
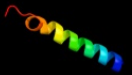
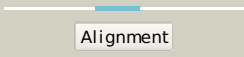

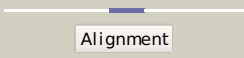
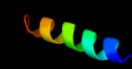
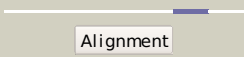
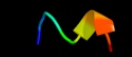
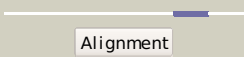

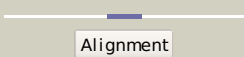

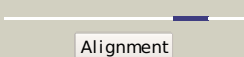

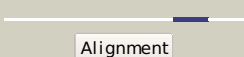



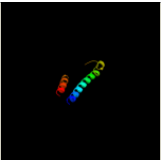


#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">c2k1aA_</a>	 Alignment		39.0	24	<b>PDB header:</b> cell adhesion <b>Chain:</b> A: <b>PDB Molecule:</b> integrin alpha-iiB; <b>PDBTitle:</b> bicelle-embedded integrin alpha(iiB) transmembrane segment
2	<a href="#">c2kncA_</a>	 Alignment		38.5	24	<b>PDB header:</b> cell adhesion <b>Chain:</b> A: <b>PDB Molecule:</b> integrin alpha-iiB; <b>PDBTitle:</b> platelet integrin alphaIIb-beta3 transmembrane-cytoplasmic2 heterocomplex
3	<a href="#">c2bbjB_</a>	 Alignment		14.4	16	<b>PDB header:</b> metal transport/membrane protein <b>Chain:</b> B: <b>PDB Molecule:</b> divalent cation transport-related protein; <b>PDBTitle:</b> crystal structure of the cora mg2+ transporter
4	<a href="#">c2jpnA_</a>	 Alignment		13.5	50	<b>PDB header:</b> hydrolase <b>Chain:</b> A: <b>PDB Molecule:</b> atp-dependent dna helicase uvsw; <b>PDBTitle:</b> solution structure of t4 bacteriophage helicase uvsw.1
5	<a href="#">c2lcyA_</a>	 Alignment		13.5	60	<b>PDB header:</b> viral protein <b>Chain:</b> A: <b>PDB Molecule:</b> virion spike glycoprotein; <b>PDBTitle:</b> nmr structure of the complete internal fusion loop from ebolavirus gp22 at ph 5.5
6	<a href="#">d2iuba2</a>	 Alignment		12.9	14	<b>Fold:</b> Transmembrane helix hairpin <b>Superfamily:</b> Magnesium transport protein CorA, transmembrane region <b>Family:</b> Magnesium transport protein CorA, transmembrane region
7	<a href="#">c3csyl_</a>	 Alignment		8.6	60	<b>PDB header:</b> immune system/viral protein <b>Chain:</b> J: <b>PDB Molecule:</b> envelope glycoprotein gp2; <b>PDBTitle:</b> crystal structure of the trimeric prefusion ebola virus glycoprotein2 in complex with a neutralizing antibody from a human survivor
8	<a href="#">c3s88J_</a>	 Alignment		7.9	60	<b>PDB header:</b> immune system/viral protein <b>Chain:</b> J: <b>PDB Molecule:</b> envelope glycoprotein; <b>PDBTitle:</b> crystal structure of sudan ebolavirus glycoprotein (strain gulu) bound2 to 16f6
9	<a href="#">d1wixa_</a>	 Alignment		6.6	27	<b>Fold:</b> CH domain-like <b>Superfamily:</b> Hook domain <b>Family:</b> Hook domain

10	<a href="#">d3ehbb2</a>	Alignment		6.0	23	<b>Fold:</b> Transmembrane helix hairpin <b>Superfamily:</b> Cytochrome c oxidase subunit II-like, transmembrane region <b>Family:</b> Cytochrome c oxidase subunit II-like, transmembrane region
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