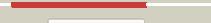
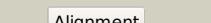
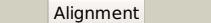
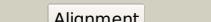


Phyre²

Email	mdejesus@rockefeller.edu
Description	RVBD1153c_(omt)_1280003_1280851
Date	Wed Jul 31 22:05:23 BST 2019
Unique Job ID	9f13648f980acc58

Detailed template information

#	Template	Alignment Coverage	3D Model	Confidence	% i.d.	Template Information
1	d2uyoA1			100.0	23	Fold: S-adenosyl-L-methionine-dependent methyltransferases Superfamily: S-adenosyl-L-methionine-dependent methyltransferases Family: ML2640-like
2	c3ieiD_			100.0	23	PDB header: transferase Chain: D: PDB Molecule: leucine carboxyl methyltransferase 1; PDBTitle: crystal structure of human leucine carboxylmethyltransferase-1 in2 complex with s-adenosyl homocysteine
3	d1rjda_			100.0	22	Fold: S-adenosyl-L-methionine-dependent methyltransferases Superfamily: S-adenosyl-L-methionine-dependent methyltransferases Family: Leucine carboxy methyltransferase Ppm1
4	c3mntA_			100.0	25	PDB header: transferase Chain: A: PDB Molecule: leucine carboxyl methyltransferase 1; PDBTitle: crystal structure of human leucine carboxyl methyltransferase 1
5	c2zwaA_			100.0	17	PDB header: transferase Chain: A: PDB Molecule: leucine carboxyl methyltransferase 2; PDBTitle: crystal structure of trna wybutosine synthesizing enzyme2 tyw4
6	c2qe6B_			99.7	13	PDB header: transferase Chain: B: PDB Molecule: uncharacterized protein tfu_2867; PDBTitle: crystal structure of a putative methyltransferase (tfu_2867) from2 thermobifida fusca yx at 1.95 a resolution
7	c3giwA_			99.3	20	PDB header: unknown function Chain: A: PDB Molecule: protein of unknown function duf574; PDBTitle: crystal structure of a duf574 family protein (sav_2177) from2 streptomyces avermitilis ma-4680 at 1.45 a resolution
8	c4uy5A_			98.5	15	PDB header: transferase Chain: A: PDB Molecule: histidine-specific methyltransferase egtd; PDBTitle: crystal structure of histidine-specific methyltransferase egtd from2 mycobacterium smegmatis
9	c5wp5A_			98.3	14	PDB header: transferase Chain: A: PDB Molecule: phosphomethylethanolamine n-methyltransferase 2; PDBTitle: arabidopsis thaliana phosphoethanolamine n-methyltransferase 22 (atpmt2) in complex with sah
10	c3ofkA_			98.1	16	PDB header: transferase Chain: A: PDB Molecule: nodulation protein s; PDBTitle: crystal structure of n-methyltransferase nods from bradyrhizobium2 japonicum wM9 in complex with s-adenosyl-l-homocysteine (sah)
11	c4krqB_			98.0	11	PDB header: transferase Chain: B: PDB Molecule: phosphoethanolamine n-methyltransferase 1; PDBTitle: semet haemonchus contortus phosphoethanolamine n-methyltransferase 12 in complex with phosphoethanolamine and s-adenosylhomocysteine

12	d1xvaa			97.9	13	Fold: S-adenosyl-L-methionine-dependent methyltransferases Superfamily: S-adenosyl-L-methionine-dependent methyltransferases Family: Glycine N-methyltransferase
13	d1r74a			97.9	17	Fold: S-adenosyl-L-methionine-dependent methyltransferases Superfamily: S-adenosyl-L-methionine-dependent methyltransferases Family: Glycine N-methyltransferase
14	c3jwgA			97.8	14	PDB header: transferase Chain: A: PDB Molecule: methyltransferase type 12; PDBTitle: crystal structure analysis of the methyltransferase domain of2 bacterial-cthen1-c
