
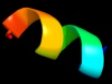





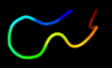





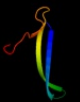




Phyre2

Email	l.a.kelley@imperial.ac.uk
Description	P0AB28
Date	Thu Jan 5 11:14:28 GMT 2012
Unique Job ID	aee7d6a36f139a37

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	c3owtC_	 Alignment		30.5	60	PDB header: protein binding Chain: C: PDB Molecule: regulatory protein sir3; PDBTitle: crystal structure of s. cerevisiae rap1-sir3 complex
2	c3s32A_	 Alignment		15.1	47	PDB header: transcription Chain: A: PDB Molecule: set1/ash2 histone methyltransferase complex subunit ash2; PDBTitle: crystal structure of ash2l n-terminal domain
3	dlwiga2	 Alignment		7.6	63	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: LIM domain
4	dlq47a1	 Alignment		7.5	40	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: LIM domain
5	dlzfoa_	 Alignment		6.8	56	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: LASP-1
6	d2dloa2	 Alignment		5.7	23	Fold: Glucocorticoid receptor-like (DNA-binding domain) Superfamily: Glucocorticoid receptor-like (DNA-binding domain) Family: LIM domain
7	dlzupa1	 Alignment		5.7	7	Fold: Ribonuclease H-like motif Superfamily: Ribonuclease H-like Family: TM1739-like
8	dles6a2	 Alignment		5.5	29	Fold: EV matrix protein Superfamily: EV matrix protein Family: EV matrix protein

9 [d2cupa2](#)

Alignment



5.4

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Fold:Glucocorticoid receptor-like (DNA-binding domain)
Superfamily:Glucocorticoid receptor-like (DNA-binding domain)
Family:LIM domain