
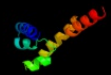
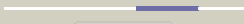


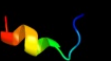



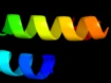








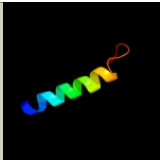
Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d1kpla_	 Alignment		12.6	15	Fold: Clc chloride channel Superfamily: Clc chloride channel Family: Clc chloride channel
2	d1rh1a2	 Alignment		11.6	14	Fold: Toxins' membrane translocation domains Superfamily: Colicin Family: Colicin
3	d1dxsa_	 Alignment		11.0	21	Fold: SAM domain-like Superfamily: SAM/Pointed domain Family: SAM (sterile alpha motif) domain
4	d2o34a1	 Alignment		10.7	7	Fold: T-fold Superfamily: ApbE-like Family: DVU1097-like
5	c2uwjG_	 Alignment		9.2	3	PDB header: chaperone Chain: G: PDB Molecule: type iii export protein pscg; PDBTitle: structure of the heterotrimeric complex which regulates2 type iii secretion needle formation
6	c2ju0B_	 Alignment		8.4	56	PDB header: metal binding protein/signaling protein Chain: B: PDB Molecule: phosphatidylinositol 4-kinase pik1; PDBTitle: structure of yeast frequenin bound to pdtins 4-kinase
7	c3fewX_	 Alignment		7.0	15	PDB header: immune system Chain: X: PDB Molecule: colicin s4; PDBTitle: structure and function of colicin s4, a colicin with a2 duplicated receptor binding domain
8	d1aa7a_	 Alignment		6.3	11	Fold: Influenza virus matrix protein M1 Superfamily: Influenza virus matrix protein M1 Family: Influenza virus matrix protein M1

9

[dlc7qa_](#)

Alignment



6.2

20

Fold:SIS domain

Superfamily:SIS domain

Family:Phosphoglucose isomerase, PGI