15	c5thyB			97.7	10	PDB header: transferase,lyase Chain: B: PDB Molecule: curj; PDBTitle: crystal structure of semet-substituted curj carbon methyltransferase
16	c5do0A			97.7	12	PDB header: transferase Chain: A: PDB Molecule: protein lysine methyltransferase 1; PDBTitle: the structure of pkmt1 from rickettsia prowazekii
17	c5dpIB			97.7	15	PDB header: transferase Chain: B: PDB Molecule: protein lysine methyltransferase 2; PDBTitle: the structure of pkmt2 from rickettsia typhi in complex with adohcy
18	c3g2qA			97.7	19	PDB header: transferase Chain: A: PDB Molecule: pcza361.24; PDBTitle: crystal structure of the glycopeptide n-methyltransferase mtfa2 complexed with sinefungin
19	d1wzna1			97.7	15	Fold: S-adenosyl-L-methionine-dependent methyltransferases Superfamily: S-adenosyl-L-methionine-dependent methyltransferases Family: CAC2371-like
20	c3gwzB			97.7	17	PDB header: transferase Chain: B: PDB Molecule: mmcr; PDBTitle: structure of the mitomycin 7-o-methyltransferase mmcr
21	d1y8ca		not modelled	97.6	9	Fold: S-adenosyl-L-methionine-dependent methyltransferases Superfamily: S-adenosyl-L-methionine-dependent methyltransferases Family: CAC2371-like
22	c3ocjA		not modelled	97.6	22	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: putative exported protein; PDBTitle: the crystal structure of a possilbe exported protein from bordetella parapertussis
23	c3jwhA		not modelled	97.6	13	PDB header: transferase Chain: A: PDB Molecule: hen1; PDBTitle: crystal structure analysis of the methyltransferase domain of2 bacterial-avhen1-c
24	d1ri5a		not modelled	97.5	15	Fold: S-adenosyl-L-methionine-dependent methyltransferases Superfamily: S-adenosyl-L-methionine-dependent methyltransferases Family: mRNA cap (Guanine N-7) methyltransferase
25	c1z3cA		not modelled	97.5	15	PDB header: transferase Chain: A: PDB Molecule: mRNA capping enzyme; PDBTitle: encephalitozoano cuniculi mRNA cap (guanine-n7)2 methyltransferasein complexed with azoadomet
26	c6clwA		not modelled	97.5	22	PDB header: biosynthetic protein Chain: A: PDB Molecule: o-methyltransferase; PDBTitle: crystal structure of tnmh
27	c4xcxA		not modelled	97.5	16	PDB header: transferase Chain: A: PDB Molecule: small rna 2'-o-methyltransferase; PDBTitle: methyltransferase domain of small rna 2'-o-methyltransferase
28	c4htfA		not modelled	97.5	9	PDB header: transferase Chain: A: PDB Molecule: s-adenosylmethionine-dependent methyltransferase; PDBTitle: crystal structure of s-adenosylmethionine-dependent methyltransferase2 from escherichia coli in complex with s-

29	d1d2ha		Alignment	not modelled	97.5	16 adenosylmethionine. Fold: S-adenosyl-L-methionine-dependent methyltransferases Superfamily: S-adenosyl-L-methionine-dependent methyltransferases Family: Glycine N-methyltransferase
30	d2bzga1		Alignment	not modelled	97.5	16 Fold: S-adenosyl-L-methionine-dependent methyltransferases Superfamily: S-adenosyl-L-methionine-dependent methyltransferases Family: Thiopurine S-methyltransferase
31	d1im8a		Alignment	not modelled	97.5	14 Fold: S-adenosyl-L-methionine-dependent methyltransferases Superfamily: S-adenosyl-L-methionine-dependent methyltransferases Family: Hypothetical protein HI0319 (YecO)
