembrane domain of m2
4	c3lhwB_	 Alignment		20.6	56	PDB header: transport protein Chain: B: PDB Molecule: m2 protein; PDBTitle: high resolution crystal structure of transmembrane domain of m2
5	c3lhwD_	 Alignment		20.6	56	PDB header: transport protein Chain: D: PDB Molecule: m2 protein; PDBTitle: high resolution crystal structure of transmembrane domain of m2
6	c3lhwA_	 Alignment		20.6	56	PDB header: transport protein Chain: A: PDB Molecule: m2 protein; PDBTitle: high resolution crystal structure of transmembrane domain of m2
7	c3c9jC_	 Alignment		16.3	56	PDB header: membrane protein Chain: C: PDB Molecule: proton channel protein m2, transmembrane segment; PDBTitle: the crystal structure of transmembrane domain of m2 protein and2 amantadine complex
8	c3c9jB_	 Alignment		16.3	56	PDB header: membrane protein Chain: B: PDB Molecule: proton channel protein m2, transmembrane segment; PDBTitle: the crystal structure of transmembrane domain of m2 protein and2 amantadine complex
9	c3c9jA_	 Alignment		15.7	56	PDB header: membrane protein Chain: A: PDB Molecule: proton channel protein m2, transmembrane segment; PDBTitle: the crystal structure of transmembrane domain of m2 protein and2 amantadine complex
10	c3c9jD_	 Alignment		14.8	56	PDB header: membrane protein Chain: D: PDB Molecule: proton channel protein m2, transmembrane segment; PDBTitle: the crystal structure of transmembrane domain of m2 protein and2 amantadine complex
11	c1nyjD_	 Alignment		6.3	50	PDB header: viral protein Chain: D: PDB Molecule: matrix protein m2; PDBTitle: the closed state structure of m2 protein h+ channel by2 solid state nmr spectroscopy

12	c1nyjB_	Alignment		6.3	50	PDB header: viral protein Chain: B: PDB Molecule: matrix protein m2; PDBTitle: the closed state structure of m2 protein h+ channel by2 solid state nmr spectroscopy
13	c2kqtB_	Alignment		6.3	50	PDB header: transport protein Chain: B: PDB Molecule: m2 protein; PDBTitle: solid-state nmr structure of the m2 transmembrane peptide of the2 influenza a virus in dmpe lipid bilayers bound to deuterated3 amantadine
14	c1mp6A_	Alignment		6.3	50	PDB header: membrane protein Chain: A: PDB Molecule: matrix protein m2; PDBTitle: structure of the transmembrane region of the m2 protein h+2 channel by solid state nmr spectroscopy
15	c1nyjA_	Alignment		6.3	50	PDB header: viral protein Chain: A: PDB Molecule: matrix protein m2; PDBTitle: the closed state structure of m2 protein h+ channel by2 solid state nmr spectroscopy
16	c1nyjC_	Alignment		6.3	50	PDB header: viral protein Chain: C: PDB Molecule: matrix protein m2; PDBTitle: the closed state structure of m2 protein h+ channel by2 solid state nmr spectroscopy
17	c2kqtD_	Alignment		6.3	50	PDB header: transport protein Chain: D: PDB Molecule: m2 protein; PDBTitle: solid-state nmr structure of the m2 transmembrane peptide of the2 influenza a virus in dmpe lipid bilayers bound to deuterated3 amantadine

18 [c2kqtC](#)

Alignment



6.3

50

PDB header:transport protein
Chain: C: **PDB Molecule:**m2 protein;
PDBTitle: solid-state nmr structure of the m2 transmembrane peptide of the2 influenza a virus in dmpe lipid bilayers bound to deuterated3 amantadine