
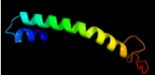



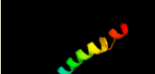







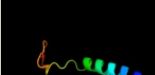








#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	<a href="#">c3qngD_</a>	 Alignment		65.9	11	<b>PDB header:</b> membrane protein, transport protein <b>Chain:</b> D: <b>PDB Molecule:</b> pts system, cellobiose-specific iic component; <b>PDBTitle:</b> crystal structure of the transporter chbc, the iic component from the 2 n,n'-diacetylchitobiose-specific phosphotransferase system
2	<a href="#">c3iz5Q_</a>	 Alignment		40.0	29	<b>PDB header:</b> ribosome <b>Chain:</b> Q: <b>PDB Molecule:</b> 60s ribosomal protein l5 (l18p); <b>PDBTitle:</b> localization of the large subunit ribosomal proteins into a 5.5 a2 cryo-em map of triticum aestivum translating 80s ribosome
3	<a href="#">c2l34B_</a>	 Alignment		37.3	30	<b>PDB header:</b> protein binding <b>Chain:</b> B: <b>PDB Molecule:</b> tyro protein tyrosine kinase-binding protein; <b>PDBTitle:</b> structure of the dap12 transmembrane homodimer
4	<a href="#">c2l35B_</a>	 Alignment		34.6	30	<b>PDB header:</b> protein binding <b>Chain:</b> B: <b>PDB Molecule:</b> tyro protein tyrosine kinase-binding protein; <b>PDBTitle:</b> structure of the dap12-nkg2c transmembrane heterotrimer
5	<a href="#">c2l34A_</a>	 Alignment		27.1	41	<b>PDB header:</b> protein binding <b>Chain:</b> A: <b>PDB Molecule:</b> tyro protein tyrosine kinase-binding protein; <b>PDBTitle:</b> structure of the dap12 transmembrane homodimer
6	<a href="#">c2l35A_</a>	 Alignment		19.2	36	<b>PDB header:</b> protein binding <b>Chain:</b> A: <b>PDB Molecule:</b> dap12-nkg2c tm; <b>PDBTitle:</b> structure of the dap12-nkg2c transmembrane heterotrimer
7	<a href="#">d1kqfb2</a>	 Alignment		14.9	28	<b>Fold:</b> Single transmembrane helix <b>Superfamily:</b> Iron-sulfur subunit of formate dehydrogenase N, transmembrane anchor <b>Family:</b> Iron-sulfur subunit of formate dehydrogenase N, transmembrane anchor
8	<a href="#">c1ciiA_</a>	 Alignment		14.5	14	<b>PDB header:</b> transmembrane protein <b>Chain:</b> A: <b>PDB Molecule:</b> colicin ia; <b>PDBTitle:</b> colicin ia
9	<a href="#">c2kncA_</a>	 Alignment		7.4	20	<b>PDB header:</b> cell adhesion <b>Chain:</b> A: <b>PDB Molecule:</b> integrin alpha-iiB; <b>PDBTitle:</b> platelet integrin alpha-iiB-beta3 transmembrane-cytoplasmic2 heterocomplex
10	<a href="#">d1r7ma1</a>	 Alignment		6.7	29	<b>Fold:</b> Homing endonuclease-like <b>Superfamily:</b> Homing endonucleases <b>Family:</b> Group I mobile intron endonuclease

11 [c1r7mA](#)

Alignment



5.7

29

**PDB header:**hydrolase/dna  
**Chain:** A: **PDB Molecule:**intron-encoded endonuclease i-scei;  
**PDBTitle:** the homing endonuclease i-scei bound to its dna recognition2 region