32	c5mgzA		Alignment	not modelled	97.5	12 PDB header: transferase Chain: A: PDB Molecule: 8-demethylnovobioc acid c(8)-methyltransferase; PDBTitle: streptomyces sphaeroides novo (8-demethylnovobioc acid2 methyltransferase) with sah
33	c6ifsb		Alignment	not modelled	97.5	11 PDB header: transferase Chain: B: PDB Molecule: ribosomal rna small subunit methyltransferase a; PDBTitle: ksga from bacillus subtilis 168
34	c3pfhD		Alignment	not modelled	97.5	16 PDB header: transferase Chain: D: PDB Molecule: n-methyltransferase; PDBTitle: x-ray crystal structure the n,n-dimethyltransferase tylm1 from streptomyces fradiae in complex with sah and ddtp-quip3n
35	c5hjIA		Alignment	not modelled	97.4	13 PDB header: transferase Chain: A: PDB Molecule: glycine sarcosine n-methyltransferase; PDBTitle: crystal structure of glycine sarcosine n-methyltransferase from2 methanohalophilus portcalensis in complex with betaine
36	c3ggdA		Alignment	not modelled	97.4	12 PDB header: transferase Chain: A: PDB Molecule: sam-dependent methyltransferase; PDBTitle: crystal structure of sam-dependent methyltransferase (yp_325210.1)2 from anabaena variabilis atcc 29413 at 2.11 a resolution
37	c4necC		Alignment	not modelled	97.4	12 PDB header: transferase/antibiotic Chain: C: PDB Molecule: putative sam-dependent methyltransferase; PDBTitle: conversion of a disulfide bond into a thioacetal group during2 echinomycin biosynthesis
38	d2ex4a1		Alignment	not modelled	97.4	10 Fold: S-adenosyl-L-methionine-dependent methyltransferases Superfamily: S-adenosyl-L-methionine-dependent methyltransferases Family: AD-003 protein-like
39	c3m70A		Alignment	not modelled	97.4	15 PDB header: structural genomics, unknown function Chain: A: PDB Molecule: stellurite resistance protein tehb homolog; PDBTitle: crystal structure of tehb from haemophilus influenzae
40	c3bgvC		Alignment	not modelled	97.4	12 PDB header: transferase Chain: C: PDB Molecule: mRNA cap guanine-n7 methyltransferase; PDBTitle: crystal structure of mRNA cap guanine-n7 methyltransferase2 in complex with sah
41	d1ve3a1		Alignment	not modelled	97.4	13 Fold: S-adenosyl-L-methionine-dependent methyltransferases Superfamily: S-adenosyl-L-methionine-dependent methyltransferases Family: CAC2371-like
42	c4a6dA		Alignment	not modelled	97.4	12 PDB header: transferase Chain: A: PDB Molecule: hydroxyindole o-methyltransferase; PDBTitle: crystal structure of human n-acetylserotonin2 methyltransferase (asmt) in complex with sam
43	c3lccA		Alignment	not modelled	97.4	13 PDB header: transferase Chain: A: PDB Molecule: putative methyl chloride transferase; PDBTitle: structure of a sam-dependent halide methyltransferase from arabidopsis2 thaliana
44	d2i6ga1		Alignment	not modelled	97.4	18 Fold: S-adenosyl-L-methionine-dependent methyltransferases Superfamily: S-adenosyl-L-methionine-dependent methyltransferases Family: TehB-like
45	c3bgdB		Alignment	not modelled	97.3	15 PDB header: transferase Chain: B: PDB Molecule: thiopurine s-methyltransferase; PDBTitle: thiopurine s-methyltransferase
46	c6mroA		Alignment	not modelled	97.3	13 PDB header: transferase Chain: A: PDB Molecule: methyl transferase from methanosarcina acetivorans; PDBTitle: crystal structure of methyl transferase from methanosarcina2 acetivorans at 1.6 angstroms resolution, northeast structural3 genomics consortium (nsg) target mvr53.
