

The Oxidation Potential of AA steps in single strand DNA Oligomers

– Electronic Supplementary Information –

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TABLE S1. Sum of the Mulliken spin densities^(a) (PCM/M06-2X/6-31G**) for the singly ionized trimers (H model).

	5'-CCA-3'	5'-ACC-3'	5'-CAA-3'	5'-AAC-3'
5'-B ₁	0.00	0.94	0.05	0.08
B ₂	0.02	0.06	0.87	0.85
B ₃ -3'	0.98	0.00	0.08	0.07

^(a)The sum has been carried out only for the atoms belonging to each nucleobase B_i, (i = 1,2,3) without including ribose and phosphate atoms.

TABLE S2. Sum of the Mulliken spin densities^(a) (PCM/M06-2X/6-31G**) for the singly ionized trimers (PO- model).

	5'-CCA-3'	5'-ACC-3'	5'-CAA-3'	5'-AAC-3'
5'-B ₁	0.00	0.94	0.05	0.07
B ₂	0.03	0.05	0.93	0.91
B ₃ -3'	0.97	0.00	0.01	0.02

^(a)The sum has been carried out only for the atoms belonging to each nucleobase B_i.

TABLE S3. Sum of the Mulliken spin densities^(a) (PCM/B3LYP-D/6-31G**) for the singly ionized trimers (H model).

	5'-CCA-3'	5'-ACC-3'	5'-CAA-3'	5'-AAC-3'
5'-B ₁	0.00	0.81	0.02	0.37
B ₂	0.36	0.18	0.55	0.56
B ₃ -3'	0.63	0.00	0.43	0.07

^(a)The sum has been carried out only for the atoms belonging to each nucleobase B_i.

TABLE S4. Sum of the Mulliken spin densities^(a) (PCM/B3LYP-D/6-31G**) for the singly ionized trimers (PO- model).

	5'-CCA-3'	5'-ACC-3'	5'-CAA-3'	5'-AAC-3'
5'-B ₁	0.03	0.79	0.16	0.36
B ₂	0.39	0.19	0.76	0.56
B ₃ -3'	0.57	0.02	0.07	0.08

^(a)The sum has been carried out only for the atoms belonging to each nucleobase B_i.

TABLE S5. Mulliken charges of neutral and singly ionized 5'-ACC-3' single strand evaluated by using the M06-2X functional. See scheme SI for the definition of the fragments.

	A1	dR1	P12	(H/Na)12	dR2	C2	P23	(H/Na)13	dR3	C3
H model										
neut.	-0.27	0.01	-0.48	0.40	0.33	-0.24	-0.47	0.40	0.60	-0.28
sing. ion.	0.53	0.09	-0.47	0.40	0.34	-0.16	-0.47	0.40	0.61	-0.27
Na model										
neut.	-0.28	-0.05	-0.67	0.72	0.21	-0.26	-0.66	0.72	0.54	-0.27
sing. ion.	0.51	0.02	-0.67	0.73	0.23	-0.16	-0.65	0.72	0.54	-0.27
PO- model										
neut.	-0.28	-0.10	-0.84	—	0.15	-0.29	-0.84	—	0.51	-0.30
sing. ion.	0.53	0.00	-0.84	—	0.17	-0.21	-0.84	—	0.49	-0.29

TABLE S6. Mulliken charges of neutral and singly ionized 5'-AAC-3' single strand evaluated by using the M06-2X functional. See scheme SI for the definition of the fragments.

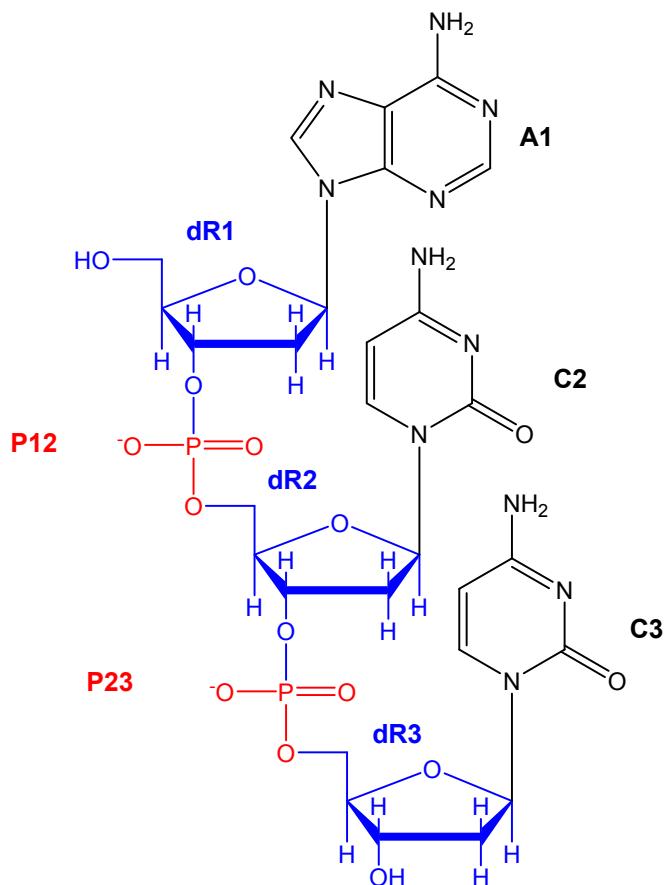
	A1	dR1	P12	(H/Na)12	dR2	A2	P23	(H/Na)13	dR3	C3
H model										
neut.	-0.28	0.02	-0.48	0.40	0.35	-0.27	-0.48	0.40	0.61	-0.28
sing. ion.	-0.20	0.04	-0.47	0.40	0.40	0.44	-0.47	0.40	0.63	-0.18
Na model										
neut.	-0.28	-0.03	-0.67	0.72	0.23	-0.30	-0.65	0.72	0.55	-0.27
sing. ion.	-0.20	-0.03	-0.67	0.73	0.29	0.44	-0.65	0.72	0.58	-0.20
PO- model										
neut.	-0.28	-0.08	-0.85	—	0.17	-0.30	-0.83	—	0.49	-0.31
sing. ion.	-0.22	-0.07	-0.83	—	0.24	0.44	-0.81	—	0.52	-0.26

TABLE S7. Mulliken charges of neutral and singly ionized 5'-ACC-3' single strand evaluated by using the B3LYP-D functional. See scheme SI for the definition of the fragments.

	A1	dR1	P12	(H/Na)12	dR2	C2	P23	(H/Na)13	dR3	C3
H model										
neut.	-0.26	0.02	-0.45	0.37	0.32	-0.24	-0.44	0.37	0.57	-0.27
sing. ion.	0.42	0.09	-0.44	0.37	0.34	-0.05	-0.44	0.37	0.58	-0.25
Na model										
neut.	-0.27	-0.04	-0.63	0.70	0.20	-0.27	-0.64	0.71	0.50	-0.28
sing. ion.	0.41	0.02	-0.62	0.71	0.22	-0.07	-0.64	0.71	0.51	-0.26
PO- model										
neut.	-0.27	-0.09	-0.82	—	0.12	-0.29	-0.83	—	0.47	-0.29
sing. ion.	0.40	-0.02	-0.81	—	0.16	-0.09	-0.82	—	0.46	-0.27

TABLE S8. Mulliken charges of neutral and singly ionized 5'-AAC-3' single strand evaluated by using the B3LYP-D functional. See scheme SI for the definition of the fragments.

	A1	dR1	P12	(H/Na)12	dR2	A2	P23	(H/Na)13	dR3	C3
H model										
neut.	-0.27	0.03	-0.45	0.37	0.35	-0.26	-0.45	0.37	0.58	-0.28
sing. ion.	0.06	0.07	-0.45	0.37	0.37	0.20	-0.44	0.37	0.61	-0.17
Na model										
neut.	-0.26	-0.03	-0.63	0.70	0.22	-0.30	-0.62	0.70	0.52	-0.30
sing. ion.	-0.01	0.00	-0.63	0.70	0.26	0.21	-0.62	0.70	0.55	-0.19
PO- model										
neut.	-0.26	-0.08	-0.83	—	0.16	-0.30	-0.81	—	0.44	-0.31
sing. ion.	0.04	-0.05	-0.81	—	0.21	0.17	-0.81	—	0.48	-0.22



SCHEME SI

Optimized Geometries: local base, helical step and conformational parameters

5'-ACC-3' (H model) neutral / B3LYP-D

step overlap area (\AA^2)
 1 A/C 3.88
 2 C/C 4.37

 Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 A/C	0.49	0.26	2.44	13.93	-0.28	34.11
2 C/C	0.32	0.46	2.83	6.43	-2.31	36.88
ave.	0.41	0.36	2.63	10.18	-1.29	35.50
s.d.	0.12	0.14	0.27	5.30	1.44	1.96

 Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 A/C	0.45	0.70	2.45	-0.45	-22.62	36.77
2 C/C	0.98	0.23	2.81	-3.61	-10.06	37.49
ave.	0.71	0.46	2.63	-2.03	-16.34	37.13
s.d.	0.38	0.33	0.25	2.23	8.88	0.51

 Sugar conformational parameters:
 Note: v0: C4'-O4'-C1'-C2'; v1: O4'-C1'-C2'-C3'; v2: C1'-C2'-C3'-C4'
 v3: C2'-C3'-C4'-O4'; v4: C3'-C4'-O4'-C1'
 tm: the amplitude of pucker; P: the phase angle of pseudorotation

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 A	-10.6	30.1	-36.9	31.8	-13.3	37.0	182.2	C3'-exo
2 C	-44.3	46.6	-31.5	6.5	23.4	47.1	132.0	C1'-exo
3 C	-45.0	44.6	-27.4	1.9	26.7	46.2	126.4	C1'-exo

5'-ACC-3' (H model) singly ionized / B3LYP-D

step overlap area (\AA^2)
 1 A/C 3.53
 2 C/C 5.19

 Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 A/C	1.69	1.30	1.63	30.05	-4.22	43.98
2 C/C	0.65	0.72	3.31	-1.38	-1.69	37.79
ave.	1.17	1.01	2.47	14.34	-2.95	40.89
s.d.	0.74	0.40	1.19	22.23	1.79	4.38

 Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 A/C	1.63	-0.65	2.18	-4.99	-35.57	53.01
2 C/C	1.33	-1.18	3.25	-2.60	2.13	37.85
ave.	1.48	-0.91	2.72	-3.79	-16.72	45.43
s.d.	0.21	0.38	0.76	1.69	26.66	10.72

 Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 A	-9.9	27.1	-32.7	28.0	-11.5	32.7	181.4	C3'-exo
2 C	-44.1	46.7	-32.0	7.3	22.7	47.1	132.8	C1'-exo
3 C	-44.9	45.2	-28.3	2.8	26.0	46.4	127.5	C1'-exo

5'-AAC-3' (H model) neutral / B3LYP-D

step overlap area (\AA^2)

1	A/A	5.51
2	A/C	4.14

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 A/A	2.86	1.96	2.92	17.20	-11.77	50.54
2 A/C	0.62	0.01	2.79	5.64	3.87	35.79
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ave.	1.74	0.98	2.86	11.42	-3.95	43.17
s.d.	1.58	1.38	0.09	8.17	11.06	10.43

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 A/A	2.84	-2.11	3.18	-13.17	-19.24	54.41
2 A/C	-0.45	-0.32	2.84	6.22	-9.08	36.42
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ave.	1.19	-1.21	3.01	-3.47	-14.16	45.42
s.d.	2.33	1.27	0.24	13.71	7.18	12.72

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 A	-10.1	27.1	-32.6	27.6	-11.1	32.6	180.9	C3'-exo
2 A	-40.9	45.8	-33.5	10.7	18.7	45.5	137.5	C1'-exo
3 C	-45.1	45.3	-28.2	2.6	26.2	46.6	127.3	C1'-exo

5'-AAC-3' (H model) singly ionized / B3LYP-D

step overlap area (\AA^2)

1	A/A	5.27
2	A/C	5.05

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 A/A	1.07	1.86	2.85	6.17	-2.67	40.93
2 A/C	1.72	0.88	2.67	8.30	-2.31	44.97
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ave.	1.39	1.37	2.76	7.24	-2.49	42.95
s.d.	0.46	0.69	0.13	1.51	0.25	2.86

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 A/A	2.87	-0.94	2.85	-3.78	-8.76	41.45
2 A/C	1.30	-1.59	2.88	-2.99	-10.74	45.74
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ave.	2.09	-1.26	2.86	-3.39	-9.75	43.60
s.d.	1.11	0.46	0.02	0.56	1.40	3.03

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 A	-3.2	25.0	-35.9	35.1	-20.1	37.0	193.7	C3'-exo
2 A	-39.4	45.8	-34.7	12.9	16.3	45.1	140.3	C1'-exo
3 C	-45.1	44.7	-27.4	1.8	26.8	46.3	126.3	C1'-exo

5'-CCA-3' (H model) neutral / B3LYP-Dstep overlap area (\AA^2)

1	C/C	4.86
2	C/A	4.60

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 C/C	0.21	0.80	2.15	20.91	-6.05	32.84
2 C/A	0.70	1.09	3.02	3.98	2.25	38.79
ave.	0.45	0.95	2.59	12.44	-1.90	35.81
s.d.	0.34	0.21	0.62	11.97	5.87	4.21

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 C/C	1.67	1.34	1.80	-9.52	-32.87	39.23
2 C/A	1.38	-0.59	3.13	3.38	-5.97	39.05
ave.	1.52	0.38	2.47	-3.07	-19.42	39.14
s.d.	0.21	1.36	0.94	9.12	19.03	0.13

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 C	-16.0	31.9	-35.0	26.6	-6.7	35.3	172.3	C2'-endo
2 C	-43.8	46.3	-31.7	7.1	22.7	46.8	132.8	C1'-exo
3 A	-42.4	44.2	-29.3	5.4	22.8	44.7	130.9	C1'-exo

5'-CCA-3' (H model) singly ionized / B3LYP-Dstep overlap area (\AA^2)

1	C/C	4.55
2	C/A	5.98

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 C/C	0.37	0.97	1.97	23.98	-6.34	36.58
2 C/A	1.06	2.01	2.75	7.25	0.07	41.60
ave.	0.72	1.49	2.36	15.61	-3.14	39.09
s.d.	0.49	0.74	0.55	11.83	4.53	3.55

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 C/C	1.67	0.92	1.71	-8.95	-33.85	43.96
2 C/A	2.79	-0.83	2.89	0.10	-10.12	42.20
ave.	2.23	0.04	2.30	-4.43	-21.98	43.08
s.d.	0.80	1.24	0.83	6.40	16.78	1.24

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 C	-15.8	31.7	-34.8	26.5	-6.8	35.1	172.4	C2'-endo
2 C	-43.5	45.7	-30.8	6.5	22.8	46.1	131.9	C1'-exo
3 A	-41.5	44.3	-30.1	7.1	21.1	44.2	132.8	C1'-exo

5'-CAA-3' (H model) neutral / B3LYP-D

step overlap area (\AA^2)
 1 C/A 4.13
 2 A/A 5.34

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 C/A	3.16	3.13	2.63	19.95	-7.07	56.81
2 A/A	0.43	0.88	3.17	-0.35	8.66	32.85
ave.	1.80	2.00	2.90	9.80	0.80	44.83
s.d.	1.93	1.60	0.38	14.36	11.13	16.94

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 C/A	3.43	-2.26	3.11	-7.17	-20.21	60.31
2 A/A	0.11	-0.80	3.28	15.00	0.61	33.95
ave.	1.77	-1.53	3.20	3.91	-9.80	47.13
s.d.	2.35	1.03	0.12	15.67	14.72	18.64

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 C	-13.8	29.1	-32.7	25.4	-7.3	32.8	174.3	C2'-endo
2 A	-37.8	43.9	-33.6	12.5	15.6	43.4	140.6	C1'-exo
3 A	-42.4	45.2	-30.6	6.8	22.0	45.3	132.5	C1'-exo

5'-CAA-3' (H model) singly ionized / B3LYP-D

step overlap area (\AA^2)
 1 C/A 4.20
 2 A/A 5.81

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 C/A	3.38	3.14	2.53	23.06	-7.64	62.39
2 A/A	1.23	1.60	2.91	2.93	1.43	40.32
ave.	2.30	2.37	2.72	13.00	-3.10	51.36
s.d.	1.52	1.09	0.27	14.24	6.42	15.60

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 C/A	3.14	-2.25	3.12	-7.10	-21.42	66.50
2 A/A	2.17	-1.48	3.04	2.07	-4.24	40.45
ave.	2.65	-1.87	3.08	-2.51	-12.83	53.48
s.d.	0.69	0.54	0.05	6.48	12.15	18.42

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 C	-14.5	30.3	-33.7	26.1	-7.4	33.9	173.8	C2'-endo
2 A	-37.9	44.2	-33.9	13.0	15.3	43.6	141.0	C1'-exo
3 A	-40.8	45.0	-31.7	9.1	19.5	44.5	135.4	C1'-exo

5'-ACC-3' (Na model) neutral / B3LYP-Dstep overlap area (\AA^2)

1	A/C	4.01
2	C/C	2.81

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 A/C	0.54	-0.06	2.77	6.57	0.85	35.83
2 C/C	0.77	-0.42	3.71	-8.75	9.60	45.82
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ave.	0.66	-0.24	3.24	-1.09	5.22	40.82
s.d.	0.16	0.26	0.66	10.84	6.19	7.06

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## Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 A/C	-0.20	-0.08	2.82	1.36	-10.57	36.41
2 C/C	-1.38	-1.74	3.37	12.04	10.99	47.53
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ave.	-0.79	-0.91	3.10	6.70	0.21	41.97
s.d.	0.83	1.17	0.38	7.55	15.24	7.86

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 A	-12.8	32.8	-39.0	32.8	-12.7	39.0	180.0	C2'-endo
2 C	-42.5	43.6	-28.2	4.1	23.9	44.5	129.4	C1'-exo
3 C	-18.9	33.7	-34.8	24.8	-3.8	35.7	167.4	C2'-endo

5'-ACC-3' (Na model) singly ionized / B3LYP-Dstep overlap area (\AA^2)

