
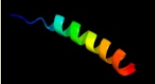
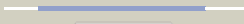




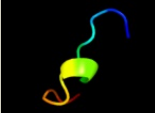





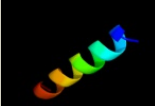
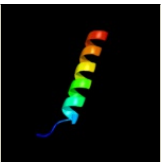


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2kncA_	 Alignment		41.5	16	PDB header: cell adhesion Chain: A: PDB Molecule: integrin alpha-iiB; PDBTitle: platelet integrin alphaIIb-beta3 transmembrane-cytoplasmic2 heterocomplex
2	c2b2hA_	 Alignment		22.0	17	PDB header: transport protein Chain: A: PDB Molecule: ammonium transporter; PDBTitle: ammonium transporter amt-1 from a. fulgidus (as)
3	c2d68A_	 Alignment		20.7	12	PDB header: cell cycle Chain: A: PDB Molecule: fop; PDBTitle: structure of the n-terminal domain of fop (fgfr1op) protein
4	c2e2zA_	 Alignment		20.4	8	PDB header: protein transport, chaperone regulator Chain: A: PDB Molecule: tim15; PDBTitle: solution nmr structure of yeast tim15, co-chaperone of2 mitochondrial hsp70
5	c3hd6A_	 Alignment		17.7	16	PDB header: membrane protein, transport protein Chain: A: PDB Molecule: ammonium transporter rh type c; PDBTitle: crystal structure of the human rhesus glycoprotein rhcg
6	d2ptza2	 Alignment		14.6	15	Fold: Enolase N-terminal domain-like Superfamily: Enolase N-terminal domain-like Family: Enolase N-terminal domain-like
7	c2jp3A_	 Alignment		7.1	28	PDB header: transcription Chain: A: PDB Molecule: fxyd domain-containing ion transport regulator 4; PDBTitle: solution structure of the human fxyd4 (chif) protein in sds2 micelles

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[c2k1aA](#)

Alignment



6.5

16

PDB header:cell adhesion
Chain: A: **PDB Molecule:**integrin alpha-iiB;
PDBTitle: bicelle-embedded integrin alpha(iiB) transmembrane segment