47	c6iv7B		Alignment	not modelled	97.3	17 PDB header: transferase Chain: B: PDB Molecule: methyltransferase lepi; PDBTitle: the crystal structure of a sam-dependent enzyme from aspergillus2 flavus
48	c5bszA		Alignment	not modelled	97.3	13 PDB header: transferase Chain: A: PDB Molecule: n-methyltransferase; PDBTitle: x-ray structure of the sugar n-methyltransferase keds8 from2 streptalloleitichus sp atcc 53650
49	c3cgkB		Alignment	not modelled	97.3	12 PDB header: transferase Chain: B: PDB Molecule: sam-dependent methyltransferase; PDBTitle: crystal structure of tehb-like sam-dependent methyltransferase2 (np_600671.1) from corynebacterium glutamicum atcc 13032 kitasato at 2.00 a resolution
50	c3mczb		Alignment	not modelled	97.2	18 PDB header: transferase Chain: B: PDB Molecule: o-methyltransferase; PDBTitle: the structure of an o-methyltransferase family protein from2 burkholderia thailandensis.
51	c4qvgC		Alignment	not modelled	97.2	19 PDB header: transferase Chain: C: PDB Molecule: sibl; PDBTitle: crystal structure of s-adenosylmethionine-dependent methyltransferase2 sibl in its apo form
52	c3lcvB		Alignment	not modelled	97.2	14 PDB header: transferase Chain: B: PDB Molecule: sisomicin-gentamicin resistance methylase sgm; PDBTitle: crystal structure of antibiotic related methyltransferase

53	c1kyzC	Alignment	not modelled	97.2	14	PDB header: transferase Chain: C PDB Molecule: caffeic acid 3-o-methyltransferase; PDBTitle: crystal structure analysis of caffeic acid/5-hydroxyferulic acid complex 3/5-o-methyltransferase ferulic acid complex
54	d1gzza2	Alignment	not modelled	97.2	17	Fold: S-adenosyl-L-methionine-dependent methyltransferases Superfamily: S-adenosyl-L-methionine-dependent methyltransferases Family: Plant O-methyltransferase, C-terminal domain
55	d1l1ea	Alignment	not modelled	97.2	13	Fold: S-adenosyl-L-methionine-dependent methyltransferases Superfamily: S-adenosyl-L-methionine-dependent methyltransferases Family: Mycolic acid cyclopropane synthase
56	c3h2bB	Alignment	not modelled	97.2	17	PDB header: transferase Chain: B PDB Molecule: sam-dependent methyltransferase; PDBTitle: crystal structure of the sam-dependent methyltransferase cg3271 from2 corynebacterium glutamicum in complex with s-adenosyl-l-homocysteine3 and pyrophosphate. northeast structural genomics consortium target4 cgr113a
57	c1zgaA	Alignment	not modelled	97.2	14	PDB header: plant protein, transferase Chain: A PDB Molecule: isoflavanone 4'-o-methyltransferase'; PDBTitle: crystal structure of isoflavanone 4'-o-methyltransferase complexed2 with (+)-6a-hydroxymaackiain
58	c4x1oA	Alignment	not modelled	97.2	9	PDB header: transferase Chain: A PDB Molecule: 16s rrna (adenine(1408)-n(1))-methyltransferase; PDBTitle: crystal structure of the 16s rrna (adenine(1408)-n(1))-2 methyltransferase from catenulisporales acidiphilia
59	c5cvvB	Alignment	not modelled	97.1	18	PDB header: transferase Chain: B PDB Molecule: (iso)eugenol o-methyltransferase; PDBTitle: coniferyl alcohol bound monolignol 4-o-methyltransferase 9
60	c4krhB	Alignment	not modelled	97.1	11	PDB header: transferase Chain: B PDB Molecule: phosphoethanolamine n-methyltransferase 2; PDBTitle: semet haemonchus contortus phosphoethanolamine n-methyltransferase 22 in complex with s-adenosyl-l-methionine