1	A/C	1.68
2	C/C	2.64

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 A/C	-1.15	-0.22	2.68	9.69	-19.23	25.28
2 C/C	1.43	-0.78	4.02	-13.95	9.67	50.88
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ave.	0.14	-0.50	3.35	-2.13	-4.78	38.08
s.d.	1.82	0.39	0.94	16.71	20.44	18.11

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## Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 A/C	2.19	3.32	1.86	-36.60	-18.44	33.09
2 C/C	-1.57	-2.61	3.36	10.92	15.74	53.46
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ave.	0.31	0.36	2.61	-12.84	-1.35	43.27
s.d.	2.66	4.19	1.06	33.60	24.17	14.40

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 A	-10.6	30.2	-37.0	32.1	-13.6	37.1	182.4	C3'-exo
2 C	-41.6	45.9	-32.4	9.2	19.9	45.5	135.4	C1'-exo
3 C	-17.0	32.4	-34.7	25.8	-5.7	35.1	170.5	C2'-endo

5'-AAC-3' (Na model) neutral / B3LYP-Dstep overlap area (\AA^2)

1	A/A	5.01
2	A/C	4.23

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 A/A	0.12	1.14	3.01	2.04	11.77	27.42
2 A/C	0.68	-0.17	2.83	5.06	5.23	37.43
ave.	0.40	0.48	2.92	3.55	8.50	32.43
s.d.	0.40	0.93	0.13	2.13	4.63	7.08

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 A/A	-0.23	0.19	3.22	23.47	-4.07	29.87
2 A/C	-0.85	-0.48	2.84	8.05	-7.79	38.11
ave.	-0.54	-0.15	3.03	15.76	-5.93	33.99
s.d.	0.44	0.47	0.27	10.90	2.64	5.83

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 A	-25.4	38.9	-36.9	23.3	1.1	39.4	159.7	C2'-endo
2 A	-34.8	44.8	-37.4	18.4	10.0	44.0	148.2	C2'-endo
3 C	-43.4	44.6	-28.7	4.2	24.2	45.3	129.3	C1'-exo

5'-AAC-3' (Na model) singly ionized / B3LYP-Dstep overlap area (\AA^2)

1	A/A	5.30
2	A/C	5.17

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 A/A	1.09	1.81	2.88	5.36	-2.31	41.34
2 A/C	1.60	0.72	2.73	7.06	-1.79	44.24
ave.	1.34	1.26	2.81	6.21	-2.05	42.79
s.d.	0.36	0.77	0.10	1.20	0.37	2.05

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 A/A	2.75	-1.02	2.89	-3.25	-7.55	41.74
2 A/C	1.08	-1.54	2.91	-2.35	-9.30	44.80
ave.	1.92	-1.28	2.90	-2.80	-8.43	43.27
s.d.	1.18	0.37	0.02	0.63	1.24	2.17

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 A	-5.9	27.6	-37.5	35.3	-18.5	38.1	190.0	C3'-exo
2 A	-35.4	45.8	-38.3	19.1	9.8	44.9	148.6	C2'-endo
3 C	-42.8	44.7	-29.5	5.5	23.1	45.2	130.9	C1'-exo

5'-CCA-3' (Na model) neutral / B3LYP-D

step overlap area (\AA^2)
 1 C/C 2.92
 2 C/A 2.33

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 C/C	-0.44	-0.05	2.57	11.10	-12.14	30.81
2 C/A	0.79	0.76	3.03	2.40	0.94	49.79
ave.	0.18	0.35	2.80	6.75	-5.60	40.30
s.d.	0.87	0.57	0.33	6.15	9.25	13.42

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 C/C	1.34	2.03	2.16	-21.07	-19.27	34.84
2 C/A	0.83	-0.77	3.07	1.12	-2.85	49.85
ave.	1.09	0.63	2.62	-9.97	-11.06	42.35
s.d.	0.36	1.98	0.65	15.69	11.61	10.62

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 C	-17.1	34.5	-37.8	29.0	-7.5	38.2	172.6	C2'-endo
2 C	-40.0	45.8	-34.0	11.6	17.5	45.2	138.8	C1'-exo
3 A	-12.2	29.3	-34.2	28.1	-10.2	34.2	178.3	C2'-endo

5'-CCA-3' (Na model) singly ionized / B3LYP-D

step overlap area (\AA^2)
 1 C/C 2.74
 2 C/A 4.56

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 C/C	-0.37	-0.11	2.69	8.74	-17.63	32.32
2 C/A	1.72	1.15	2.81	8.49	7.40	57.84
ave.	0.68	0.52	2.75	8.62	-5.12	45.08
s.d.	1.48	0.89	0.08	0.17	17.70	18.05

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 C/C	1.71	1.50	2.28	-28.61	-14.18	37.70
2 C/A	0.82	-1.35	3.12	7.57	-8.69	58.84
ave.	1.26	0.08	2.70	-10.52	-11.43	48.27
s.d.	0.63	2.02	0.59	25.58	3.88	14.95

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 C	-17.9	35.4	-38.5	29.2	-7.2	38.9	171.9	C2'-endo
2 C	-41.1	47.0	-34.2	11.4	18.3	46.1	138.0	C1'-exo
3 A	10.1	10.9	-26.0	32.5	-27.1	32.3	216.3	C4'-endo

5'-CAA-3' (Na model) neutral / B3LYP-D

step overlap area (\AA^2)
 1 C/A 2.17
 2 A/A 5.06

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 C/A	-0.02	0.71	2.68	11.37	-9.89	35.44
2 A/A	1.08	0.06	3.23	-3.52	8.22	49.20
ave.	0.53	0.38	2.96	3.93	-0.83	42.32
s.d.	0.78	0.46	0.39	10.53	12.81	9.73

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 C/A	2.06	1.15	2.30	-15.42	-17.73	38.42
2 A/A	-0.52	-1.53	3.12	9.78	4.18	49.96
ave.	0.77	-0.19	2.71	-2.82	-6.77	44.19
s.d.	1.82	1.90	0.58	17.82	15.49	8.16

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 C	-17.3	34.5	-37.8	28.8	-7.3	38.1	172.3	C2'-endo
2 A	-34.0	43.4	-35.8	16.9	10.6	42.6	147.2	C2'-endo
3 A	-12.2	29.6	-34.5	28.4	-10.4	34.5	178.4	C2'-endo

5'-CAA-3' (Na model) singly ionized / B3LYP-D

step overlap area (\AA^2)
 1 C/A 0.03
 2 A/A 5.71

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 C/A	-1.79	0.30	2.24	16.43	-27.56	21.40
2 A/A	1.66	-0.12	3.03	-0.17	8.60	55.09
ave.	-0.07	0.09	2.64	8.13	-9.48	38.25
s.d.	2.44	0.29	0.56	11.74	25.56	23.82

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 C/A	2.20	3.77	0.29	-49.08	-29.26	38.41
2 A/A	-0.58	-1.78	2.98	9.23	0.18	55.71
ave.	0.81	0.99	1.64	-19.93	-14.54	47.06
s.d.	1.97	3.92	1.90	41.23	20.82	12.23

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 C	-15.6	32.2	-35.7	27.7	-7.7	35.9	173.6	C2'-endo
2 A	-36.2	46.1	-37.3	17.6	11.3	44.7	146.5	C2'-endo
3 A	1.8	18.1	-29.6	31.4	-21.1	31.8	201.7	C3'-exo

5'-ACC-3' (PO- model) neutral / B3LYP-Dstep overlap area (\AA^2)

1	A/C	4.04
2	C/C	2.09

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 A/C	0.53	-0.07	2.80	5.69	2.10	35.44
2 C/C	1.11	-0.82	4.22	-16.78	13.85	49.31
ave.	0.82	-0.44	3.51	-5.54	7.97	42.38
s.d.	0.41	0.53	1.00	15.89	8.31	9.81

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 A/C	-0.37	-0.16	2.84	3.41	-9.27	35.94
2 C/C	-1.97	-2.52	3.38	15.74	19.08	53.62
ave.	-1.17	-1.34	3.11	9.58	4.90	44.78
s.d.	1.13	1.67	0.38	8.72	20.05	12.50

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 A	-17.2	35.7	-39.4	30.7	-8.6	39.6	173.6	C2'-endo
2 C	-40.8	44.7	-31.2	8.3	20.2	44.5	134.6	C1'-exo
3 C	-14.2	30.8	-34.6	27.3	-8.4	34.8	175.0	C2'-endo

5'-ACC-3' (PO- model) singly ionized / B3LYP-Dstep overlap area (\AA^2)

1	A/C	1.72
2	C/C	0.78

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 A/C	-1.14	-0.25	2.71	8.85	-18.76	25.44
2 C/C	1.18	-1.28	4.78	-26.30	16.67	49.27
ave.	0.02	-0.77	3.75	-8.73	-1.05	37.35
s.d.	1.64	0.73	1.46	24.86	25.05	16.85

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 A/C	2.18	3.27	1.96	-35.94	-16.96	32.72
2 C/C	-2.54	-3.13	3.28	18.07	28.52	57.76
ave.	-0.18	0.07	2.62	-8.93	5.78	45.24
s.d.	3.34	4.53	0.93	38.19	32.16	17.71

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 A	-11.1	30.7	-37.2	32.0	-13.3	37.3	181.7	C3'-exo
2 C	-39.3	47.0	-35.9	14.2	15.4	45.8	141.6	C1'-exo
3 C	-10.3	27.7	-33.5	28.5	-11.7	33.5	181.2	C3'-exo

5'-AAC-3' (PO- model) neutral / B3LYP-Dstep overlap area (\AA^2)

1	A/A	5.73
2	A/C	4.91

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 A/A	0.63	0.59	3.22	-1.29	8.42	35.27
2 A/C	1.16	-1.33	4.06	-16.27	23.24	49.71
ave.	0.90	-0.37	3.64	-8.78	15.83	42.49
s.d.	0.38	1.35	0.60	10.60	10.48	10.21

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 A/A	-0.24	-1.19	3.24	13.65	2.09	36.26
2 A/C	-2.86	-2.25	2.77	25.41	17.79	56.79
ave.	-1.55	-1.72	3.01	19.53	9.94	46.52
s.d.	1.85	0.74	0.33	8.31	11.10	14.52

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 A	-29.6	42.7	-38.8	23.0	3.8	42.5	156.0	C2'-endo
2 A	-33.5	45.0	-38.3	19.9	8.2	44.0	150.4	C2'-endo
3 C	-8.4	26.1	-32.9	29.0	-13.2	33.0	184.3	C3'-exo

5'-AAC-3' (PO- model) singly ionized / B3LYP-Dstep overlap area (\AA^2)

1	A/A	7.39
2	A/C	5.14

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 A/A	1.39	0.31	3.05	-0.53	5.07	37.53
2 A/C	1.72	-1.55	3.60	-11.45	12.82	54.63
ave.	1.56	-0.62	3.33	-5.99	8.94	46.08
s.d.	0.23	1.31	0.39	7.72	5.48	12.09

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 A/A	-0.14	-2.21	3.05	7.83	0.82	37.87
2 A/C	-2.33	-2.44	2.82	13.59	12.14	57.07
ave.	-1.23	-2.32	2.94	10.71	6.48	47.47
s.d.	1.55	0.16	0.16	4.08	8.00	13.58

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 A	-30.6	44.6	-40.6	24.5	3.4	44.2	156.6	C2'-endo
2 A	-37.4	47.0	-37.3	17.1	12.3	45.4	145.3	C2'-endo
3 C	-9.8	27.0	-33.0	28.2	-11.8	33.0	181.8	C3'-exo

5'-CCA-3' (PO- model) neutral / B3LYP-D

step overlap area (\AA^2)
 1 C/C 2.60
 2 C/A 2.42

 Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 C/C	-0.73	0.02	2.57	11.17	-11.76	29.18
2 C/A	0.84	0.73	3.05	1.82	1.33	49.93
ave.	0.06	0.37	2.81	6.49	-5.21	39.56
s.d.	1.11	0.50	0.33	6.61	9.25	14.67

 Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 C/C	1.50	2.67	2.02	-21.38	-20.31	33.30
2 C/A	0.77	-0.87	3.09	1.57	-2.15	49.98
ave.	1.14	0.90	2.56	-9.90	-11.23	41.64
s.d.	0.52	2.50	0.76	16.23	12.84	11.80

 Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 C	-17.2	34.3	-37.4	28.5	-7.2	37.7	172.2	C2'-endo
2 C	-37.0	45.2	-35.9	15.2	13.4	44.4	143.9	C1'-exo
3 A	-10.2	28.1	-34.1	29.2	-12.1	34.1	181.7	C3'-exo

5'-CCA-3' (PO- model) singly ionized / B3LYP-D

step overlap area (\AA^2)
 1 C/C 2.73
 2 C/A 4.52

 Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 C/C	-0.38	-0.08	2.72	8.22	-17.58	32.61
2 C/A	1.70	1.14	2.83	8.14	7.74	57.77
ave.	0.66	0.53	2.77	8.18	-4.92	45.19
s.d.	1.47	0.86	0.08	0.05	17.90	17.79

 Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 C/C	1.76	1.47	2.31	-28.38	-13.27	37.81
2 C/A	0.80	-1.35	3.14	7.93	-8.34	58.76
ave.	1.28	0.06	2.72	-10.23	-10.80	48.29
s.d.	0.68	1.99	0.59	25.68	3.49	14.81

 Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 C	-18.2	35.7	-38.7	29.3	-7.1	39.2	171.6	C2'-endo
2 C	-38.5	46.9	-36.4	15.1	14.4	45.7	142.9	C1'-exo
3 A	11.2	9.7	-25.2	32.2	-27.6	32.1	218.4	C4'-endo

5'-CAA-3' (PO- model) neutral / B3LYP-Dstep overlap area (\AA^2)

1	C/A	1.94
2	A/A	4.91

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 C/A	-0.07	0.70	2.68	11.14	-9.67	35.73
2 A/A	0.99	0.10	3.25	-3.66	8.81	48.46
ave.	0.46	0.40	2.97	3.74	-0.43	42.10
s.d.	0.75	0.42	0.40	10.47	13.07	9.00

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 C/A	2.01	1.20	2.30	-15.00	-17.28	38.57
2 A/A	-0.53	-1.45	3.15	10.62	4.41	49.34
ave.	0.74	-0.13	2.72	-2.19	-6.44	43.95
s.d.	1.80	1.87	0.60	18.11	15.34	7.62

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 C	-17.6	34.9	-38.0	28.9	-7.2	38.3	172.1	C2'-endo
2 A	-30.8	42.4	-37.3	20.2	6.4	42.0	152.6	C2'-endo
3 A	-10.2	28.1	-34.2	29.3	-12.2	34.2	181.7	C3'-exo

5'-CAA-3' (PO- model) singly ionized / B3LYP-Dstep overlap area (\AA^2)

1	C/A	0.00
2	A/A	5.75

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 C/A	-1.83	0.38	2.28	15.65	-28.18	21.60
2 A/A	1.67	-0.12	3.03	-0.20	9.14	54.97
ave.	-0.08	0.13	2.66	7.72	-9.52	38.28
s.d.	2.48	0.35	0.53	11.21	26.39	23.60

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 C/A	2.36	3.76	0.29	-49.87	-27.69	38.64
2 A/A	-0.62	-1.80	2.97	9.82	0.22	55.67
ave.	0.87	0.98	1.63	-20.02	-13.74	47.15
s.d.	2.11	3.93	1.90	42.21	19.73	12.04

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 C	-15.6	32.4	-35.9	27.9	-7.9	36.2	173.7	C2'-endo
2 A	-33.3	45.7	-39.3	21.4	7.0	44.6	151.9	C2'-endo
3 A	2.9	17.2	-29.2	31.7	-22.0	31.9	203.7	C3'-exo

5'-ACC-3' (H model) neutral / M06-2Xstep overlap area (\AA^2)

1	A/C	3.90
2	C/C	3.56

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 A/C	0.47	0.18	2.52	11.60	-0.45	33.60
2 C/C	0.06	0.02	2.71	7.22	-7.19	34.92
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ave.	0.26	0.10	2.62	9.41	-3.82	34.26
s.d.	0.28	0.12	0.14	3.10	4.77	0.93

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## Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 A/C	0.35	0.59	2.54	-0.75	-19.36	35.50
2 C/C	0.87	0.74	2.62	-11.67	-11.73	36.33
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ave.	0.61	0.66	2.58	-6.21	-15.54	35.92
s.d.	0.36	0.11	0.06	7.73	5.40	0.59

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 A	-15.2	33.4	-37.9	30.3	-9.5	38.0	175.7	C2'-endo
2 C	-45.0	47.4	-32.3	7.2	23.4	47.9	132.5	C1'-exo
3 C	-46.7	44.5	-25.5	-1.0	29.6	47.1	122.8	C1'-exo

5'-ACC-3' singly ionized H model / M06-2Xstep overlap area (\AA^2)

1	A/C	3.06
2	C/C	4.60

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 A/C	1.65	1.43	1.52	32.67	-5.82	43.81
2 C/C	0.23	0.63	3.26	-0.92	-2.20	34.51
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ave.	0.94	1.03	2.39	15.88	-4.01	39.16
s.d.	1.00	0.56	1.22	23.75	2.56	6.57

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## Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 A/C	1.78	-0.58	2.04	-6.77	-38.02	54.46
2 C/C	1.40	-0.52	3.20	-3.70	1.54	34.59
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ave.	1.59	-0.55	2.62	-5.23	-18.24	44.53
s.d.	0.27	0.04	0.82	2.17	27.97	14.05

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 A	-12.2	28.1	-32.1	26.1	-8.8	32.1	176.9	C2'-endo
2 C	-44.2	46.9	-32.3	7.7	22.5	47.2	133.2	C1'-exo
3 C	-46.1	45.3	-27.5	1.4	27.7	47.1	125.7	C1'-exo

5'-AAC-3' neutral H model / M06-2X

step overlap area (\AA^2)
 1 A/A 5.40
 2 A/C 4.17

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 A/A	2.78	2.10	3.01	17.61	-15.86	51.36
2 A/C	0.61	0.01	2.73	6.72	2.99	35.08
ave.	1.70	1.05	2.87	12.16	-6.44	43.22
s.d.	1.53	1.48	0.20	7.70	13.33	11.51

