
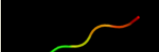
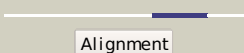

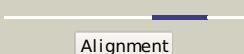



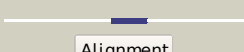

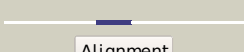
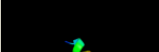






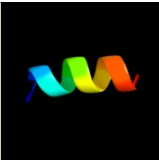


| # | Template | Alignment Coverage | 3D Model | Confidence | % i.d. | Template Information |
|---|-------------------------|--|---|------------|--------|---|
| 1 | c2jx5A_ |  Alignment |  | 9.8 | 38 | PDB header: ribosomal protein Chain: A: PDB Molecule: glub(s27a); PDBTitle: solution structure of the ubiquitin domain n-terminal to2 the s27a ribosomal subunit of giardia lamblia |
| 2 | c2y69Z_ |  Alignment |  | 9.0 | 23 | PDB header: electron transport Chain: Z: PDB Molecule: cytochrome c oxidase polypeptide 8h; PDBTitle: bovine heart cytochrome c oxidase re-refined with molecular2 oxygen |
| 3 | d1v54m_ |  Alignment |  | 8.8 | 23 | Fold: Single transmembrane helix Superfamily: Mitochondrial cytochrome c oxidase subunit VIIIb (aka IX) Family: Mitochondrial cytochrome c oxidase subunit VIIIb (aka IX) |
| 4 | c3nzzA_ |  Alignment |  | 6.9 | 55 | PDB header: cell invasion Chain: A: PDB Molecule: cell invasion protein sipd; PDBTitle: crystal structure of the salmonella type iii secretion system tip2 protein sipd |
| 5 | c1g2cN_ |  Alignment |  | 6.1 | 47 | PDB header: viral protein Chain: N: PDB Molecule: fusion protein (f); PDBTitle: human respiratory syncytial virus fusion protein core |
| 6 | d1sdia_ |  Alignment |  | 5.2 | 27 | Fold: YcfC-like Superfamily: YcfC-like Family: YcfC-like |
| 7 | c2gohA_ |  Alignment |  | 5.0 | 54 | PDB header: viral protein Chain: A: PDB Molecule: vpu protein; PDBTitle: three-dimensional structure of the trans-membrane domain of2 vpu from hiv-1 in aligned phospholipid bicelles |
| 8 | c2gofA_ |  Alignment |  | 5.0 | 54 | PDB header: viral protein Chain: A: PDB Molecule: vpu protein; PDBTitle: three-dimensional structure of the trans-membrane domain of2 vpu from hiv-1 in aligned phospholipid bicelles |
| 9 | c1pjeA_ |  Alignment |  | 5.0 | 54 | PDB header: viral protein Chain: A: PDB Molecule: vpu protein; PDBTitle: structure of the channel-forming trans-membrane domain of2 virus protein "u"(vpu) from hiv-1 |

10 [c1pi7A_](#)

Alignment



5.0

54

PDB header:viral protein
Chain: A: **PDB Molecule:**vpu protein;
PDBTitle: structure of the channel-forming trans-membrane domain of2 virus protein "u" (vpu) from hiv-1