61	c2p8jA	Alignment	not modelled	97.1	10	PDB header: transferase Chain: A PDB Molecule: s-adenosylmethionine-dependent methyltransferase; PDBTitle: crystal structure of s-adenosylmethionine-dependent methyltransferase2 (np_349143.1) from clostridium acetobutylicum at 2.00 a resolution
62	c3gjyA	Alignment	not modelled	97.1	20	PDB header: transferase Chain: A PDB Molecule: spermidine synthase; PDBTitle: crystal structure of a probable spermidine synthase from2 corynebacterium glutamicum atcc 13032
63	c4hgyC	Alignment	not modelled	97.1	16	PDB header: transferase Chain: C PDB Molecule: ccbj; PDBTitle: structure of the ccbj methyltransferase from streptomyces caelestis
64	c2r3sA	Alignment	not modelled	97.1	10	PDB header: transferase Chain: A PDB Molecule: uncharacterized protein; PDBTitle: crystal structure of a putative o-methyltransferase (npun_r0239) from2 nostoc punctiforme pcc 73102 at 2.15 a resolution
65	d1xtpa	Alignment	not modelled	97.1	16	Fold: S-adenosyl-L-methionine-dependent methyltransferases Superfamily: S-adenosyl-L-methionine-dependent methyltransferases Family: AD-003 protein-like
66	c4iv0B	Alignment	not modelled	97.1	13	PDB header: transferase Chain: B PDB Molecule: phosphoethanolamine n-methyltransferase, putative; PDBTitle: crystal structure of n-methyl transferase from plasmodium vivax2 complexed with s-adenosyl methionine and phosphate
67	c3adnB	Alignment	not modelled	97.1	20	PDB header: transferase Chain: B PDB Molecule: spermidine synthase; PDBTitle: crystal structure of spermidine synthase from e. coli.
68	c6dcba	Alignment	not modelled	97.1	13	PDB header: transferase/rna Chain: A PDB Molecule: 7sk snrna methylphosphate capping enzyme; PDBTitle: structure of methylphosphate capping enzyme methyltransferase domain2 in complex with 5' end of 7sk rna
69	c3dtmA	Alignment	not modelled	97.1	12	PDB header: structural genomics, unknown function Chain: A PDB Molecule: putative methyltransferase mm_2633; PDBTitle: crystal structure of putative methyltransferase-mm_2633 from2 methanoscarcina maezi .
70	c3dp7B	Alignment	not modelled	97.1	11	PDB header: transferase Chain: B PDB Molecule: sam-dependent methyltransferase; PDBTitle: crystal structure of sam-dependent methyltransferase from bacteroides2 vulgatus atcc 8482
71	c4qdkB	Alignment	not modelled	97.1	10	PDB header: transferase Chain: B PDB Molecule: magnesium-protoporphyrin o-methyltransferase; PDBTitle: crystal structure of magnesium protoporphyrin ix methyltransferase2 (chl) from synechocystis pcc 6803 with bound sah
72	c3bxoA	Alignment	not modelled	97.1	15	PDB header: transferase Chain: A PDB Molecule: n,n-dimethyltransferase; PDBTitle: crystal structure of streptomyces venezuelae desvi
73	c2fk8A	Alignment	not modelled	97.1	17	PDB header: transferase Chain: A PDB Molecule: methoxy mycolic acid synthase 4; PDBTitle: crystal structure of hma (mmaa4) from mycobacterium tuberculosis2 complexed with s-adenosylmethionine
74	c1x1aA	Alignment	not modelled	97.1	11	PDB header: transferase Chain: A PDB Molecule: crtf-related protein; PDBTitle: crystal structure of bchu complexed with s-adenosyl-l-methionine
75	c2ip2B	Alignment	not modelled	97.1	19	PDB header: transferase Chain: B PDB Molecule: probable phenazine-specific methyltransferase; PDBTitle: structure of the pyocyanin biosynthetic protein phzm