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 A/A	3.11	-1.97	3.02	-17.29	-19.20	56.22
2 A/C	-0.35	-0.17	2.79	4.90	-11.00	35.82
ave.	1.38	-1.07	2.91	-6.20	-15.10	46.02
s.d.	2.45	1.27	0.17	15.69	5.80	14.43

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 A	-12.8	29.7	-34.3	27.9	-9.6	34.4	177.3	C2'-endo
2 A	-42.1	46.6	-33.8	10.5	19.5	46.3	136.8	C1'-exo
3 C	-46.9	45.6	-27.0	0.4	28.7	47.6	124.5	C1'-exo

5'-AAC-3' singly ionized H model / M06-2X

step overlap area (\AA^2)
 1 A/A 4.98
 2 A/C 5.00

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 A/A	0.82	1.84	2.85	6.70	-1.43	37.23
2 A/C	1.66	0.88	2.42	12.15	-3.39	43.49
ave.	1.24	1.36	2.63	9.43	-2.41	40.36
s.d.	0.59	0.68	0.30	3.85	1.39	4.42

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 A/A	3.00	-0.51	2.88	-2.22	-10.39	37.84
2 A/C	1.39	-1.29	2.69	-4.47	-16.00	45.20
ave.	2.20	-0.90	2.79	-3.34	-13.19	41.52
s.d.	1.14	0.55	0.13	1.59	3.97	5.20

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 A	-4.0	25.2	-35.5	34.3	-19.1	36.3	192.5	C3'-exo
2 A	-40.6	46.4	-34.6	12.6	17.2	45.7	139.3	C1'-exo
3 C	-47.1	44.3	-24.9	-1.8	30.3	47.2	121.8	C1'-exo

5'-CCA-3' neutral / H model / M06-2X

step overlap area (\AA^2)
 1 C/C 4.98
 2 C/A 6.87

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 C/C	0.04	0.95	2.56	10.36	2.00	29.72
2 C/A	1.84	0.99	2.78	8.85	-2.97	44.93
ave.	0.94	0.97	2.67	9.60	-0.49	37.32
s.d.	1.27	0.03	0.16	1.07	3.52	10.76

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 C/C	1.44	1.47	2.49	3.75	-19.44	31.49
2 C/A	1.50	-1.67	3.00	-3.84	-11.43	45.84
ave.	1.47	-0.10	2.75	-0.04	-15.44	38.67
s.d.	0.04	2.22	0.36	5.37	5.66	10.15

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 C	-34.0	42.6	-35.3	16.6	10.7	42.1	146.9	C2'-endo
2 C	-43.2	47.6	-34.3	10.3	20.3	47.5	136.3	C1'-exo
3 A	-44.9	45.0	-28.1	2.9	26.0	46.2	127.4	C1'-exo

5'-CCA-3' singly ionized / H model / M06-2X

step overlap area (\AA^2)
 1 C/C 4.90
 2 C/A 1.41

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 C/C	0.22	0.43	2.77	6.55	-1.44	32.14
2 C/A	0.30	0.10	2.30	13.69	-11.09	38.66
ave.	0.26	0.26	2.53	10.12	-6.26	35.40
s.d.	0.05	0.23	0.33	5.05	6.83	4.61

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 C/C	0.97	0.56	2.74	-2.56	-11.68	32.82
2 C/A	0.95	0.59	2.18	-15.79	-19.49	42.35
ave.	0.96	0.58	2.46	-9.18	-15.58	37.58
s.d.	0.01	0.02	0.40	9.36	5.53	6.74

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 C	-35.2	43.6	-35.7	16.4	11.6	43.0	146.1	C2'-endo
2 C	-45.3	48.3	-33.2	8.1	23.0	48.5	133.3	C1'-exo
3 A	-43.9	45.6	-29.6	5.3	23.7	45.9	130.2	C1'-exo

5'-CAA-3' neutral / H model / M06-2X

step overlap area (\AA^2)
 1 C/A 1.45
 2 A/A 4.68

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 C/A	-0.59	0.49	2.48	17.64	-19.41	27.27
2 A/A	1.01	0.07	3.17	-2.30	5.94	51.11
ave.	0.21	0.28	2.83	7.67	-6.73	39.19
s.d.	1.13	0.30	0.49	14.10	17.93	16.86

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 C/A	2.47	2.47	1.29	-33.05	-30.04	37.66
2 A/A	-0.31	-1.32	3.11	6.86	2.66	51.48
ave.	1.08	0.58	2.20	-13.10	-13.69	44.57
s.d.	1.97	2.68	1.29	28.22	23.12	9.77

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 C	-18.7	33.6	-35.1	25.3	-4.2	35.9	168.1	C2'-endo
2 A	-42.3	46.5	-33.2	9.7	20.1	46.2	135.8	C1'-exo
3 A	-17.3	32.5	-34.5	25.6	-5.4	35.1	170.0	C2'-endo

5'-CAA-3' singly ionized H model / M06-2X

step overlap area (\AA^2)
 1 C/A 0.00
 2 A/A 3.83

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 C/A	-1.63	0.61	1.82	29.17	-30.96	25.91
2 A/A	1.35	2.11	2.85	5.87	-0.18	55.91
ave.	-0.14	1.36	2.33	17.52	-15.57	40.91
s.d.	2.10	1.06	0.73	16.47	21.76	21.22

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 C/A	1.43	2.61	-0.34	-43.71	-41.19	49.49
2 A/A	2.25	-1.14	2.96	-0.19	-6.24	56.19
ave.	1.84	0.73	1.31	-21.95	-23.72	52.84
s.d.	0.58	2.65	2.33	30.77	24.71	4.74

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 C	-18.6	33.3	-34.9	25.1	-4.2	35.6	168.1	C2'-endo
2 A	-39.3	46.3	-35.8	14.6	15.1	45.4	141.9	C1'-exo
3 A	-17.7	32.7	-34.4	25.3	-4.9	35.0	169.2	C2'-endo

5'-ACC-3' (Na model) neutral / M06-2X

step overlap area (\AA^2)
 1 A/C 3.78
 2 C/C 2.86

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 A/C	0.43	0.07	2.65	9.57	-0.47	34.64
2 C/C	0.15	0.27	2.92	3.43	-7.12	42.91
ave.	0.29	0.17	2.78	6.50	-3.80	38.78
s.d.	0.20	0.14	0.20	4.34	4.71	5.85

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 A/C	0.16	0.44	2.67	-0.77	-15.70	35.90
2 C/C	0.97	0.09	2.85	-9.64	-4.64	43.60
ave.	0.57	0.26	2.76	-5.20	-10.17	39.75
s.d.	0.57	0.24	0.13	6.27	7.82	5.45

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 A	-14.2	33.6	-39.0	32.1	-11.4	39.1	178.0	C2'-endo
2 C	-42.4	48.1	-35.5	12.0	18.8	47.5	138.3	C1'-exo
3 C	-24.0	36.5	-34.5	21.7	1.3	36.9	159.3	C2'-endo

5'-ACC-3' (Na model) singly ionized / M06-2X

step overlap area (\AA^2)
 1 A/C 4.51
 2 C/C 3.32

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 A/C	1.44	0.98	2.16	17.73	-3.77	42.02
2 C/C	0.27	0.59	3.04	1.87	-4.68	43.08
ave.	0.86	0.78	2.60	9.80	-4.22	42.55
s.d.	0.83	0.28	0.63	11.22	0.65	0.75

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 A/C	1.51	-0.71	2.45	-4.98	-23.46	45.60
2 C/C	1.21	-0.20	2.98	-6.35	-2.54	43.36
ave.	1.36	-0.46	2.71	-5.67	-13.00	44.48
s.d.	0.21	0.36	0.37	0.96	14.80	1.58

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 A	-8.7	30.2	-38.8	35.3	-16.7	39.0	186.3	C3'-exo
2 C	-41.0	48.1	-37.0	14.6	16.2	47.3	141.5	C1'-exo
3 C	-24.2	36.7	-34.7	21.8	1.4	37.1	159.2	C2'-endo

5'-AAC-3' (Na model) neutral / M06-2X

step overlap area (\AA^2)
 1 A/A 5.05
 2 A/C 3.60

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 A/A	0.20	1.31	2.94	3.61	10.45	26.42
2 A/C	0.82	0.12	2.76	6.87	1.01	44.68
ave.	0.51	0.72	2.85	5.24	5.73	35.55
s.d.	0.44	0.84	0.13	2.30	6.68	12.91

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 A/A	0.33	0.39	3.22	21.70	-7.50	28.61
2 A/C	0.08	-0.54	2.85	1.32	-8.97	45.19
ave.	0.20	-0.07	3.03	11.51	-8.23	36.90
s.d.	0.18	0.66	0.26	14.41	1.04	11.72

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 A	-27.3	39.7	-36.6	22.1	3.0	39.8	157.0	C2'-endo
2 A	-36.3	45.6	-37.2	17.6	11.5	44.7	146.5	C2'-endo
3 C	-25.6	37.6	-34.7	21.0	2.7	37.7	157.3	C2'-endo

5'-AAC-3' (Na model) singly ionized / M06-2X

step overlap area (\AA^2)
 1 A/A 5.43
 2 A/C 5.31

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 A/A	0.86	1.58	2.88	4.85	2.12	35.70
2 A/C	1.53	0.55	2.44	12.07	-2.15	42.49
ave.	1.20	1.06	2.66	8.46	-0.02	39.10
s.d.	0.48	0.72	0.31	5.11	3.02	4.80

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 A	-21.8	37.5	-38.2	27.0	-3.5	39.4	166.2	C2'-endo
2 A	-37.7	47.0	-38.0	17.8	12.0	45.8	146.0	C2'-endo
3 C	-45.2	45.7	-28.8	3.6	25.7	46.7	128.1	C1'-exo

5'-CCA-3' (Na model) neutral / M06-2X

step overlap area (\AA^2)

1	C/C	1.89
2	C/A	1.50

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 C/C	-0.84	-0.37	2.33	13.77	-16.10	26.71
2 C/A	0.68	0.54	2.95	1.98	-3.62	50.16

ave.	-0.08	0.08	2.64	7.87	-9.86	38.43
s.d.	1.07	0.65	0.44	8.34	8.82	16.58

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 C/C	1.14	2.97	1.68	-29.63	-25.34	33.97
2 C/A	0.87	-0.67	2.93	-4.26	-2.33	50.31

ave.	1.01	1.15	2.30	-16.95	-13.84	42.14
s.d.	0.19	2.57	0.88	17.94	16.27	11.56

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 C	-19.8	36.2	-38.0	27.7	-5.0	38.8	168.8	C2'-endo
2 C	-42.4	48.4	-35.8	12.4	18.5	47.7	138.7	C1'-exo
3 A	-14.7	31.0	-34.5	27.0	-7.9	34.7	174.2	C2'-endo

5'-CCA-3' (Na model) singly ionized / M06-2X

step overlap area (\AA^2)

1	C/C	1.38
2	C/A	1.64

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 C/C	-1.10	-0.54	2.62	9.87	-20.35	24.52
2 C/A	1.07	0.36	2.41	11.26	-11.19	54.34

ave.	-0.01	-0.09	2.51	10.56	-15.77	39.43
s.d.	1.53	0.64	0.15	0.98	6.47	21.08

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 C/C	1.81	3.26	1.95	-38.94	-18.88	33.24
2 C/A	0.86	-0.65	2.46	-11.97	-12.04	56.44

ave.	1.34	1.30	2.21	-25.45	-15.46	44.84
s.d.	0.67	2.76	0.36	19.07	4.84	16.41

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 C	-20.1	36.2	-37.8	27.3	-4.6	38.7	168.1	C2'-endo
2 C	-42.8	47.7	-34.2	10.4	20.0	47.2	136.4	C1'-exo
3 A	-10.7	28.1	-33.6	28.5	-11.4	33.6	180.6	C3'-exo

5'-CAA-3' (Na model) neutral / M06-2Xstep overlap area (\AA^2)

1	C/A	0.87
2	A/A	4.79

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 C/A	-0.72	0.11	2.46	13.44	-15.83	28.77
2 A/A	1.10	-0.03	3.17	-2.44	4.59	51.63
ave.	0.19	0.04	2.81	5.50	-5.62	40.20
s.d.	1.28	0.10	0.50	11.23	14.44	16.16

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 C/A	1.83	2.58	1.69	-27.76	-23.57	35.35
2 A/A	-0.33	-1.41	3.10	5.25	2.80	51.87
ave.	0.75	0.59	2.40	-11.25	-10.39	43.61
s.d.	1.53	2.82	1.00	23.35	18.64	11.68

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 C	-19.7	36.0	-37.8	27.6	-5.0	38.5	168.7	C2'-endo
2 A	-35.9	45.1	-36.8	17.1	11.6	44.2	146.4	C2'-endo
3 A	-14.1	30.7	-34.5	27.4	-8.5	34.7	175.2	C2'-endo

5'-CAA-3' (Na model) singly ionized / M06-2Xstep overlap area (\AA^2)

1	C/A	0.00
2	A/A	5.61

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 C/A	-1.72	0.64	2.17	18.97	-31.63	25.46
2 A/A	1.38	-0.12	3.04	-0.69	5.24	50.27
ave.	-0.17	0.26	2.60	9.14	-13.20	37.86
s.d.	2.19	0.54	0.62	13.91	26.07	17.55

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 C/A	2.16	3.05	0.09	-48.49	-29.09	44.56
2 A/A	-0.48	-1.66	3.00	6.14	0.81	50.53
ave.	0.84	0.70	1.54	-21.17	-14.14	47.55
s.d.	1.87	3.33	2.06	38.64	21.14	4.22

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 C	-19.8	34.9	-36.2	25.9	-3.9	37.1	167.4	C2'-endo
2 A	-30.2	42.1	-37.2	21.0	5.5	41.5	153.6	C2'-endo
3 A	-15.2	31.3	-34.5	26.8	-7.5	34.7	173.4	C2'-endo

5'-ACC-3' (PO- model) neutral / M06-2X

step overlap area (\AA^2)
 1 A/C 4.22
 2 C/C 1.07

 Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 A/C	0.59	-0.08	2.76	6.37	1.70	34.48
2 C/C	1.08	-1.15	4.34	-19.31	14.17	48.66

 ~~~~~  
 ave. 0.83 -0.61 3.55 -6.47 7.93 41.57  
 s.d. 0.35 0.76 1.12 18.16 8.82 10.03  
 \*\*\*\*  
 Local base helical parameters  

| step  | X-disp | Y-disp | h-Rise | Incl. | Tip    | h-Twist |
|-------|--------|--------|--------|-------|--------|---------|
| 1 A/C | -0.35  | -0.15  | 2.81   | 2.84  | -10.63 | 35.08   |
| 2 C/C | -2.36  | -2.68  | 3.30   | 16.11 | 21.96  | 53.92   |

 ~~~~~  
 ave. -1.35 -1.41 3.06 9.47 5.66 44.50
 s.d. 1.43 1.79 0.35 9.39 23.05 13.32

 Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 A	-26.1	40.9	-39.4	25.7	-0.0	41.6	161.2	C2'-endo
2 C	-43.0	45.8	-30.8	6.7	22.5	45.9	132.2	C1'-exo
3 C	-16.8	32.3	-34.7	25.9	-5.9	35.1	170.7	C2'-endo

5'-ACC-3' (PO- model) singly ionized / M06-2X

step overlap area (\AA^2)
 1 A/C 1.74
 2 C/C 0.19

 Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 A/C	-1.15	-0.30	2.81	7.85	-19.51	25.54
2 C/C	1.06	-1.52	5.11	-31.63	17.04	47.84

 ~~~~~  
 ave. -0.04 -0.91 3.96 -11.89 -1.24 36.69  
 s.d. 1.56 0.87 1.63 27.92 25.85 15.77  
 \*\*\*\*  
 Local base helical parameters  

| step  | X-disp | Y-disp | h-Rise | Incl.  | Tip    | h-Twist |
|-------|--------|--------|--------|--------|--------|---------|
| 1 A/C | 2.29   | 3.16   | 2.09   | -37.13 | -14.94 | 32.98   |
| 2 C/C | -2.80  | -3.41  | 3.25   | 18.33  | 34.02  | 59.19   |

 ~~~~~  
 ave. -0.25 -0.12 2.67 -9.40 9.54 46.08
 s.d. 3.60 4.65 0.82 39.21 34.62 18.53

 Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 A	-12.9	32.1	-37.9	31.8	-12.0	37.9	179.4	C2'-endo
2 C	-41.1	47.8	-35.5	12.8	17.4	46.7	139.5	C1'-exo
3 C	-14.1	30.3	-34.1	27.0	-8.3	34.3	175.1	C2'-endo

5'-AAC-3' (PO- model) neutral / M06-2X

step overlap area (\AA^2)
 1 A/A 5.78
 2 A/C 4.92

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 A/A	0.74	0.49	3.19	-0.68	7.64	37.10
2 A/C	1.17	-1.38	4.05	-16.08	20.29	49.33
ave.	0.95	-0.44	3.62	-8.38	13.96	43.22
s.d.	0.31	1.32	0.61	10.89	8.94	8.65

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 A/A	-0.21	-1.23	3.22	11.85	1.06	37.86
2 A/C	-2.83	-2.33	2.85	22.60	17.92	55.34
ave.	-1.52	-1.78	3.03	17.23	9.49	46.60
s.d.	1.85	0.78	0.26	7.60	11.92	12.36

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 A	-31.7	44.1	-39.2	22.4	5.5	43.7	154.0	C2'-endo
2 A	-34.1	45.1	-38.0	19.5	8.9	44.1	149.6	C2'-endo
3 C	-12.8	29.0	-33.3	26.9	-9.1	33.4	176.8	C2'-endo

5'-AAC-3' (PO- model) singly ionized / M06-2X

step overlap area (\AA^2)
 1 A/A 7.04
 2 A/C 4.97

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 A/A	1.53	0.34	3.05	0.20	4.99	41.93
2 A/C	1.81	0.05	2.62	10.82	1.32	55.74
ave.	1.67	0.19	2.83	5.51	3.15	48.84
s.d.	0.20	0.20	0.30	7.51	2.59	9.77

