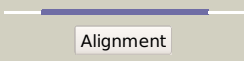
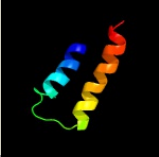
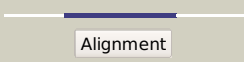
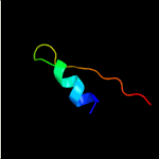
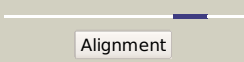
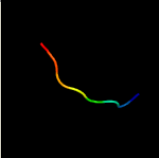
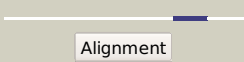
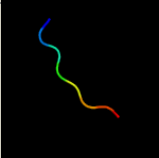


Phyre2

Email	mdejesus@rockefeller.edu
Description	RVBD2452c_(-)_2752858_2753004
Date	Wed Aug 7 12:50:07 BST 2019
Unique Job ID	9d9e494f79d7409a

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c2kk1A_	 Alignment		11.9	26	PDB header: transferase Chain: A: PDB Molecule: tyrosine-protein kinase abl2; PDBTitle: solution structure of c-terminal domain of tyrosine-protein2 kinase abl2 from homo sapiens, northeast structural3 genomics consortium (nesg) target hr5537a
2	c6cp8B_	 Alignment		7.8	42	PDB header: toxin/antitoxin Chain: B: PDB Molecule: cdia; PDBTitle: contact-dependent growth inhibition toxin-immunity protein complex2 from from e. coli 3006
3	c1pyuD_	 Alignment		6.5	50	PDB header: lyase Chain: D: PDB Molecule: aspartate 1-decarboxylase alfa chain; PDBTitle: processed aspartate decarboxylase mutant with ser25 mutated to cys
4	c1vc3B_	 Alignment		6.3	75	PDB header: lyase Chain: B: PDB Molecule: l-aspartate-alpha-decarboxylase heavy chain; PDBTitle: crystal structure of l-aspartate-alpha-decarboxylase

5

[c1uheA](#)

Alignment



6.0

50

PDB header: lyase
Chain: A: **PDB Molecule:** aspartate 1-decarboxylase alpha chain;
PDB Title: crystal structure of aspartate decarboxylase, isoasparagine complex