76	c3ujcA		Alignment	not modelled	97.1	15	PDB header: transferase Chain: A: PDB Molecule: phosphoethanolamine n-methyltransferase; PDBTitle: phosphoethanolamine methyltransferase mutant (h132a) from plasmidum2 falciparum in complex with phosphocholine
77	c5kn4B		Alignment	not modelled	97.0	9	PDB header: transferase Chain: B: PDB Molecule: pavine n-methyltransferase; PDBTitle: pavine n-methyltransferase apoenzyme ph 6.0
78	d1fp2a2		Alignment	not modelled	97.0	13	Fold: S-adenosyl-L-methionine-dependent methyltransferases Superfamily: S-adenosyl-L-methionine-dependent methyltransferases Family: Plant O-methyltransferase, C-terminal domain
79	c4iscA		Alignment	not modelled	97.0	13	PDB header: transferase Chain: A: PDB Molecule: methyltransferase; PDBTitle: crystal structure of a putative methyltransferase from pseudomonas2 syringae
80	d2b2ca1		Alignment	not modelled	97.0	18	Fold: S-adenosyl-L-methionine-dependent methyltransferases Superfamily: S-adenosyl-L-methionine-dependent methyltransferases Family: Spermidine synthase
81	d2fk8a1		Alignment	not modelled	97.0	18	Fold: S-adenosyl-L-methionine-dependent methyltransferases Superfamily: S-adenosyl-L-methionine-dependent methyltransferases Family: Mycolic acid cyclopropane synthase
82	c6gkvB		Alignment	not modelled	97.0	10	PDB header: transferase Chain: B: PDB Molecule: coclaurine n-methyltransferase; PDBTitle: crystal structure of cochlaurine n-methyltransferase (cnmt) bound to n-2 methylheliamine and sah
83	d1iy9a		Alignment	not modelled	97.0	17	Fold: S-adenosyl-L-methionine-dependent methyltransferases Superfamily: S-adenosyl-L-methionine-dependent methyltransferases Family: Spermidine synthase
84	c3p9kD		Alignment	not modelled	97.0	15	PDB header: transferase Chain: D: PDB Molecule: caffic acid o-methyltransferase; PDBTitle: crystal structure of perennial ryegrass lpmot1 complexed with s-2 adenosyl-l-homocysteine and coniferaldehyde
85	c2xvmB		Alignment	not modelled	97.0	16	PDB header: transferase Chain: B: PDB Molecule: tellurite resistance protein tehb; PDBTitle: crystal structure of the tellurite detoxification protein2 tehb from e. coli in complex with sah
86	c3b89A		Alignment	not modelled	97.0	15	PDB header: transferase Chain: A: PDB Molecule: 16s rrna methylase; PDBTitle: crystal structure of rrna methylase from escherichia coli
87	c3d2IC		Alignment	not modelled	97.0	12	PDB header: transferase Chain: C: PDB Molecule: sam-dependent methyltransferase; PDBTitle: crystal structure of sam-dependent methyltransferase (zp_00538691.1)2 from exiguobacterium sp. 255-15 at 1.90 a resolution
88	c1fpqA		Alignment	not modelled	97.0	15	PDB header: transferase Chain: A: PDB Molecule: isoliquiritigenin 2'-o-methyltransferase; PDBTitle: crystal structure analysis of selenomethionine substituted chalcone o-2 methyltransferase
89	c6i5zA		Alignment	not modelled	97.0	21	PDB header: transferase Chain: A: PDB Molecule: o-methyltransferase 1; PDBTitle: papaver somniferum o-methyltransferase
90	d1tpya		Alignment	not modelled	97.0	13	Fold: S-adenosyl-L-methionine-dependent methyltransferases Superfamily: S-adenosyl-L-methionine-dependent methyltransferases Family: Mycolic acid cyclopropane synthase