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 A/A	-0.01	-2.10	3.07	6.94	-0.28	42.21
2 A/C	-0.01	-1.39	2.90	1.40	-11.45	56.72
ave.	-0.01	-1.75	2.99	4.17	-5.87	49.46
s.d.	0.00	0.50	0.12	3.92	7.90	10.25

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 A	-32.1	45.5	-40.9	24.0	4.7	45.1	155.2	C2'-endo
2 A	-39.2	47.4	-36.3	15.1	14.6	45.9	142.4	C1'-exo
3 C	2.4	18.0	-30.1	32.5	-22.2	32.7	202.9	C3'-exo

5'-CCA-3' (PO- model) neutral / M06-2X

step overlap area (\AA^2)
 1 C/C 1.86
 2 C/A 1.85

 Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 C/C	-0.84	-0.30	2.42	12.27	-15.92	28.06
2 C/A	0.78	0.59	2.98	2.19	-0.86	50.39
ave.	-0.03	0.14	2.70	7.23	-8.39	39.22
s.d.	1.14	0.63	0.39	7.13	10.65	15.79

 Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 C/C	1.28	2.80	1.83	-28.65	-22.08	34.40
2 C/A	0.74	-0.77	3.00	-1.01	-2.57	50.44
ave.	1.01	1.01	2.41	-14.83	-12.33	42.42
s.d.	0.38	2.53	0.82	19.54	13.79	11.34

 Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 C	-20.8	37.0	-38.4	27.6	-4.4	39.3	167.7	C2'-endo
2 C	-38.7	47.1	-37.4	16.1	13.9	46.2	144.0	C2'-endo
3 A	-11.8	29.2	-34.4	28.6	-10.8	34.5	179.1	C2'-endo

5'-CCA-3' (PO- model) singly ionized / M06-2X

step overlap area (\AA^2)
 1 C/C 1.37
 2 C/A 2.05

 Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 C/C	-1.07	-0.54	2.72	7.56	-18.21	25.19
2 C/A	1.22	0.42	2.43	11.90	-6.02	55.41
ave.	0.08	-0.06	2.58	9.73	-12.11	40.30
s.d.	1.62	0.68	0.20	3.07	8.62	21.37

 Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 C/C	1.83	3.11	2.21	-35.60	-14.78	31.89
2 C/A	0.70	-0.77	2.57	-6.37	-12.59	56.87
ave.	1.27	1.17	2.39	-20.98	-13.68	44.38
s.d.	0.80	2.75	0.25	20.67	1.54	17.66

 Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 C	-20.4	36.6	-38.1	27.4	-4.5	38.9	167.9	C2'-endo
2 C	-42.9	47.5	-33.4	9.5	20.6	46.9	135.3	C1'-exo
3 A	10.1	11.6	-27.0	33.5	-27.7	33.2	215.7	C3'-exo

5'-CAA-3' (PO- model) neutral / M06-2X

step overlap area (\AA^2)
 1 C/A 0.72
 2 A/A 4.81

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 C/A	-0.78	0.18	2.45	13.79	-17.19	29.34
2 A/A	1.08	0.00	3.18	-2.47	5.98	50.96
ave.	0.15	0.09	2.82	5.66	-5.60	40.15
s.d.	1.32	0.13	0.51	11.50	16.38	15.29

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 C/A	1.93	2.57	1.61	-29.29	-23.50	36.55
2 A/A	-0.40	-1.40	3.11	6.92	2.86	51.34
ave.	0.77	0.58	2.36	-11.19	-10.32	43.94
s.d.	1.64	2.81	1.06	25.60	18.64	10.46

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 C	-20.6	36.8	-38.2	27.5	-4.4	39.1	167.7	C2'-endo
2 A	-33.1	44.4	-38.2	20.1	8.0	43.7	150.9	C2'-endo
3 A	-12.7	29.9	-34.6	28.4	-10.0	34.7	177.8	C2'-endo

5'-CAA-3' (PO- model) singly ionized / M06-2X

step overlap area (\AA^2)
 1 C/A 0.00
 2 A/A 5.16

Local base step parameters

step	Shift	Slide	Rise	Tilt	Roll	Twist
1 C/A	-1.85	0.68	2.38	15.49	-35.03	25.04
2 A/A	1.31	-0.15	2.97	1.72	4.33	52.90
ave.	-0.27	0.27	2.67	8.61	-15.35	38.97
s.d.	2.23	0.59	0.42	9.74	27.83	19.70

Local base helical parameters

step	X-disp	Y-disp	h-Rise	Incl.	Tip	h-Twist
1 C/A	2.50	3.10	0.20	-53.07	-23.47	45.50
2 A/A	-0.42	-1.37	2.99	4.85	-1.92	53.09
ave.	1.04	0.87	1.59	-24.11	-12.70	49.30
s.d.	2.06	3.16	1.97	40.95	15.24	5.37

Sugar conformational parameters:

base	v0	v1	v2	v3	v4	tm	P	Puckering
1 C	-20.2	35.6	-36.8	26.3	-3.9	37.7	167.3	C2'-endo
2 A	-22.7	39.6	-40.0	28.4	-3.9	41.2	166.2	C2'-endo
3 A	1.3	18.5	-29.7	31.3	-20.7	31.8	200.8	C3'-exo

Optimized Geometries. Cartesian coordinates (\AA), PDB format

COMPND ACC / M06-2X / H MODEL / NEUTRAL

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ATOM  1 O5' DA A 1  0.000  0.000  0.000  1.00  0.00   O
ATOM  2 C5' DA A 1  0.000  0.000  1.415  1.00  0.00   C
ATOM  3 C4' DA A 1  1.436  0.000  1.905  1.00  0.00   C
ATOM  4 O4' DA A 1  2.090 -1.207  1.507  1.00  0.00   O
ATOM  5 C3' DA A 1  2.295  1.128  1.326  1.00  0.00   C
ATOM  6 O3' DA A 1  3.346  1.362  2.298  1.00  0.00   O
ATOM  7 C2' DA A 1  2.872  0.494  0.076  1.00  0.00   C
ATOM  8 C1' DA A 1  3.122 -0.927  0.571  1.00  0.00   C
ATOM  9 N9 DA A 1  3.128 -1.919 -0.479  1.00  0.00   N
ATOM 10 C8 DA A 1  2.477 -1.914 -1.693  1.00  0.00   C
ATOM 11 N7 DA A 1  2.798 -2.920 -2.466  1.00  0.00   N
ATOM 12 C5 DA A 1  3.703 -3.643 -1.713  1.00  0.00   C
ATOM 13 C6 DA A 1  4.483 -4.782 -1.986  1.00  0.00   C
ATOM 14 N6 DA A 1  4.502 -5.355 -3.214  1.00  0.00   N
ATOM 15 N1 DA A 1  5.280 -5.265 -1.024  1.00  0.00   N
ATOM 16 C2 DA A 1  5.358 -4.595  0.135  1.00  0.00   C
ATOM 17 N3 DA A 1  4.730 -3.479  0.493  1.00  0.00   N
ATOM 18 C4 DA A 1  3.915 -3.046 -0.475  1.00  0.00   C
ATOM 19 P DC A 2  4.449  2.494  2.078  1.00  0.00   P
ATOM 20 OP1 DC A 2  3.747  3.891  2.384  1.00  0.00   O
ATOM 21 OP2 DC A 2  5.104  2.544  0.755  1.00  0.00   O
ATOM 22 O5' DC A 2  5.388  2.268  3.338  1.00  0.00   O
ATOM 23 C5' DC A 2  5.607  1.010  4.001  1.00  0.00   C
ATOM 24 C4' DC A 2  6.725  0.219  3.361  1.00  0.00   C
ATOM 25 O4' DC A 2  6.268 -0.392  2.157  1.00  0.00   O
ATOM 26 C3' DC A 2  7.968  1.059  2.970  1.00  0.00   C
ATOM 27 O3' DC A 2  9.104  0.373  3.539  1.00  0.00   O
ATOM 28 C2' DC A 2  7.983  0.966  1.449  1.00  0.00   C
ATOM 29 C1' DC A 2  7.401 -0.435  1.282  1.00  0.00   C
ATOM 30 N1 DC A 2  6.977 -0.850 -0.031  1.00  0.00   N
ATOM 31 C2 DC A 2  7.334 -2.158 -0.472  1.00  0.00   C
ATOM 32 O2 DC A 2  8.010 -2.873  0.259  1.00  0.00   O
ATOM 33 N3 DC A 2  6.897 -2.550 -1.700  1.00  0.00   N
ATOM 34 C4 DC A 2  6.163 -1.745 -2.444  1.00  0.00   C
ATOM 35 N4 DC A 2  5.790 -2.190 -3.666  1.00  0.00   N
ATOM 36 C5 DC A 2  5.786 -0.422 -2.029  1.00  0.00   C
ATOM 37 C6 DC A 2  6.219 -0.027 -0.808  1.00  0.00   C
ATOM 38 P DC A 3  10.570  1.002  3.484  1.00  0.00   P
ATOM 39 OP1 DC A 3  10.521  2.344  4.344  1.00  0.00   O
ATOM 40 OP2 DC A 3  11.130  1.288  2.148  1.00  0.00   O
ATOM 41 O5' DC A 3  11.377 -0.008  4.406  1.00  0.00   O
ATOM 42 C5' DC A 3  11.026 -1.388  4.610  1.00  0.00   C
ATOM 43 C4' DC A 3  11.498 -2.267  3.478  1.00  0.00   C
ATOM 44 O4' DC A 3  10.720 -2.022  2.309  1.00  0.00   O
ATOM 45 C3' DC A 3  12.984 -2.084  3.073  1.00  0.00   C
ATOM 46 O3' DC A 3  13.618 -3.328  3.312  1.00  0.00   O
ATOM 47 C2' DC A 3  12.889 -1.749  1.582  1.00  0.00   C
ATOM 48 C1' DC A 3  11.563 -2.411  1.224  1.00  0.00   C
ATOM 49 N1 DC A 3  10.963 -2.021 -0.031  1.00  0.00   N
ATOM 50 C2 DC A 3  10.721 -3.008 -1.022  1.00  0.00   C
ATOM 51 O2 DC A 3  11.086 -4.163 -0.819  1.00  0.00   O
ATOM 52 N3 DC A 3  10.116 -2.613 -2.177  1.00  0.00   N
ATOM 53 C4 DC A 3  9.735 -1.358 -2.337  1.00  0.00   C
ATOM 54 N4 DC A 3  9.128 -1.028 -3.498  1.00  0.00   N
ATOM 55 C5 DC A 3  9.969 -0.337 -1.356  1.00  0.00   C
ATOM 56 C6 DC A 3  10.589 -0.728 -0.217  1.00  0.00   C
END

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COMPND ACC / M06-2X / H MODEL / SINGLY IONIZED

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ATOM  1 O5' DA A 1  0.000  0.000  0.000  1.00  0.00   O
ATOM  2 C5' DA A 1  0.000  0.000  1.418  1.00  0.00   C
ATOM  3 C4' DA A 1  1.441  0.000  1.890  1.00  0.00   C
ATOM  4 O4' DA A 1  2.067 -1.240  1.524  1.00  0.00   O
ATOM  5 C3' DA A 1  2.285  1.102  1.242  1.00  0.00   C
ATOM  6 O3' DA A 1  3.317  1.478  2.184  1.00  0.00   O
ATOM  7 C2' DA A 1  2.913  0.402  0.054  1.00  0.00   C
ATOM  8 C1' DA A 1  3.072 -1.025  0.568  1.00  0.00   C
ATOM  9 N9 DA A 1  2.943 -1.997 -0.521  1.00  0.00   N
ATOM 10 C8 DA A 1  1.930 -2.114 -1.407  1.00  0.00   C
ATOM 11 N7 DA A 1  2.165 -3.001 -2.391  1.00  0.00   N
ATOM 12 C5 DA A 1  3.388 -3.462 -2.120  1.00  0.00   C
ATOM 13 C6 DA A 1  4.254 -4.396 -2.781  1.00  0.00   C
ATOM 14 N6 DA A 1  3.898 -5.004 -3.889  1.00  0.00   N
ATOM 15 N1 DA A 1  5.470 -4.652 -2.239  1.00  0.00   N
ATOM 16 C2 DA A 1  5.808 -4.033 -1.132  1.00  0.00   C
ATOM 17 N3 DA A 1  5.075 -3.136 -0.389  1.00  0.00   N
ATOM 18 C4 DA A 1  3.926 -2.878 -0.933  1.00  0.00   C
ATOM 19 P DC A 2  4.239  2.745  1.849  1.00  0.00   P
ATOM 20 OP1 DC A 2  3.611  3.992  2.611  1.00  0.00   O
ATOM 21 OP2 DC A 2  4.403  3.051  0.415  1.00  0.00   O
ATOM 22 O5' DC A 2  5.547  2.441  2.701  1.00  0.00   O
ATOM 23 C5' DC A 2  6.086  1.134  2.957  1.00  0.00   C
ATOM 24 C4' DC A 2  6.984  0.658  1.839  1.00  0.00   C
ATOM 25 O4' DC A 2  6.196  0.254  0.723  1.00  0.00   O
ATOM 26 C3' DC A 2  7.973  1.723  1.300  1.00  0.00   C
ATOM 27 O3' DC A 2  9.268  1.092  1.314  1.00  0.00   O
ATOM 28 C2' DC A 2  7.488  1.963 -0.127  1.00  0.00   C
ATOM 29 C1' DC A 2  6.967  0.568 -0.445  1.00  0.00   C
ATOM 30 N1 DC A 2  6.133  0.373 -1.607  1.00  0.00   N
ATOM 31 C2 DC A 2  6.150 -0.921 -2.200  1.00  0.00   C
ATOM 32 O2 DC A 2  7.004 -1.727 -1.843  1.00  0.00   O
ATOM 33 N3 DC A 2  5.182 -1.221 -3.109  1.00  0.00   N
ATOM 34 C4 DC A 2  4.346 -0.274 -3.526  1.00  0.00   C
ATOM 35 N4 DC A 2  3.394 -0.631 -4.397  1.00  0.00   N
ATOM 36 C5 DC A 2  4.420  1.086 -3.072  1.00  0.00   C
ATOM 37 C6 DC A 2  5.305  1.346 -2.083  1.00  0.00   C
ATOM 38 P DC A 3  10.600  1.887  0.942  1.00  0.00   P
ATOM 39 OP1 DC A 3  10.754  3.043  2.031  1.00  0.00   O
ATOM 40 OP2 DC A 3  10.681  2.469 -0.413  1.00  0.00   O
ATOM 41 O5' DC A 3  11.719  0.832  1.333  1.00  0.00   O
ATOM 42 C5' DC A 3  11.509 -0.587  1.456  1.00  0.00   C
ATOM 43 C4' DC A 3  11.576 -1.275  0.115  1.00  0.00   C
ATOM 44 O4' DC A 3  10.436 -0.925 -0.665  1.00  0.00   O
ATOM 45 C3' DC A 3  12.824 -0.924 -0.737  1.00  0.00   C
ATOM 46 O3' DC A 3  13.497 -2.146 -0.984  1.00  0.00   O
ATOM 47 C2' DC A 3  12.210 -0.330 -2.007  1.00  0.00   C
ATOM 48 C1' DC A 3  10.863 -1.044 -2.022  1.00  0.00   C
ATOM 49 N1 DC A 3  9.847 -0.518 -2.904  1.00  0.00   N

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ATOM 50 C2 DC A 3 9.219 -1.392 -3.834 1.00 0.00 C
 ATOM 51 O2 DC A 3 9.577 -2.563 -3.893 1.00 0.00 O
 ATOM 52 N3 DC A 3 8.257 -0.868 -4.643 1.00 0.00 N
 ATOM 53 C4 DC A 3 7.906 0.402 -4.533 1.00 0.00 C
 ATOM 54 N4 DC A 3 6.938 0.854 -5.352 1.00 0.00 N
 ATOM 55 C5 DC A 3 8.514 1.307 -3.598 1.00 0.00 C
 ATOM 56 C6 DC A 3 9.481 0.790 -2.805 1.00 0.00 C
 END

