
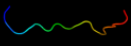








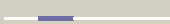
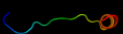


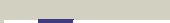








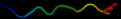





Detailed template information

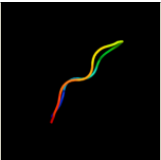
#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2hzqA_	 Alignment		17.6	33	PDB header: transport protein Chain: A: PDB Molecule: apolipoprotein d; PDBTitle: crystal structure of human apolipoprotein d (apod) in2 complex with progesterone
2	c1vd3A_	 Alignment		14.2	40	PDB header: hydrolase Chain: A: PDB Molecule: rnase ngr3; PDBTitle: ribonuclease nt in complex with 2'-ump
3	d1iyba_	 Alignment		13.8	37	Fold: Ribonuclease Rh-like Superfamily: Ribonuclease Rh-like Family: Ribonuclease Rh-like
4	c2pqxA_	 Alignment		11.2	17	PDB header: hydrolase Chain: A: PDB Molecule: ribonuclease i; PDBTitle: e. coli rnase 1 (in vivo folded)
5	d1kxoA_	 Alignment		10.5	27	Fold: Lipocalins Superfamily: Lipocalins Family: Retinol binding protein-like
6	d1bbpA_	 Alignment		10.2	26	Fold: Lipocalins Superfamily: Lipocalins Family: Retinol binding protein-like
7	d1dixa_	 Alignment		9.1	30	Fold: Ribonuclease Rh-like Superfamily: Ribonuclease Rh-like Family: Ribonuclease Rh-like
8	d1n0sA_	 Alignment		8.4	29	Fold: Lipocalins Superfamily: Lipocalins Family: Retinol binding protein-like
9	d1sgla_	 Alignment		8.1	23	Fold: Ribonuclease Rh-like Superfamily: Ribonuclease Rh-like Family: Ribonuclease Rh-like
10	d1jy5a_	 Alignment		7.5	23	Fold: Ribonuclease Rh-like Superfamily: Ribonuclease Rh-like Family: Ribonuclease Rh-like
11	d1z24a1	 Alignment		7.4	18	Fold: Lipocalins Superfamily: Lipocalins Family: Retinol binding protein-like

12	d1sg1x2	Alignment		6.9	46	Fold: TNF receptor-like Superfamily: TNF receptor-like Family: TNF receptor-like
13	d1gkab_	Alignment		5.9	31	Fold: Lipocalins Superfamily: Lipocalins Family: Retinol binding protein-like
14	d1ucda_	Alignment		5.9	30	Fold: Ribonuclease Rh-like Superfamily: Ribonuclease Rh-like Family: Ribonuclease Rh-like
15	c3d3zA_	Alignment		5.6	27	PDB header: hydrolase Chain: A: PDB Molecule: actibind; PDBTitle: crystal structure of actibind a t2 rnase
16	d2a0sa1	Alignment		5.5	25	Fold: T-fold Superfamily: Tetrahydrobiopterin biosynthesis enzymes-like Family: 6-pyruvoyl tetrahydropterin synthase

17

[dly13a_](#)

Alignment



5.5

25

Fold:T-fold
Superfamily:Tetrahydrobiopterin biosynthesis enzymes-like
Family:6-pyruvoyl tetrahydropterin synthase