91	c2gs9A		Alignment	not modelled	97.0	11	PDB header: transferase Chain: A: PDB Molecule: hypothetical protein tt1324; PDBTitle: crystal structure of tt1324 from thermus thermophilis hb8
92	c5i2hB		Alignment	not modelled	97.0	18	PDB header: transferase Chain: B: PDB Molecule: o-methyltransferase family 2; PDBTitle: crystal structure of o-methyltransferase family 2 protein plim_11472 from planctomyces limnophilus dsm 3776 complex with apigenin
93	c1tw3A		Alignment	not modelled	96.9	20	PDB header: transferase Chain: A: PDB Molecule: carminomycin 4-o-methyltransferase; PDBTitle: crystal structure of carminomycin-4-o-methyltransferase (dnrk) in2 complex with s-adenosyl-l-homocystein (sah) and 4-methoxy-e-3 rhodomycin t (m-et)
94	c1xdwA		Alignment	not modelled	96.9	18	PDB header: transferase Chain: A: PDB Molecule: protein rdmb; PDBTitle: crystal structure of aclacinomycin-10-hydroxylase (rdmb) in complex2 with sinefungin (sfg)
95	c3g5tA		Alignment	not modelled	96.9	19	PDB header: transferase Chain: A: PDB Molecule: trans-aconitate 3-methyltransferase; PDBTitle: crystal structure of trans-aconitate 3-methyltransferase from yeast
96	c5xohA		Alignment	not modelled	96.9	15	PDB header: transferase Chain: A: PDB Molecule: bergaptol o-methyltransferase; PDBTitle: crystal structure of bergaptol o-methyltransferase complex
97	c6c5bB		Alignment	not modelled	96.9	15	PDB header: transferase Chain: B: PDB Molecule: methyltransferase; PDBTitle: crystal structure analysis of laphzm
98	c4p7cB		Alignment	not modelled	96.9	10	PDB header: transferase Chain: B: PDB Molecule: trna (mo5u34)-methyltransferase; PDBTitle: crystal structure of putative methyltransferase from pseudomonas2 syringae pv. tomato
99	d1qyra		Alignment	not modelled	96.9	10	Fold: S-adenosyl-L-methionine-dependent methyltransferases Superfamily: S-adenosyl-L-methionine-dependent methyltransferases Family: rRNA adenine dimethylase-like
100	c5z9oA		Alignment	not modelled	96.9	12	PDB header: transferase Chain: A: PDB Molecule: cyclopropane-fatty-acyl-phospholipid synthase; PDBTitle: the crystal structure of cyclopropane-fatty-acyl-

						phospholipid synthase2 from lactobacillus acidophilus
101	c4kifB	Alignment	not modelled	96.9	13	PDB header: transferase Chain: B: PDB Molecule: methyltransferase mppj; PDBTitle: crystal structure of methyltransferase from streptomyces hygroscopicus2 complexed with phenylpyruvic acid
102	c2gpyB	Alignment	not modelled	96.9	15	PDB header: structural genomics, unknown function Chain: B: PDB Molecule: o-methyltransferase; PDBTitle: crystal structure of putative o-methyltransferase from bacillus2 halodurans
103	c5cm2Z	Alignment	not modelled	96.8	13	PDB header: transferase Chain: Z: PDB Molecule: tRNA methyltransferase; PDBTitle: insights into molecular plasticity in protein complexes from trm9-2 trm112 tRNA modifying enzyme crystal structure
104	c2hteB	Alignment	not modelled	96.8	13	PDB header: transferase Chain: B: PDB Molecule: spermidine synthase; PDBTitle: the crystal structure of spermidine synthase from p. falciparum in2 complex with 5'-methylthioadenosine
105	c4kvzA	Alignment	not modelled	96.8	7	PDB header: transferase Chain: A: PDB Molecule: baml; PDBTitle: crystal structure of the plantazolicin methyltransferase baml in2 complex with sah