COMPND AAC / M06-2X / H MODEL / NEUTRAL

ATOM 1 O5' DA A 1 0.000 0.000 0.000 1.00 0.00 O
 ATOM 2 C5' DA A 1 0.000 0.000 1.416 1.00 0.00 C
 ATOM 3 C4' DA A 1 1.437 0.000 1.902 1.00 0.00 C
 ATOM 4 O4' DA A 1 2.071 -1.237 1.568 1.00 0.00 O
 ATOM 5 C3' DA A 1 2.297 1.090 1.257 1.00 0.00 C
 ATOM 6 O3' DA A 1 3.333 1.423 2.217 1.00 0.00 O
 ATOM 7 C2' DA A 1 2.913 0.380 0.067 1.00 0.00 C
 ATOM 8 C1' DA A 1 3.098 -1.035 0.609 1.00 0.00 C
 ATOM 9 N9 DA A 1 3.019 -2.051 -0.417 1.00 0.00 N
 ATOM 10 C8 DA A 1 2.054 -2.195 -1.389 1.00 0.00 C
 ATOM 11 N7 DA A 1 2.277 -3.195 -2.203 1.00 0.00 N
 ATOM 12 C5 DA A 1 3.452 -3.750 -1.736 1.00 0.00 C
 ATOM 13 C6 DA A 1 4.243 -4.830 -2.176 1.00 0.00 C
 ATOM 14 N6 DA A 1 3.911 -5.571 -3.252 1.00 0.00 N
 ATOM 15 N1 DA A 1 5.353 -5.137 -1.487 1.00 0.00 N
 ATOM 16 C2 DA A 1 5.665 -4.395 -0.412 1.00 0.00 C
 ATOM 17 N3 DA A 1 5.020 -3.351 0.094 1.00 0.00 N
 ATOM 18 C4 DA A 1 3.923 -3.066 -0.619 1.00 0.00 C
 ATOM 19 P DA A 2 4.256 2.702 1.966 1.00 0.00 P
 ATOM 20 OP1 DA A 2 3.596 3.918 2.756 1.00 0.00 O
 ATOM 21 OP2 DA A 2 4.470 3.074 0.554 1.00 0.00 O
 ATOM 22 O5' DA A 2 5.539 2.372 2.849 1.00 0.00 O
 ATOM 23 C5' DA A 2 6.015 1.050 3.147 1.00 0.00 C
 ATOM 24 C4' DA A 2 6.918 0.525 2.055 1.00 0.00 C
 ATOM 25 O4' DA A 2 6.131 0.224 0.904 1.00 0.00 O
 ATOM 26 C3' DA A 2 8.006 1.518 1.580 1.00 0.00 C
 ATOM 27 O3' DA A 2 9.228 0.755 1.520 1.00 0.00 O
 ATOM 28 C2' DA A 2 7.543 1.895 0.179 1.00 0.00 C
 ATOM 29 C1' DA A 2 6.919 0.570 -0.238 1.00 0.00 C
 ATOM 30 N9 DA A 2 6.099 0.590 -1.412 1.00 0.00 N
 ATOM 31 C8 DA A 2 5.382 1.633 -1.964 1.00 0.00 C
 ATOM 32 N7 DA A 2 4.776 1.315 -3.078 1.00 0.00 N
 ATOM 33 C5 DA A 2 5.113 -0.011 -3.280 1.00 0.00 C
 ATOM 34 C6 DA A 2 4.862 -0.910 -4.335 1.00 0.00 C
 ATOM 35 N6 DA A 2 4.184 -0.532 -5.446 1.00 0.00 N
 ATOM 36 N1 DA A 2 5.372 -2.146 -4.266 1.00 0.00 N
 ATOM 37 C2 DA A 2 6.166 -2.450 -3.228 1.00 0.00 C
 ATOM 38 N3 DA A 2 6.514 -1.686 -2.196 1.00 0.00 N
 ATOM 39 C4 DA A 2 5.943 -0.479 -2.265 1.00 0.00 C
 ATOM 40 P DC A 3 10.622 1.416 1.124 1.00 0.00 P
 ATOM 41 OP2 DC A 3 10.713 2.092 -0.189 1.00 0.00 O
 ATOM 42 OP1 DC A 3 10.944 2.394 2.345 1.00 0.00 O
 ATOM 43 O5' DC A 3 11.614 0.205 1.369 1.00 0.00 O
 ATOM 44 C5' DC A 3 11.241 -1.182 1.287 1.00 0.00 C
 ATOM 45 C4' DC A 3 11.233 -1.679 -0.137 1.00 0.00 C
 ATOM 46 O4' DC A 3 10.133 -1.119 -0.852 1.00 0.00 O
 ATOM 47 C3' DC A 3 12.511 -1.346 -0.950 1.00 0.00 C
 ATOM 48 O3' DC A 3 13.071 -2.587 -1.340 1.00 0.00 O
 ATOM 49 C2' DC A 3 11.959 -0.555 -2.139 1.00 0.00 C
 ATOM 50 C1' DC A 3 10.550 -1.130 -2.218 1.00 0.00 C
 ATOM 51 N1 DC A 3 9.595 -0.419 -3.038 1.00 0.00 N
 ATOM 52 C2 DC A 3 8.961 -1.110 -4.107 1.00 0.00 C
 ATOM 53 O2 DC A 3 9.236 -2.288 -4.310 1.00 0.00 O
 ATOM 54 N3 DC A 3 8.075 -0.413 -4.872 1.00 0.00 N
 ATOM 55 C4 DC A 3 7.801 0.850 -4.602 1.00 0.00 C
 ATOM 56 N4 DC A 3 6.918 1.484 -5.408 1.00 0.00 N
 ATOM 57 C5 DC A 3 8.418 1.572 -3.524 1.00 0.00 C
 ATOM 58 C6 DC A 3 9.314 0.888 -2.774 1.00 0.00 C
 END

COMPND AAC / M06-2X / H MODEL / SINGLY IONIZED

ATOM 1 O5' DA A 1 0.000 0.000 0.000 1.00 0.00 O
 ATOM 2 C5' DA A 1 0.000 0.000 1.415 1.00 0.00 C
 ATOM 3 C4' DA A 1 1.436 0.000 1.909 1.00 0.00 C
 ATOM 4 O4' DA A 1 2.114 -1.187 1.491 1.00 0.00 O
 ATOM 5 C3' DA A 1 2.298 1.138 1.363 1.00 0.00 C
 ATOM 6 O3' DA A 1 3.383 1.295 2.311 1.00 0.00 O
 ATOM 7 C2' DA A 1 2.818 0.564 0.056 1.00 0.00 C
 ATOM 8 C1' DA A 1 3.001 -0.909 0.424 1.00 0.00 C
 ATOM 9 N9 DA A 1 2.750 -1.826 -0.678 1.00 0.00 N
 ATOM 10 C8 DA A 1 2.043 -1.653 -1.844 1.00 0.00 C
 ATOM 11 N7 DA A 1 1.958 -2.739 -2.576 1.00 0.00 N
 ATOM 12 C5 DA A 1 2.622 -3.692 -1.839 1.00 0.00 C
 ATOM 13 C6 DA A 1 2.900 -0.060 -2.068 1.00 0.00 C
 ATOM 14 N6 DA A 1 2.529 -5.680 -3.195 1.00 0.00 N
 ATOM 15 N1 DA A 1 3.570 -5.742 -1.125 1.00 0.00 N
 ATOM 16 C2 DA A 1 3.974 -5.089 -0.029 1.00 0.00 C
 ATOM 17 N3 DA A 1 3.808 -3.803 0.290 1.00 0.00 N
 ATOM 18 C4 DA A 1 3.121 -3.155 -0.655 1.00 0.00 C
 ATOM 19 P DA A 2 4.486 2.433 2.133 1.00 0.00 P
 ATOM 20 OP1 DA A 2 3.801 3.813 2.532 1.00 0.00 O
 ATOM 21 OP2 DA A 2 5.109 2.554 0.799 1.00 0.00 O
 ATOM 22 O5' DA A 2 5.471 2.124 3.341 1.00 0.00 O
 ATOM 23 C5' DA A 2 5.625 0.851 3.991 1.00 0.00 C
 ATOM 24 C4' DA A 2 6.649 -0.010 3.288 1.00 0.00 C
 ATOM 25 O4' DA A 2 6.102 -0.481 2.051 1.00 0.00 O
 ATOM 26 C3' DA A 2 7.966 0.715 2.918 1.00 0.00 C
 ATOM 27 O3' DA A 2 9.018 -0.200 3.277 1.00 0.00 O
 ATOM 28 C2' DA A 2 7.874 0.852 1.402 1.00 0.00 C
 ATOM 29 C1' DA A 2 7.151 -0.451 1.095 1.00 0.00 C
 ATOM 30 N9 DA A 2 6.563 -0.578 -0.223 1.00 0.00 N
 ATOM 31 C8 DA A 2 6.019 0.371 -1.023 1.00 0.00 C
 ATOM 32 N7 DA A 2 5.462 -0.118 -2.139 1.00 0.00 N
 ATOM 33 C5 DA A 2 5.670 -1.439 -2.054 1.00 0.00 C
 ATOM 34 C6 DA A 2 5.366 -2.534 -2.927 1.00 0.00 C
 ATOM 35 N6 DA A 2 4.719 -2.358 -4.057 1.00 0.00 N
 ATOM 36 N1 DA A 2 5.755 -3.782 -2.563 1.00 0.00 N
 ATOM 37 C2 DA A 2 6.389 -3.937 -1.422 1.00 0.00 C
 ATOM 38 N3 DA A 2 6.717 -2.985 -0.486 1.00 0.00 N
 ATOM 39 C4 DA A 2 6.354 -1.792 -0.858 1.00 0.00 C
 ATOM 40 P DC A 3 10.567 0.141 3.102 1.00 0.00 P
 ATOM 41 OP1 DC A 3 10.866 1.187 4.269 1.00 0.00 O
 ATOM 42 OP2 DC A 3 11.020 0.614 1.775 1.00 0.00 O
 ATOM 43 O5' DC A 3 11.226 -1.192 3.653 1.00 0.00 O
 ATOM 44 C5' DC A 3 10.606 -2.487 3.565 1.00 0.00 C
 ATOM 45 C4' DC A 3 10.730 -3.090 2.187 1.00 0.00 C
 ATOM 46 O4' DC A 3 9.922 -2.368 1.258 1.00 0.00 O
 ATOM 47 C3' DC A 3 12.165 -3.114 1.600 1.00 0.00 C

ATOM 48 O3' DC A 3 12.504 -4.478 1.435 1.00 0.00 O
 ATOM 49 C2' DC A 3 12.004 -2.379 0.264 1.00 0.00 C
 ATOM 50 C1' DC A 3 10.524 -2.620 -0.011 1.00 0.00 C
 ATOM 51 N1 DC A 3 9.895 -1.800 -1.026 1.00 0.00 N
 ATOM 52 C2 DC A 3 9.152 -2.435 -2.055 1.00 0.00 C
 ATOM 53 O2 DC A 3 9.071 -3.660 -2.075 1.00 0.00 O
 ATOM 54 N3 DC A 3 8.525 -1.646 -2.972 1.00 0.00 N
 ATOM 55 C4 DC A 3 8.674 -0.324 -2.935 1.00 0.00 C
 ATOM 56 N4 DC A 3 8.051 0.392 -3.877 1.00 0.00 N
 ATOM 57 C5 DC A 3 9.444 0.343 -1.925 1.00 0.00 C
 ATOM 58 C6 DC A 3 10.028 -0.443 -0.991 1.00 0.00 C
 END

COMPND CCA / M06-2X / H MODEL / NEUTRAL
 ATOM 1 O5' DC A 1 0.000 0.000 0.000 1.00 0.00 O
 ATOM 2 C5' DC A 1 0.000 0.000 1.414 1.00 0.00 C
 ATOM 3 C4' DC A 1 1.422 0.000 1.951 1.00 0.00 C
 ATOM 4 O4' DC A 1 2.072 -1.165 1.435 1.00 0.00 O
 ATOM 5 C3' DC A 1 2.261 1.197 1.468 1.00 0.00 C
 ATOM 6 O3' DC A 1 3.085 1.588 2.589 1.00 0.00 O
 ATOM 7 C2' DC A 1 3.106 0.606 0.348 1.00 0.00 C
 ATOM 8 C1' DC A 1 3.345 -0.789 0.908 1.00 0.00 C
 ATOM 9 N1 DC A 1 3.779 -1.816 -0.015 1.00 0.00 N
 ATOM 10 C2 DC A 1 4.344 -2.987 0.570 1.00 0.00 C
 ATOM 11 O2 DC A 1 4.461 -3.031 1.789 1.00 0.00 O
 ATOM 12 N3 DC A 1 4.722 -3.997 -0.261 1.00 0.00 N
 ATOM 13 C4 DC A 1 4.540 -3.892 -1.565 1.00 0.00 C
 ATOM 14 N4 DC A 1 4.990 -4.894 -2.341 1.00 0.00 N
 ATOM 15 C5 DC A 1 3.918 -2.754 -2.183 1.00 0.00 C
 ATOM 16 C6 DC A 1 3.561 -1.745 -1.359 1.00 0.00 C
 ATOM 17 P DC A 2 3.978 2.907 2.521 1.00 0.00 P
 ATOM 18 OP1 DC A 2 2.968 4.134 2.390 1.00 0.00 O
 ATOM 19 OP2 DC A 2 4.988 2.975 1.446 1.00 0.00 O
 ATOM 20 O5' DC A 2 4.527 3.005 4.008 1.00 0.00 O
 ATOM 21 C5' DC A 2 4.542 1.934 4.970 1.00 0.00 C
 ATOM 22 C4' DC A 2 5.816 1.129 4.856 1.00 0.00 C
 ATOM 23 O4' DC A 2 5.777 0.332 3.677 1.00 0.00 O
 ATOM 24 C3' DC A 2 7.090 2.002 4.742 1.00 0.00 C
 ATOM 25 O3' DC A 2 8.039 1.441 5.675 1.00 0.00 O
 ATOM 26 C2' DC A 2 7.536 1.771 3.305 1.00 0.00 C
 ATOM 27 C1' DC A 2 7.114 0.314 3.155 1.00 0.00 C
 ATOM 28 N1 DC A 2 7.110 -0.272 1.839 1.00 0.00 N
 ATOM 29 C2 DC A 2 7.395 -1.668 1.744 1.00 0.00 C
 ATOM 30 O2 DC A 2 7.650 -2.293 2.766 1.00 0.00 O
 ATOM 31 N3 DC A 2 7.390 -2.233 0.505 1.00 0.00 N
 ATOM 32 C4 DC A 2 7.073 -1.515 -0.557 1.00 0.00 C
 ATOM 33 N4 DC A 2 7.125 -2.119 -1.758 1.00 0.00 N
 ATOM 34 C5 DC A 2 6.706 -0.127 -0.486 1.00 0.00 C
 ATOM 35 C6 DC A 2 6.743 0.446 0.739 1.00 0.00 C
 ATOM 36 P DA A 3 9.462 2.107 5.943 1.00 0.00 P
 ATOM 37 OP1 DA A 3 9.189 3.440 6.777 1.00 0.00 O
 ATOM 38 OP2 DA A 3 10.300 2.424 4.770 1.00 0.00 O
 ATOM 39 O5' DA A 3 10.072 1.110 7.019 1.00 0.00 O
 ATOM 40 C5' DA A 3 9.701 -0.275 7.144 1.00 0.00 C
 ATOM 41 C4' DA A 3 10.417 -1.137 6.134 1.00 0.00 C
 ATOM 42 O4' DA A 3 9.929 -0.845 4.824 1.00 0.00 O
 ATOM 43 C3' DA A 3 11.956 -0.952 6.089 1.00 0.00 C
 ATOM 44 O3' DA A 3 12.519 -2.230 6.324 1.00 0.00 O
 ATOM 45 C2' DA A 3 12.197 -0.468 4.657 1.00 0.00 C
 ATOM 46 C1' DA A 3 11.015 -11.116 3.945 1.00 0.00 C
 ATOM 47 N9 DA A 3 10.721 -0.629 2.625 1.00 0.00 N
 ATOM 48 C8 DA A 3 10.562 0.678 2.216 1.00 0.00 C
 ATOM 49 N7 DA A 3 10.300 0.798 0.939 1.00 0.00 N
 ATOM 50 C5 DA A 3 10.306 -0.503 0.471 1.00 0.00 C
 ATOM 51 C6 DA A 3 10.124 -1.060 -0.810 1.00 0.00 C
 ATOM 52 N6 DA A 3 9.873 -0.291 -1.898 1.00 0.00 N
 ATOM 53 N1 DA A 3 10.240 -2.386 -0.959 1.00 0.00 N
 ATOM 54 C2 DA A 3 10.499 -3.128 0.128 1.00 0.00 C
 ATOM 55 N3 DA A 3 10.679 -2.733 1.384 1.00 0.00 N
 ATOM 56 C4 DA A 3 10.568 -1.405 1.498 1.00 0.00 C
 END

COMPND CCA / M06-2X / H MODEL / SINGLY IONIZED
 ATOM 1 O5' DC A 1 0.000 0.000 0.000 1.00 0.00 O
 ATOM 2 C5' DC A 1 0.000 0.000 1.414 1.00 0.00 C
 ATOM 3 C4' DC A 1 1.421 0.000 1.953 1.00 0.00 C
 ATOM 4 O4' DC A 1 2.071 -1.170 1.446 1.00 0.00 O
 ATOM 5 C3' DC A 1 2.264 1.194 1.465 1.00 0.00 C
 ATOM 6 O3' DC A 1 3.082 1.590 2.590 1.00 0.00 O
 ATOM 7 C2' DC A 1 3.114 0.593 0.356 1.00 0.00 C
 ATOM 8 C1' DC A 1 3.353 -0.794 0.941 1.00 0.00 C
 ATOM 9 N1 DC A 1 3.825 -1.829 0.047 1.00 0.00 N
 ATOM 10 C2 DC A 1 4.528 -2.917 0.642 1.00 0.00 C
 ATOM 11 O2 DC A 1 4.712 -2.905 1.854 1.00 0.00 O
 ATOM 12 N3 DC A 1 4.965 -3.916 -0.172 1.00 0.00 N
 ATOM 13 C4 DC A 1 4.745 -3.865 -1.473 1.00 0.00 C
 ATOM 14 N4 DC A 1 5.259 -4.849 -2.233 1.00 0.00 N
 ATOM 15 C5 DC A 1 4.019 -2.797 -2.104 1.00 0.00 C
 ATOM 16 C6 DC A 1 3.577 -1.812 -1.293 1.00 0.00 C
 ATOM 17 P DC A 2 4.037 2.864 2.525 1.00 0.00 P
 ATOM 18 OP1 DC A 2 3.095 4.137 2.348 1.00 0.00 O
 ATOM 19 OP2 DC A 2 5.082 2.871 1.481 1.00 0.00 O
 ATOM 20 O5' DC A 2 4.552 2.955 4.026 1.00 0.00 O
 ATOM 21 C5' DC A 2 4.493 1.887 4.987 1.00 0.00 C
 ATOM 22 C4' DC A 2 5.703 0.991 4.878 1.00 0.00 C
 ATOM 23 O4' DC A 2 5.650 0.245 3.663 1.00 0.00 O
 ATOM 24 C3' DC A 2 7.057 1.746 4.858 1.00 0.00 C
 ATOM 25 O3' DC A 2 7.880 1.087 5.846 1.00 0.00 O
 ATOM 26 C2' DC A 2 7.576 1.479 3.450 1.00 0.00 C
 ATOM 27 C1' DC A 2 7.008 0.078 3.252 1.00 0.00 C
 ATOM 28 N1 DC A 2 7.060 -0.486 1.925 1.00 0.00 N
 ATOM 29 C2 DC A 2 7.583 -1.795 1.770 1.00 0.00 C
 ATOM 30 O2 DC A 2 7.991 -2.402 2.758 1.00 0.00 O
 ATOM 31 N3 DC A 2 7.672 -2.301 0.511 1.00 0.00 N
 ATOM 32 C4 DC A 2 7.233 -1.605 -0.526 1.00 0.00 C
 ATOM 33 N4 DC A 2 7.362 -2.161 -1.744 1.00 0.00 N
 ATOM 34 C5 DC A 2 6.679 -0.287 -0.401 1.00 0.00 C
 ATOM 35 C6 DC A 2 6.619 0.228 0.849 1.00 0.00 C
 ATOM 36 P DA A 3 9.349 1.584 6.216 1.00 0.00 P
 ATOM 37 OP1 DA A 3 9.199 3.018 6.891 1.00 0.00 O
 ATOM 38 OP2 DA A 3 10.325 1.655 5.108 1.00 0.00 O
 ATOM 39 O5' DA A 3 9.711 0.621 7.425 1.00 0.00 O
 ATOM 40 C5' DA A 3 9.128 -0.673 7.657 1.00 0.00 C
 ATOM 41 C4' DA A 3 9.873 -1.753 6.911 1.00 0.00 C
 ATOM 42 O4' DA A 3 9.620 -1.624 5.506 1.00 0.00 O
 ATOM 43 C3' DA A 3 11.412 -1.731 7.089 1.00 0.00 C
 ATOM 44 O3' DA A 3 11.782 -3.025 7.526 1.00 0.00 O
 ATOM 45 C2' DA A 3 11.913 -1.442 5.671 1.00 0.00 C

ATOM 46 C1' DA A 3 10.793 -2.083 4.864 1.00 0.00 C
 ATOM 47 N9 DA A 3 10.733 -1.712 3.463 1.00 0.00 N
 ATOM 48 C8 DA A 3 10.555 -0.476 2.958 1.00 0.00 C
 ATOM 49 N7 DA A 3 10.442 -0.446 1.614 1.00 0.00 N
 ATOM 50 C5 DA A 3 10.554 -1.724 1.256 1.00 0.00 C
 ATOM 51 C6 DA A 3 10.522 -2.387 -0.016 1.00 0.00 C
 ATOM 52 N6 DA A 3 10.409 -1.722 -1.141 1.00 0.00 N
 ATOM 53 N1 DA A 3 10.639 -3.741 -0.048 1.00 0.00 N
 ATOM 54 C2 DA A 3 10.786 -4.381 1.088 1.00 0.00 C
 ATOM 55 N3 DA A 3 10.853 -3.873 2.364 1.00 0.00 N
 ATOM 56 C4 DA A 3 10.720 -2.582 2.387 1.00 0.00 C
 END

COMPND CAA / M06-2X / H MODEL / NEUTRAL
 ATOM 1 O5' DC A 1 0.000 0.000 0.000 1.00 0.00 O
 ATOM 2 C5' DC A 1 0.000 0.000 1.417 1.00 0.00 C
 ATOM 3 C4' DC A 1 1.439 0.000 1.896 1.00 0.00 C
 ATOM 4 O4' DC A 1 2.059 -1.243 1.567 1.00 0.00 O
 ATOM 5 C3' DC A 1 2.295 1.083 1.229 1.00 0.00 C
 ATOM 6 O3' DC A 1 3.297 1.475 2.200 1.00 0.00 O
 ATOM 7 C2' DC A 1 2.945 0.342 0.075 1.00 0.00 C
 ATOM 8 C1' DC A 1 3.158 -1.043 0.684 1.00 0.00 C
 ATOM 9 N1 DC A 1 3.192 -2.123 -0.285 1.00 0.00 N
 ATOM 10 C2 DC A 1 4.216 -3.101 -0.200 1.00 0.00 C
 ATOM 11 O2 DC A 1 5.002 -3.068 0.742 1.00 0.00 O
 ATOM 12 N3 DC A 1 4.281 -4.032 -1.192 1.00 0.00 N
 ATOM 13 C4 DC A 1 3.341 -4.085 -2.121 1.00 0.00 C
 ATOM 14 N4 DC A 1 3.482 -4.995 -3.099 1.00 0.00 N
 ATOM 15 C5 DC A 1 2.190 -3.224 -2.122 1.00 0.00 C
 ATOM 16 C6 DC A 1 2.168 -2.257 -1.179 1.00 0.00 C
 ATOM 17 P DA A 2 4.203 2.751 1.914 1.00 0.00 P
 ATOM 18 OP2 DA A 2 4.920 2.803 0.620 1.00 0.00 O
 ATOM 19 OP1 DA A 2 3.203 3.980 2.111 1.00 0.00 O
 ATOM 20 O5' DA A 2 5.110 2.814 3.214 1.00 0.00 O
 ATOM 21 C5' DA A 2 5.401 1.711 4.087 1.00 0.00 C
 ATOM 22 C4' DA A 2 6.606 0.937 3.606 1.00 0.00 C
 ATOM 23 O4' DA A 2 6.250 0.186 2.445 1.00 0.00 O
 ATOM 24 C3' DA A 2 7.813 1.818 3.195 1.00 0.00 C
 ATOM 25 O3' DA A 2 8.963 1.219 3.838 1.00 0.00 O
 ATOM 26 C2' DA A 2 7.871 1.633 1.684 1.00 0.00 C
 ATOM 27 C1' DA A 2 7.390 0.193 1.587 1.00 0.00 C
 ATOM 28 N9 DA A 2 7.025 -0.286 0.290 1.00 0.00 N
 ATOM 29 C8 DA A 2 6.337 0.357 -0.713 1.00 0.00 C
 ATOM 30 N7 DA A 2 6.174 -0.377 -1.786 1.00 0.00 N
 ATOM 31 C5 DA A 2 6.805 -1.570 -1.477 1.00 0.00 C
 ATOM 32 C6 DA A 2 7.019 -2.757 -2.200 1.00 0.00 C
 ATOM 33 N6 DA A 2 6.572 -2.915 -3.472 1.00 0.00 N
 ATOM 34 N1 DA A 2 7.748 -3.730 -1.637 1.00 0.00 N
 ATOM 35 C2 DA A 2 8.231 -3.533 -0.402 1.00 0.00 C
 ATOM 36 N3 DA A 2 8.082 -2.476 0.391 1.00 0.00 N
 ATOM 37 C4 DA A 2 7.351 -1.529 -0.200 1.00 0.00 C
 ATOM 38 P DA A 3 10.430 1.819 3.668 1.00 0.00 P
 ATOM 39 OP2 DA A 3 10.610 2.860 2.642 1.00 0.00 O
 ATOM 40 OP1 DA A 3 10.748 2.212 5.181 1.00 0.00 O
 ATOM 41 O5' DA A 3 11.321 0.535 3.344 1.00 0.00 O
 ATOM 42 C5' DA A 3 11.244 -0.648 4.160 1.00 0.00 C
 ATOM 43 C4' DA A 3 11.968 -1.764 3.436 1.00 0.00 C
 ATOM 44 O4' DA A 3 11.265 -2.078 2.233 1.00 0.00 O
 ATOM 45 C3' DA A 3 13.408 -1.439 3.019 1.00 0.00 C
 ATOM 46 O3' DA A 3 14.117 -2.663 3.086 1.00 0.00 O
 ATOM 47 C2' DA A 3 13.238 -0.974 1.577 1.00 0.00 C
 ATOM 48 C1' DA A 3 12.089 -1.861 1.098 1.00 0.00 C
 ATOM 49 N9 DA A 3 11.288 -1.269 0.048 1.00 0.00 N
 ATOM 50 C8 DA A 3 10.600 -0.080 0.108 1.00 0.00 C
 ATOM 51 N7 DA A 3 9.889 0.175 -0.960 1.00 0.00 N
 ATOM 52 C5 DA A 3 10.118 -0.916 -1.778 1.00 0.00 C
 ATOM 53 C6 DA A 3 9.635 -1.279 -3.051 1.00 0.00 C
 ATOM 54 N6 DA A 3 8.779 -0.496 -3.757 1.00 0.00 N
 ATOM 55 N1 DA A 3 10.059 -2.429 -3.596 1.00 0.00 N
 ATOM 56 C2 DA A 3 10.892 -3.204 -2.887 1.00 0.00 C
 ATOM 57 N3 DA A 3 11.393 -2.995 -1.673 1.00 0.00 N
 ATOM 58 C4 DA A 3 10.975 -1.829 -1.167 1.00 0.00 C
 END

COMPND CAA / M06-2X / H MODEL / SINGLY IONIZED
 ATOM 1 O5' DC A 1 0.000 0.000 0.000 1.00 0.00 O
 ATOM 2 C5' DC A 1 0.000 0.000 1.417 1.00 0.00 C
 ATOM 3 C4' DC A 1 1.439 0.000 1.894 1.00 0.00 C
 ATOM 4 O4' DC A 1 2.063 -1.244 1.569 1.00 0.00 O
 ATOM 5 C3' DC A 1 2.302 1.081 1.235 1.00 0.00 C
 ATOM 6 O3' DC A 1 3.291 1.462 2.223 1.00 0.00 O
 ATOM 7 C2' DC A 1 2.966 0.340 0.088 1.00 0.00 C
 ATOM 8 C1' DA A 1 3.166 -1.046 0.699 1.00 0.00 C
 ATOM 9 N1 DA A 1 3.175 -2.127 -0.280 1.00 0.00 N
 ATOM 10 C2 DC A 1 4.323 -2.929 -0.427 1.00 0.00 C
 ATOM 11 O2 DC A 1 5.313 -2.691 0.274 1.00 0.00 O
 ATOM 12 N3 DC A 1 4.297 -3.927 -1.345 1.00 0.00 N
 ATOM 13 C4 DC A 1 3.202 -4.142 -2.066 1.00 0.00 C
 ATOM 14 N4 DC A 1 3.241 -5.131 -2.969 1.00 0.00 N
 ATOM 15 C5 DA A 1 2.002 -3.371 -1.909 1.00 0.00 C
 ATOM 16 C6 DC A 1 2.042 -2.374 -0.997 1.00 0.00 C
 ATOM 17 P DA A 2 4.191 2.751 1.981 1.00 0.00 P
 ATOM 18 OP1 DA A 2 3.167 3.969 2.083 1.00 0.00 O
 ATOM 19 OP2 DA A 2 4.998 2.783 0.741 1.00 0.00 O
 ATOM 20 O5' DA A 2 4.996 2.848 3.344 1.00 0.00 O
 ATOM 21 C5' DA A 2 5.277 1.754 4.234 1.00 0.00 C
 ATOM 22 C4' DA A 2 6.615 1.127 3.913 1.00 0.00 C
 ATOM 23 O4' DA A 2 6.506 0.355 2.712 1.00 0.00 O
 ATOM 24 C3' DA A 2 7.741 2.150 3.653 1.00 0.00 C
 ATOM 25 O3' DA A 2 8.919 1.616 4.292 1.00 0.00 O
 ATOM 26 C2' DA A 2 7.929 2.090 2.142 1.00 0.00 C
 ATOM 27 C1' DA A 2 7.676 0.601 1.946 1.00 0.00 C
 ATOM 28 N9 DA A 2 7.437 0.118 0.603 1.00 0.00 N
 ATOM 29 C8 DA A 2 6.775 0.697 -0.430 1.00 0.00 C
 ATOM 30 N7 DA A 2 6.590 -0.119 -1.479 1.00 0.00 N
 ATOM 31 C5 DA A 2 7.155 -1.268 -1.100 1.00 0.00 C
 ATOM 32 C6 DA A 2 7.260 -2.558 -1.719 1.00 0.00 C
 ATOM 33 N6 DA A 2 6.757 -2.792 -2.909 1.00 0.00 N
 ATOM 34 N1 DA A 2 7.884 -3.555 -1.046 1.00 0.00 N
 ATOM 35 C2 DA A 2 8.330 -3.306 0.166 1.00 0.00 C
 ATOM 36 N3 DA A 2 8.273 -2.141 0.883 1.00 0.00 N
 ATOM 37 C4 DA A 2 7.698 -1.185 0.214 1.00 0.00 C
 ATOM 38 P DA A 3 10.136 2.566 4.687 1.00 0.00 P
 ATOM 39 OP1 DA A 3 9.631 3.139 6.088 1.00 0.00 O
 ATOM 40 OP2 DA A 3 10.559 3.570 3.694 1.00 0.00 O
 ATOM 41 O5' DA A 3 11.283 1.506 4.986 1.00 0.00 O
 ATOM 42 C5' DA A 3 11.033 0.320 5.767 1.00 0.00 C
 ATOM 43 C4' DA A 3 11.729 -0.848 5.097 1.00 0.00 C

ATOM 44 O4' DA A 3 11.206 -1.007 3.777 1.00 0.00 O
 ATOM 45 C3' DA A 3 13.249 -0.709 4.944 1.00 0.00 C
 ATOM 46 O3' DA A 3 13.774 -2.020 5.053 1.00 0.00 O
 ATOM 47 C2' DA A 3 13.395 -0.169 3.525 1.00 0.00 C
 ATOM 48 C1' DA A 3 12.240 -0.867 2.812 1.00 0.00 C
 ATOM 49 N9 DA A 3 11.716 -0.132 1.679 1.00 0.00 N
 ATOM 50 C8 DA A 3 11.387 1.202 1.634 1.00 0.00 C
 ATOM 51 N7 DA A 3 10.766 1.554 0.535 1.00 0.00 N
 ATOM 52 C5 DA A 3 10.670 0.382 -0.181 1.00 0.00 C
 ATOM 53 C6 DA A 3 10.074 0.074 -1.424 1.00 0.00 C
 ATOM 54 N6 DA A 3 9.428 0.999 -2.146 1.00 0.00 N
 ATOM 55 N1 DA A 3 10.125 -1.198 -1.861 1.00 0.00 N
 ATOM 56 C2 DA A 3 10.763 -2.107 -1.105 1.00 0.00 C
 ATOM 57 N3 DA A 3 11.336 -1.948 0.085 1.00 0.00 N
 ATOM 58 C4 DA A 3 11.249 -0.680 0.506 1.00 0.00 C
 END

COMPND ACC / M06-2X / NA MODEL / NEUTRAL

ATOM 1 O5' DA A 1 7.107 0.774 -0.532 1.00 0.00 O
 ATOM 2 C5' DA A 1 6.770 1.663 -1.581 1.00 0.00 C
 ATOM 3 C4' DA A 1 5.261 1.822 -1.641 1.00 0.00 C
 ATOM 4 O4' DA A 1 4.650 0.568 -1.961 1.00 0.00 O
 ATOM 5 C3' DA A 1 4.607 2.257 -0.326 1.00 0.00 C
 ATOM 6 O3' DA A 1 3.372 2.891 -0.685 1.00 0.00 O
 ATOM 7 C2' DA A 1 4.335 0.923 0.349 1.00 0.00 C
 ATOM 8 C1' DA A 1 3.900 0.090 -0.852 1.00 0.00 C
 ATOM 9 N9 DA A 1 4.108 -1.333 -0.684 1.00 0.00 N
 ATOM 10 C8 DA A 1 5.063 -1.990 0.057 1.00 0.00 C
 ATOM 11 N7 DA A 1 4.888 -3.286 0.115 1.00 0.00 N
 ATOM 12 C5 DA A 1 3.755 -3.501 -0.646 1.00 0.00 C
 ATOM 13 C6 DA A 1 2.995 -4.652 -0.929 1.00 0.00 C
 ATOM 14 N6 DA A 1 3.273 -5.848 -0.356 1.00 0.00 N
 ATOM 15 N1 DA A 1 1.930 -4.541 -1.733 1.00 0.00 N
 ATOM 16 C2 DA A 1 1.590 -3.322 -2.175 1.00 0.00 C
 ATOM 17 N3 DA A 1 2.179 -2.155 -1.932 1.00 0.00 N
 ATOM 18 C4 DA A 1 3.260 -2.306 -1.159 1.00 0.00 C
 ATOM 19 P DC A 2 2.464 3.610 0.456 1.00 0.00 P
 ATOM 20 OP1 DC A 2 3.094 4.898 0.918 1.00 0.00 O
 ATOM 21 OP2 DC A 2 2.127 2.659 1.585 1.00 0.00 O
 ATOM 22 O5' DC A 2 1.159 3.958 -0.431 1.00 0.00 O
 ATOM 23 C5' DC A 2 0.772 3.348 -1.662 1.00 0.00 C
 ATOM 24 C4' DC A 2 -0.236 2.234 -1.468 1.00 0.00 C
 ATOM 25 O4' DC A 2 0.424 1.036 -1.052 1.00 0.00 O
 ATOM 26 C3' DC A 2 -1.306 2.532 -0.390 1.00 0.00 C
 ATOM 27 O3' DC A 2 -2.576 2.160 -0.938 1.00 0.00 O
 ATOM 28 C2' DC A 2 -0.922 1.577 0.732 1.00 0.00 C
 ATOM 29 C1' DC A 2 -0.442 0.401 -0.110 1.00 0.00 C
 ATOM 30 N1 DC A 2 0.272 -0.668 0.553 1.00 0.00 N
 ATOM 31 C2 DC A 2 -0.053 -2.015 0.224 1.00 0.00 C
 ATOM 32 O2 DC A 2 -0.956 -2.241 -0.574 1.00 0.00 O
 ATOM 33 N3 DC A 2 0.666 -3.004 0.828 1.00 0.00 N
 ATOM 34 C4 DC A 2 1.655 -2.710 1.650 1.00 0.00 C
 ATOM 35 N4 DC A 2 2.315 -3.742 2.228 1.00 0.00 N
 ATOM 36 C5 DC A 2 2.024 -1.363 1.981 1.00 0.00 C
 ATOM 37 C6 DC A 2 1.298 -0.377 1.398 1.00 0.00 C
 ATOM 38 P DC A 3 -3.939 2.689 -0.210 1.00 0.00 P
 ATOM 39 OP1 DC A 3 -4.372 3.999 -0.814 1.00 0.00 O
 ATOM 40 OP2 DC A 3 -3.805 2.663 1.290 1.00 0.00 O
 ATOM 41 O5' DC A 3 -4.933 1.476 -0.627 1.00 0.00 O
 ATOM 42 C5' DC A 3 -5.043 1.084 -1.995 1.00 0.00 C
 ATOM 43 C4' DC A 3 -5.432 -0.379 -2.032 1.00 0.00 C
 ATOM 44 O4' DC A 3 -4.391 -1.144 -1.420 1.00 0.00 O
 ATOM 45 C3' DC A 3 -6.726 -0.717 -1.276 1.00 0.00 C
 ATOM 46 O3' DC A 3 -7.368 -1.738 -2.021 1.00 0.00 O
 ATOM 47 C2' DC A 3 -6.207 -1.241 0.058 1.00 0.00 C
 ATOM 48 C1' DC A 3 -4.925 -1.947 -0.376 1.00 0.00 C
 ATOM 49 N1 DC A 3 -3.926 -2.089 0.672 1.00 0.00 N
 ATOM 50 C2 DC A 3 -3.428 -3.380 0.987 1.00 0.00 C
 ATOM 51 O2 DC A 3 -3.868 -4.357 0.387 1.00 0.00 O
 ATOM 52 N3 DC A 3 -2.487 -3.479 1.968 1.00 0.00 N
 ATOM 53 C4 DC A 3 -2.038 -2.396 2.572 1.00 0.00 C
 ATOM 54 N4 DC A 3 -1.091 -2.550 3.524 1.00 0.00 N
 ATOM 55 C5 DC A 3 -2.523 -1.077 2.281 1.00 0.00 C
 ATOM 56 C6 DC A 3 -3.449 -0.979 1.299 1.00 0.00 C
 END