106	d1kpga	Alignment	not modelled	96.8	13	Fold: S-adenosyl-L-methionine-dependent methyltransferases Superfamily: S-adenosyl-L-methionine-dependent methyltransferases Family: Mycolic acid cyclopropane synthase
107	c6ectA	Alignment	not modelled	96.8	10	PDB header: transferase Chain: A: PDB Molecule: stie protein; PDBTitle: stie o-mt residues 961-1257
108	c6bqcA	Alignment	not modelled	96.8	14	PDB header: transferase Chain: A: PDB Molecule: cyclopropane-fatty-acyl-phospholipid synthase; PDBTitle: cyclopropane fatty acid synthase from e. coli
109	c3merA	Alignment	not modelled	96.8	12	PDB header: structural genomics, unknown function Chain: A: PDB Molecule: slr1183 protein; PDBTitle: crystal structure of the methyltransferase slr1183 from synechocystis2 sp. pcc 6803, northeast structural genomics consortium target sgr145
110	d1fp1d2	Alignment	not modelled	96.8	11	Fold: S-adenosyl-L-methionine-dependent methyltransferases Superfamily: S-adenosyl-L-methionine-dependent methyltransferases Family: Plant O-methyltransferase, C-terminal domain
111	c4ineB	Alignment	not modelled	96.8	15	PDB header: transferase Chain: B: PDB Molecule: protein pmt-2; PDBTitle: crystal structure of n-methyl transferase (pmt-2) from caenorhabditis2 elegans complexed with s-adenosyl homocysteine and3 phosphoethanolamine
112	c3qnhA	Alignment	not modelled	96.7	15	PDB header: transferase Chain: A: PDB Molecule: arsenic methyltransferase; PDBTitle: arm arsenic(iii) s-adenosylmethionine methyltransferase with as(iii)
113	c4jxjA	Alignment	not modelled	96.7	12	PDB header: transferase Chain: A: PDB Molecule: ribosomal RNA small subunit methyltransferase a; PDBTitle: crystal structure of ribosomal RNA small subunit methyltransferase a2 from rickettsia bellii determined by iodide sad phasing
114	c4gekG	Alignment	not modelled	96.7	16	PDB header: transferase Chain: G: PDB Molecule: tRNA (cmo5u34)-methyltransferase; PDBTitle: crystal structure of wild-type cmoa from e.coli
115	c2cmgA	Alignment	not modelled	96.7	9	PDB header: transferase Chain: A: PDB Molecule: spermidine synthase; PDBTitle: crystal structure of spermidine synthase from helicobacter2 pylori
116	c3tqsB	Alignment	not modelled	96.7	10	PDB header: transferase Chain: B: PDB Molecule: ribosomal RNA small subunit methyltransferase a; PDBTitle: structure of the dimethyladenosine transferase (ksga) from coxiella2 burnetii
117	c5ubba	Alignment	not modelled	96.7	11	PDB header: transferase Chain: A: PDB Molecule: alpha n-terminal protein methyltransferase 1b; PDBTitle: crystal structure of human alpha n-terminal protein methyltransferase2 1b
118	c4qttB	Alignment	not modelled	96.7	9	PDB header: transferase Chain: B: PDB Molecule: putative methyltransferase bud23; PDBTitle: structure of s. cerevisiae bud23-trm112 complex involved in formation2 of m7g1575 on 18s rRNA (apo-form)
119	c4d7kB	Alignment	not modelled	96.6	20	PDB header: transferase Chain: B: PDB Molecule: SAM-dependent methyltransferases; PDBTitle: crystal structure of n,n-8-amino-8-demethyl-d-riboflavin2 dimethyltransferase (rosa) from streptomyces davawensis
120	d2o07a1	Alignment	not modelled	96.6	17	Fold: S-adenosyl-L-methionine-dependent methyltransferases Superfamily: S-adenosyl-L-methionine-dependent methyltransferases Family: Spermidine synthase