COMPND ACC / M06-2X / NA MODEL / SINGLY IONIZED

ATOM 1 O5' DA A 1 6.783 0.071 -0.671 1.00 0.00 O
 ATOM 2 C5' DA A 1 6.629 1.032 -1.702 1.00 0.00 C
 ATOM 3 C4' DA A 1 5.176 1.466 -1.774 1.00 0.00 C
 ATOM 4 O4' DA A 1 4.339 0.336 -2.059 1.00 0.00 O
 ATOM 5 C3' DA A 1 4.595 2.049 -0.483 1.00 0.00 C
 ATOM 6 O3' DA A 1 3.452 2.815 -0.881 1.00 0.00 O
 ATOM 7 C2' DA A 1 4.148 0.799 0.257 1.00 0.00 C
 ATOM 8 C1' DA A 1 3.605 -0.022 -0.909 1.00 0.00 C
 ATOM 9 N9 DA A 1 3.699 -1.470 -0.694 1.00 0.00 N
 ATOM 10 C8 DA A 1 4.606 -2.192 -0.003 1.00 0.00 C
 ATOM 11 N7 DA A 1 4.349 -3.512 0.028 1.00 0.00 N
 ATOM 12 C5 DA A 1 3.229 -3.636 -0.685 1.00 0.00 C
 ATOM 13 C6 DA A 1 2.392 -4.755 -1.016 1.00 0.00 C
 ATOM 14 N6 DA A 1 2.660 -5.968 -0.588 1.00 0.00 N
 ATOM 15 N1 DA A 1 1.303 -4.544 -1.794 1.00 0.00 N
 ATOM 16 C2 DA A 1 1.049 -3.321 -2.203 1.00 0.00 C
 ATOM 17 N3 DA A 1 1.757 -2.169 -1.962 1.00 0.00 N
 ATOM 18 C4 DA A 1 2.785 -2.380 -1.196 1.00 0.00 C
 ATOM 19 P DC A 2 2.578 3.662 0.202 1.00 0.00 P
 ATOM 20 OP1 DC A 2 3.226 4.987 0.507 1.00 0.00 O
 ATOM 21 OP2 DC A 2 2.262 2.840 1.432 1.00 0.00 O
 ATOM 22 O5' DC A 2 1.254 3.934 -0.685 1.00 0.00 O
 ATOM 23 C5' DC A 2 0.820 3.203 -1.831 1.00 0.00 C
 ATOM 24 C4' DC A 2 -0.205 2.139 -1.493 1.00 0.00 C
 ATOM 25 O4' DC A 2 0.450 0.987 -0.956 1.00 0.00 O
 ATOM 26 C3' DC A 2 -1.251 2.563 -0.437 1.00 0.00 C
 ATOM 27 O3' DC A 2 -2.519 2.082 -0.893 1.00 0.00 O
 ATOM 28 C2' DC A 2 -0.814 1.782 0.799 1.00 0.00 C
 ATOM 29 C1' DC A 2 -0.370 0.500 0.107 1.00 0.00 C
 ATOM 30 N1 DC A 2 0.381 -0.480 0.866 1.00 0.00 N
 ATOM 31 C2 DC A 2 0.217 -1.850 0.524 1.00 0.00 C
 ATOM 32 O2 DC A 2 -0.597 -2.158 -0.344 1.00 0.00 O
 ATOM 33 N3 DC A 2 1.002 -2.768 1.152 1.00 0.00 N
 ATOM 34 C4 DC A 2 1.851 -2.385 2.103 1.00 0.00 C
 ATOM 35 N4 DC A 2 2.572 -3.339 2.704 1.00 0.00 N
 ATOM 36 C5 DC A 2 2.029 -1.014 2.478 1.00 0.00 C
 ATOM 37 C6 DC A 2 1.278 -0.100 1.817 1.00 0.00 C
 ATOM 38 P DC A 3 -3.887 2.637 -0.200 1.00 0.00 P
 ATOM 39 OP1 DC A 3 -4.233 3.996 -0.752 1.00 0.00 O
 ATOM 40 OP2 DC A 3 -3.830 2.536 1.303 1.00 0.00 O
 ATOM 41 O5' DC A 3 -4.901 1.482 -0.717 1.00 0.00 O
 ATOM 42 C5' DC A 3 -4.900 1.057 -2.081 1.00 0.00 C

ATOM 43 C4' DC A 3 -5.154 -0.436 -2.112 1.00 0.00 C
 ATOM 44 O4' DC A 3 -4.100 -1.101 -1.412 1.00 0.00 O
 ATOM 45 C3' DC A 3 -6.468 -0.877 -1.447 1.00 0.00 C
 ATOM 46 O3' DC A 3 -6.961 -1.960 -2.218 1.00 0.00 O
 ATOM 47 C2' DC A 3 -6.006 -1.336 -0.069 1.00 0.00 C
 ATOM 48 C1' DC A 3 -4.642 -1.937 -0.396 1.00 0.00 C
 ATOM 49 N1 DC A 3 -3.714 -1.984 0.723 1.00 0.00 N
 ATOM 50 C2 DC A 3 -3.133 -3.224 1.097 1.00 0.00 C
 ATOM 51 O2 DC A 3 -3.447 -4.242 0.488 1.00 0.00 O
 ATOM 52 N3 DC A 3 -2.260 -3.228 2.143 1.00 0.00 N
 ATOM 53 C4 DC A 3 -1.945 -2.099 2.753 1.00 0.00 C
 ATOM 54 N4 DC A 3 -1.038 -2.154 3.745 1.00 0.00 N
 ATOM 55 C5 DC A 3 -2.517 -0.830 2.402 1.00 0.00 C
 ATOM 56 C6 DC A 3 -3.384 -0.829 1.364 1.00 0.00 C
 END

COMPND AAC / M06-2X / NA MODEL / NEUTRAL

ATOM 1 O5' DA A 1 6.648 2.170 -1.018 1.00 0.00 O
 ATOM 2 C5' DA A 1 5.907 2.540 -2.164 1.00 0.00 C
 ATOM 3 C4' DA A 1 4.447 2.223 -1.912 1.00 0.00 C
 ATOM 4 O4' DA A 1 4.271 0.803 -1.857 1.00 0.00 O
 ATOM 5 C3' DA A 1 3.929 2.765 -0.572 1.00 0.00 C
 ATOM 6 O3' DA A 1 2.561 3.139 -0.762 1.00 0.00 O
 ATOM 7 C2' DA A 1 4.019 1.547 0.339 1.00 0.00 C
 ATOM 8 C1' DA A 1 3.634 0.449 -0.642 1.00 0.00 C
 ATOM 9 N9 DA A 1 4.078 -0.874 -0.252 1.00 0.00 N
 ATOM 10 C8 DA A 1 4.741 -1.260 0.889 1.00 0.00 C
 ATOM 11 N7 DA A 1 4.926 -2.551 0.984 1.00 0.00 N
 ATOM 12 C5 DA A 1 4.349 -3.053 -0.166 1.00 0.00 C
 ATOM 13 C6 DA A 1 4.142 -4.364 -0.634 1.00 0.00 C
 ATOM 14 N6 DA A 1 4.507 -5.447 0.096 1.00 0.00 N
 ATOM 15 N1 DA A 1 3.509 -4.540 -1.800 1.00 0.00 N
 ATOM 16 C2 DA A 1 3.040 -3.457 -2.438 1.00 0.00 C
 ATOM 17 N3 DA A 1 3.125 -2.177 -2.085 1.00 0.00 N
 ATOM 18 C4 DA A 1 3.802 -2.033 -0.939 1.00 0.00 C
 ATOM 19 P DA A 2 1.852 4.040 0.390 1.00 0.00 P
 ATOM 20 OP1 DA A 2 2.619 5.321 0.597 1.00 0.00 O
 ATOM 21 OP2 DA A 2 1.618 3.255 1.662 1.00 0.00 O
 ATOM 22 O5' DA A 2 0.431 4.335 -0.324 1.00 0.00 O
 ATOM 23 C5' DA A 2 0.016 3.886 -1.613 1.00 0.00 C
 ATOM 24 C4' DA A 2 -0.848 2.646 -1.504 1.00 0.00 C
 ATOM 25 O4' DA A 2 -0.032 1.517 -1.176 1.00 0.00 O
 ATOM 26 C3' DA A 2 -1.922 2.732 -0.399 1.00 0.00 C
 ATOM 27 O3' DA A 2 -3.096 2.099 -0.919 1.00 0.00 O
 ATOM 28 C2' DA A 2 -1.326 1.875 0.709 1.00 0.00 C
 ATOM 29 C1' DA A 2 -0.689 0.790 -0.145 1.00 0.00 C
 ATOM 30 N9 DA A 2 0.247 -0.071 0.526 1.00 0.00 N
 ATOM 31 C8 DA A 2 1.051 0.206 1.612 1.00 0.00 C
 ATOM 32 N7 DA A 2 1.706 -0.839 2.055 1.00 0.00 N
 ATOM 33 C5 DA A 2 1.309 -1.867 1.219 1.00 0.00 C
 ATOM 34 C6 DA A 2 1.576 -3.250 1.199 1.00 0.00 C
 ATOM 35 N6 DA A 2 2.328 -3.858 2.158 1.00 0.00 N
 ATOM 36 N1 DA A 2 1.025 -4.000 0.238 1.00 0.00 N
 ATOM 37 C2 DA A 2 0.196 -3.413 -0.637 1.00 0.00 C
 ATOM 38 N3 DA A 2 -0.182 -2.139 -0.695 1.00 0.00 N
 ATOM 39 C4 DA A 2 0.410 -1.412 0.259 1.00 0.00 C
 ATOM 40 P DC A 3 -4.520 2.219 -0.131 1.00 0.00 P
 ATOM 41 OP1 DC A 3 -5.258 3.453 -0.581 1.00 0.00 O
 ATOM 42 OP2 DC A 3 -4.349 2.066 1.358 1.00 0.00 O
 ATOM 43 O5' DC A 3 -5.215 0.857 -0.673 1.00 0.00 O
 ATOM 44 C5' DC A 3 -5.155 0.490 -2.051 1.00 0.00 C
 ATOM 45 C4' DC A 3 -5.119 -1.022 -2.128 1.00 0.00 C
 ATOM 46 O4' DC A 3 -3.938 -1.489 -1.470 1.00 0.00 O
 ATOM 47 C3' DC A 3 -6.307 -1.723 -1.447 1.00 0.00 C
 ATOM 48 O3' DC A 3 -6.615 -2.854 -2.247 1.00 0.00 O
 ATOM 49 C2' DC A 3 -5.724 -2.134 -0.101 1.00 0.00 C
 ATOM 50 C1' DC A 3 -4.285 -2.458 -0.490 1.00 0.00 C
 ATOM 51 N1 DC A 3 -3.330 -2.389 0.606 1.00 0.00 N
 ATOM 52 C2 DC A 3 -2.548 -3.532 0.912 1.00 0.00 C
 ATOM 53 O2 DC A 3 -2.686 -4.555 0.247 1.00 0.00 O
 ATOM 54 N3 DC A 3 -1.673 -3.443 1.954 1.00 0.00 N
 ATOM 55 C4 DC A 3 -1.545 -2.315 2.627 1.00 0.00 C
 ATOM 56 N4 DC A 3 -0.670 -2.298 3.659 1.00 0.00 N
 ATOM 57 C5 DC A 3 -2.306 -1.134 2.328 1.00 0.00 C
 ATOM 58 C6 DC A 3 -3.179 -1.225 1.296 1.00 0.00 C
 END

COMPND AAC / M06-2X / NA MODEL / SINGLY IONIZED

ATOM 1 O5' DA A 1 6.511 2.148 -0.495 1.00 0.00 O
 ATOM 2 C5' DA A 1 5.955 2.736 -1.656 1.00 0.00 C
 ATOM 3 C4' DA A 1 4.456 2.512 -1.643 1.00 0.00 C
 ATOM 4 O4' DA A 1 4.173 1.114 -1.785 1.00 0.00 O
 ATOM 5 C3' DA A 1 3.772 2.938 -0.338 1.00 0.00 C
 ATOM 6 O3' DA A 1 2.439 3.329 -0.681 1.00 0.00 O
 ATOM 7 C2' DA A 1 3.752 1.641 0.460 1.00 0.00 C
 ATOM 8 C1' DA A 1 3.456 0.647 -0.655 1.00 0.00 C
 ATOM 9 N9 DA A 1 3.872 -0.714 -0.370 1.00 0.00 N
 ATOM 10 C8 DA A 1 4.723 -1.186 0.601 1.00 0.00 C
 ATOM 11 N7 DA A 1 4.953 -2.476 0.521 1.00 0.00 N
 ATOM 12 C5 DA A 1 4.227 -2.875 -0.577 1.00 0.00 C
 ATOM 13 C6 DA A 1 4.035 -4.135 -1.190 1.00 0.00 C
 ATOM 14 N6 DA A 1 4.587 -5.252 -0.695 1.00 0.00 N
 ATOM 15 N1 DA A 1 3.256 -4.203 -2.280 1.00 0.00 N
 ATOM 16 C2 DA A 1 2.663 -3.082 -2.713 1.00 0.00 C
 ATOM 17 N3 DA A 1 2.738 -1.847 -2.214 1.00 0.00 N
 ATOM 18 C4 DA A 1 3.542 -1.804 -1.146 1.00 0.00 C
 ATOM 19 P DA A 2 1.522 4.064 0.441 1.00 0.00 P
 ATOM 20 OP1 DA A 2 2.119 5.389 0.838 1.00 0.00 O
 ATOM 21 OP2 DA A 2 1.238 3.150 1.615 1.00 0.00 O
 ATOM 22 O5' DA A 2 0.176 4.296 -0.423 1.00 0.00 O
 ATOM 23 C5' DA A 2 -0.095 3.769 -1.721 1.00 0.00 C
 ATOM 24 C4' DA A 2 -0.902 2.489 -1.648 1.00 0.00 C
 ATOM 25 O4' DA A 2 -0.069 1.408 -1.199 1.00 0.00 O
 ATOM 26 C3' DA A 2 -2.085 2.548 -0.658 1.00 0.00 C
 ATOM 27 O3' DA A 2 -3.162 1.845 -1.277 1.00 0.00 O
 ATOM 28 C2' DA A 2 -1.560 1.758 0.536 1.00 0.00 C
 ATOM 29 C1' DA A 2 -0.806 0.682 -0.230 1.00 0.00 C
 ATOM 30 N9 DA A 2 0.120 -0.125 0.544 1.00 0.00 N
 ATOM 31 C8 DA A 2 0.845 0.217 1.636 1.00 0.00 C
 ATOM 32 N7 DA A 2 1.608 -0.781 2.114 1.00 0.00 N
 ATOM 33 C5 DA A 2 1.351 -1.806 1.293 1.00 0.00 C
 ATOM 34 C6 DA A 2 1.774 -3.176 1.279 1.00 0.00 C
 ATOM 35 N6 DA A 2 2.592 -3.656 2.188 1.00 0.00 N
 ATOM 36 N1 DA A 2 1.316 -3.987 0.294 1.00 0.00 N
 ATOM 37 C2 DA A 2 0.481 -3.492 -0.593 1.00 0.00 C
 ATOM 38 N3 DA A 2 -0.026 -2.218 -0.678 1.00 0.00 N
 ATOM 39 C4 DA A 2 0.426 -1.448 0.271 1.00 0.00 C
 ATOM 40 P DC A 3 -4.652 1.865 -0.622 1.00 0.00 P

ATOM 41 OP1 DC A 3 -5.230 3.256 -0.653 1.00 0.00 O
 ATOM 42 OP2 DC A 3 -4.679 1.223 0.748 1.00 0.00 O
 ATOM 43 O5' DC A 3 -5.411 0.953 -1.720 1.00 0.00 O
 ATOM 44 C5' DC A 3 -4.775 0.042 -2.616 1.00 0.00 C
 ATOM 45 C4' DC A 3 -4.550 -1.324 -2.010 1.00 0.00 C
 ATOM 46 O4' DC A 3 -3.500 -1.282 -1.042 1.00 0.00 O
 ATOM 47 C3' DC A 3 -5.785 -1.932 -1.299 1.00 0.00 C
 ATOM 48 O3' DC A 3 -6.047 -3.173 -1.934 1.00 0.00 O
 ATOM 49 C2' DC A 3 -5.297 -2.110 0.140 1.00 0.00 C
 ATOM 50 C1' DC A 3 -3.807 -2.311 -0.109 1.00 0.00 C
 ATOM 51 N1 DC A 3 -2.928 -2.205 1.040 1.00 0.00 N
 ATOM 52 C2 DC A 3 -1.997 -3.243 1.289 1.00 0.00 C
 ATOM 53 O2 DC A 3 -1.944 -4.206 0.527 1.00 0.00 O
 ATOM 54 N3 DC A 3 -1.168 -3.115 2.363 1.00 0.00 N
 ATOM 55 C4 DC A 3 -1.267 -2.058 3.166 1.00 0.00 C
 ATOM 56 N4 DC A 3 -0.447 -2.013 4.223 1.00 0.00 N
 ATOM 57 C5 DC A 3 -2.182 -0.983 2.923 1.00 0.00 C
 ATOM 58 C6 DC A 3 -2.992 -1.107 1.842 1.00 0.00 C
 END

COMPND CCA / M06-2X / NA MODEL / NEUTRAL

ATOM 1 O5' DC A 1 -7.314 -0.919 -0.443 1.00 0.00 O
 ATOM 2 C5' DC A 1 -6.920 -1.858 -1.430 1.00 0.00 C
 ATOM 3 C4' DC A 1 -5.405 -1.906 -1.487 1.00 0.00 C
 ATOM 4 O4' DC A 1 -4.898 -0.663 -1.975 1.00 0.00 O
 ATOM 5 C3' DC A 1 -4.730 -2.120 -0.125 1.00 0.00 C
 ATOM 6 O3' DC A 1 -3.492 -2.794 -0.388 1.00 0.00 O
 ATOM 7 C2' DC A 1 -4.461 -0.695 0.332 1.00 0.00 C
 ATOM 8 C1' DC A 1 -4.071 -0.044 -0.993 1.00 0.00 C
 ATOM 9 N1 DC A 1 -4.276 1.392 -1.040 1.00 0.00 N
 ATOM 10 C2 DC A 1 -3.224 2.227 -1.496 1.00 0.00 C
 ATOM 11 O2 DC A 1 -2.210 1.715 -1.960 1.00 0.00 O
 ATOM 12 N3 DC A 1 -3.397 3.574 -1.382 1.00 0.00 N
 ATOM 13 C4 DC A 1 -4.552 4.065 -0.965 1.00 0.00 C
 ATOM 14 N4 DC A 1 -4.644 5.399 -0.817 1.00 0.00 N
 ATOM 15 C5 DC A 1 -5.692 3.245 -0.664 1.00 0.00 C
 ATOM 16 C6 DC A 1 -5.494 1.908 -0.714 1.00 0.00 C
 ATOM 17 P DC A 2 -2.622 -3.369 0.860 1.00 0.00 P
 ATOM 18 OP1 DC A 2 -3.337 -4.512 1.534 1.00 0.00 O
 ATOM 19 OP2 DC A 2 -2.210 -2.266 1.811 1.00 0.00 O
 ATOM 20 O5' DC A 2 -1.351 -3.950 0.048 1.00 0.00 O
 ATOM 21 C5' DC A 2 -0.948 -3.598 -1.276 1.00 0.00 C
 ATOM 22 C4' DC A 2 0.111 -2.515 -1.298 1.00 0.00 C
 ATOM 23 O4' DC A 2 -0.485 -1.233 -1.092 1.00 0.00 O
 ATOM 24 C3' DC A 2 1.200 -2.668 -0.209 1.00 0.00 C
 ATOM 25 O3' DC A 2 2.462 -2.456 -0.859 1.00 0.00 O
 ATOM 26 C2' DC A 2 0.884 -1.515 0.736 1.00 0.00 C
 ATOM 27 C1' DC A 2 0.436 -0.485 -0.293 1.00 0.00 C
 ATOM 28 N1 DC A 2 -0.208 0.717 0.183 1.00 0.00 N
 ATOM 29 C2 DC A 2 0.232 1.971 -0.318 1.00 0.00 C
 ATOM 30 O2 DC A 2 1.160 2.012 -1.119 1.00 0.00 O
 ATOM 31 N3 DC A 2 -0.387 3.093 0.151 1.00 0.00 N
 ATOM 32 C4 DC A 2 -1.402 2.994 0.984 1.00 0.00 C
 ATOM 33 N4 DC A 2 -1.939 4.150 1.458 1.00 0.00 N
 ATOM 34 C5 DC A 2 -1.911 1.740 1.457 1.00 0.00 C
 ATOM 35 C6 DC A 2 -1.269 0.626 1.023 1.00 0.00 C
 ATOM 36 P DA A 3 3.848 -2.871 -0.107 1.00 0.00 P
 ATOM 37 OP1 DA A 3 4.350 -4.189 -0.636 1.00 0.00 O
 ATOM 38 OP2 DA A 3 3.712 -2.756 1.387 1.00 0.00 O
 ATOM 39 O5' DA A 3 4.794 -11.644 -0.601 1.00 0.00 O
 ATOM 40 C5' DA A 3 4.959 -1.405 -1.999 1.00 0.00 C
 ATOM 41 C4' DA A 3 5.519 -0.010 -2.187 1.00 0.00 C
 ATOM 42 O4' DA A 3 4.565 0.953 -1.729 1.00 0.00 O
 ATOM 43 C3' DA A 3 6.813 0.278 -1.419 1.00 0.00 C
 ATOM 44 O3' DA A 3 7.539 1.210 -2.203 1.00 0.00 O
 ATOM 45 C2' DA A 3 6.299 0.915 -0.134 1.00 0.00 C
 ATOM 46 C1' DA A 3 5.077 1.689 -0.632 1.00 0.00 C
 ATOM 47 N9 DA A 3 4.043 1.839 0.370 1.00 0.00 N
 ATOM 48 C8 DA A 3 3.407 0.830 1.053 1.00 0.00 C
 ATOM 49 N7 DA A 3 2.479 1.253 1.875 1.00 0.00 N
 ATOM 50 C5 DA A 3 2.498 2.624 1.718 1.00 0.00 C
 ATOM 51 C6 DA A 3 1.715 3.654 2.269 1.00 0.00 C
 ATOM 52 N6 DA A 3 0.757 3.408 3.201 1.00 0.00 N
 ATOM 53 N1 DA A 3 1.949 4.917 1.886 1.00 0.00 N
 ATOM 54 C2 DA A 3 2.911 5.137 0.977 1.00 0.00 C
 ATOM 55 N3 DA A 3 3.704 4.258 0.370 1.00 0.00 N
 ATOM 56 C4 DA A 3 3.448 3.009 0.775 1.00 0.00 C
 END

COMPND CCA / M06-2X / NA MODEL / SINGLY IONIZED

ATOM 1 O5' DC A 1 -7.306 -0.737 -0.602 1.00 0.00 O
 ATOM 2 C5' DC A 1 -6.900 1.695 -1.565 1.00 0.00 C
 ATOM 3 C4' DC A 1 -5.387 -1.790 -1.557 1.00 0.00 C
 ATOM 4 O4' DC A 1 -4.823 -0.565 -2.034 1.00 0.00 O
 ATOM 5 C3' DC A 1 -4.779 -2.012 -0.166 1.00 0.00 C
 ATOM 6 O3' DC A 1 -3.556 -2.734 -0.365 1.00 0.00 O
 ATOM 7 C2' DC A 1 -4.479 -0.592 0.288 1.00 0.00 C
 ATOM 8 C1' DC A 1 -4.014 0.033 -1.026 1.00 0.00 C
 ATOM 9 N1 DC A 1 -4.172 1.476 -1.090 1.00 0.00 N
 ATOM 10 C2 DC A 1 -3.073 2.284 -1.480 1.00 0.00 C
 ATOM 11 O2 DC A 1 -2.045 1.749 -1.885 1.00 0.00 O
 ATOM 12 N3 DC A 1 -3.217 3.635 -1.371 1.00 0.00 N
 ATOM 13 C4 DC A 1 -4.386 4.157 -1.036 1.00 0.00 C
 ATOM 14 N4 DC A 1 -4.456 5.492 -0.908 1.00 0.00 N
 ATOM 15 C5 DC A 1 -5.565 3.365 -0.814 1.00 0.00 C
 ATOM 16 C6 DC A 1 -5.396 2.024 -0.847 1.00 0.00 C
 ATOM 17 P DC A 2 -2.769 3.334 0.925 1.00 0.00 P
 ATOM 18 OP1 DC A 2 -3.588 -4.401 1.606 1.00 0.00 O
 ATOM 19 OP2 DC A 2 -2.303 -2.237 1.860 1.00 0.00 O
 ATOM 20 O5' DC A 2 -1.515 -4.023 0.174 1.00 0.00 O
 ATOM 21 C5' DC A 2 -1.081 -3.764 -1.159 1.00 0.00 C
 ATOM 22 C4' DC A 2 0.012 -2.719 -1.222 1.00 0.00 C
 ATOM 23 O4' DC A 2 -0.541 -1.412 -1.043 1.00 0.00 O
 ATOM 24 C3' DC A 2 1.121 -2.876 -0.149 1.00 0.00 C
 ATOM 25 O3' DC A 2 2.366 -2.803 -0.859 1.00 0.00 O
 ATOM 26 C2' DC A 2 0.891 -1.659 0.737 1.00 0.00 C
 ATOM 27 C1' DC A 2 0.431 -0.664 -0.319 1.00 0.00 C
 ATOM 28 N1 DC A 2 -0.150 0.570 0.162 1.00 0.00 N
 ATOM 29 C2 DC A 2 0.434 1.797 -0.225 1.00 0.00 C
 ATOM 30 O2 DC A 2 1.423 1.795 -0.956 1.00 0.00 O
 ATOM 31 N3 DC A 2 -0.078 2.944 0.303 1.00 0.00 N
 ATOM 32 C4 DC A 2 -1.148 2.899 1.083 1.00 0.00 C
 ATOM 33 N4 DC A 2 -1.564 4.065 1.620 1.00 0.00 N
 ATOM 34 C5 DC A 2 -1.800 1.676 1.442 1.00 0.00 C
 ATOM 35 C6 DC A 2 -1.243 0.533 0.969 1.00 0.00 C
 ATOM 36 P DA A 3 3.816 -2.876 -0.123 1.00 0.00 P
 ATOM 37 OP1 DA A 3 4.570 -4.080 -0.616 1.00 0.00 O
 ATOM 38 OP2 DA A 3 3.709 -2.699 1.367 1.00 0.00 O

ATOM 39 O5' DA A 3 4.493 -1.504 -0.693 1.00 0.00 O
 ATOM 40 C5' DA A 3 4.608 -1.313 -2.100 1.00 0.00 C
 ATOM 41 C4' DA A 3 5.227 0.048 -2.344 1.00 0.00 C
 ATOM 42 O4' DA A 3 4.369 1.065 -1.810 1.00 0.00 O
 ATOM 43 C3' DA A 3 6.592 0.275 -1.689 1.00 0.00 C
 ATOM 44 O3' DA A 3 7.275 1.196 -2.521 1.00 0.00 O
 ATOM 45 C2' DA A 3 6.222 0.908 -0.351 1.00 0.00 C
 ATOM 46 C1' DA A 3 4.977 1.716 -0.724 1.00 0.00 C
 ATOM 47 N9 DA A 3 4.015 1.817 0.381 1.00 0.00 N
 ATOM 48 C8 DA A 3 3.489 0.813 1.110 1.00 0.00 C
 ATOM 49 N7 DA A 3 2.566 1.209 2.010 1.00 0.00 N
 ATOM 50 C5 DA A 3 2.489 2.525 1.826 1.00 0.00 C
 ATOM 51 C6 DA A 3 1.701 3.553 2.445 1.00 0.00 C
 ATOM 52 N6 DA A 3 0.876 3.291 3.431 1.00 0.00 N
 ATOM 53 N1 DA A 3 1.831 4.831 2.003 1.00 0.00 N
 ATOM 54 C2 DA A 3 2.678 5.068 1.029 1.00 0.00 C
 ATOM 55 N3 DA A 3 3.502 4.190 0.366 1.00 0.00 N
 ATOM 56 C4 DA A 3 3.364 2.968 0.785 1.00 0.00 C
 END

COMPND CAA / M06-2X / NA MODEL / NEUTRAL
 ATOM 1 O5' DC A 1 -7.276 -0.856 -0.308 1.00 0.00 O
 ATOM 2 C5' DC A 1 -6.931 -1.817 -1.290 1.00 0.00 C
 ATOM 3 C4' DC A 1 -5.419 -1.909 -1.379 1.00 0.00 C
 ATOM 4 O4' DC A 1 -4.891 -0.684 -1.892 1.00 0.00 O
 ATOM 5 C3' DC A 1 -4.725 -2.124 -0.028 1.00 0.00 C
 ATOM 6 O3' DC A 1 -3.510 -2.838 -0.296 1.00 0.00 O
 ATOM 7 C2' DC A 1 -4.408 -0.702 0.407 1.00 0.00 C
 ATOM 8 C1' DC A 1 -4.031 -0.076 -0.933 1.00 0.00 C
 ATOM 9 N1 DC A 1 -4.193 1.366 -0.993 1.00 0.00 N
 ATOM 10 C2 DC A 1 -3.164 2.150 -1.576 1.00 0.00 C
 ATOM 11 O2 DC A 1 -2.228 1.592 -2.139 1.00 0.00 O
 ATOM 12 N3 DC A 1 -3.267 3.505 -1.465 1.00 0.00 N
 ATOM 13 C4 DC A 1 -4.362 4.049 -0.963 1.00 0.00 C
 ATOM 14 N4 DC A 1 -4.386 5.388 -0.826 1.00 0.00 N
 ATOM 15 C5 DC A 1 -5.507 3.281 -0.557 1.00 0.00 C
 ATOM 16 C6 DC A 1 -5.362 1.937 -0.588 1.00 0.00 C
 ATOM 17 P DA A 2 -2.703 -3.486 0.957 1.00 0.00 P
 ATOM 18 OP1 DA A 2 -3.507 -4.590 1.597 1.00 0.00 O
 ATOM 19 OP2 DA A 2 -2.242 -2.427 1.938 1.00 0.00 O
 ATOM 20 O5' DA A 2 -1.431 -4.107 0.179 1.00 0.00 O
 ATOM 21 C5' DA A 2 -1.086 -3.951 -1.196 1.00 0.00 C
 ATOM 22 C4' DA A 2 0.058 -2.973 -1.373 1.00 0.00 C
 ATOM 23 O4' DA A 2 -0.421 -1.627 -1.267 1.00 0.00 O
 ATOM 24 C3' DA A 2 1.171 -3.120 -0.311 1.00 0.00 C
 ATOM 25 O3' DA A 2 2.414 -2.999 -1.014 1.00 0.00 O
 ATOM 26 C2' DA A 2 0.907 -1.927 0.598 1.00 0.00 C
 ATOM 27 C1' DA A 2 0.480 -0.899 -0.439 1.00 0.00 C
 ATOM 28 N9 DA A 2 -0.177 0.275 0.065 1.00 0.00 N
 ATOM 29 C8 DA A 2 -1.150 0.340 1.032 1.00 0.00 C
 ATOM 30 N7 DA A 2 -1.534 1.565 1.307 1.00 0.00 N
 ATOM 31 C5 DA A 2 -0.755 2.352 0.478 1.00 0.00 C
 ATOM 32 C6 DA A 2 -0.663 3.741 0.292 1.00 0.00 C
 ATOM 33 N6 DA A 2 -1.414 4.617 1.014 1.00 0.00 N
 ATOM 34 N1 DA A 2 0.244 4.217 -0.574 1.00 0.00 N
 ATOM 35 C2 DA A 2 1.014 3.340 -1.235 1.00 0.00 C
 ATOM 36 N3 DA A 2 1.007 2.011 -1.170 1.00 0.00 N
 ATOM 37 C4 DA A 2 0.096 1.571 -0.298 1.00 0.00 C
 ATOM 38 P DA A 3 3.848 -3.102 -0.242 1.00 0.00 P
 ATOM 39 OP1 DA A 3 4.667 -4.193 -0.876 1.00 0.00 O
 ATOM 40 OP2 DA A 3 3.669 -3.158 1.249 1.00 0.00 O
 ATOM 41 O5' DA A 3 4.479 -1.640 -0.601 1.00 0.00 O
 ATOM 42 C5' DA A 3 4.670 -1.316 -1.978 1.00 0.00 C
 ATOM 43 C4' DA A 3 5.278 0.067 -2.083 1.00 0.00 C
 ATOM 44 O4' DA A 3 4.363 1.039 -1.570 1.00 0.00 O
 ATOM 45 C3' DA A 3 6.582 0.261 -1.300 1.00 0.00 C
 ATOM 46 O3' DA A 3 7.340 1.214 -2.026 1.00 0.00 O
 ATOM 47 C2' DA A 3 6.089 0.834 0.023 1.00 0.00 C
 ATOM 48 C1' DA A 3 4.896 1.681 -0.425 1.00 0.00 C
 ATOM 49 N9 DA A 3 3.857 1.793 0.578 1.00 0.00 N
 ATOM 50 C8 DA A 3 3.153 0.757 1.145 1.00 0.00 C
 ATOM 51 N7 DA A 3 2.199 1.147 1.953 1.00 0.00 N
 ATOM 52 C5 DA A 3 2.273 2.526 1.911 1.00 0.00 C
 ATOM 53 C6 DA A 3 1.501 3.544 2.502 1.00 0.00 C
 ATOM 54 N6 DA A 3 0.445 3.277 3.317 1.00 0.00 N
 ATOM 55 N1 DA A 3 1.831 4.822 2.264 1.00 0.00 N
 ATOM 56 C2 DA A 3 2.851 5.075 1.432 1.00 0.00 C
 ATOM 57 N3 DA A 3 3.625 4.214 0.777 1.00 0.00 N
 ATOM 58 C4 DA A 3 3.291 2.948 1.056 1.00 0.00 C
 END

COMPND CAA / M06-2X / NA MODEL / SINGLY IONIZED
 ATOM 1 O5' DC A 1 -6.984 -1.619 -0.498 1.00 0.00 O
 ATOM 2 C5' DC A 1 -6.409 -2.463 -1.482 1.00 0.00 C
 ATOM 3 C4' DC A 1 -4.906 -2.262 -1.469 1.00 0.00 C
 ATOM 4 O4' DC A 1 -4.593 -0.945 -1.932 1.00 0.00 O
 ATOM 5 C3' DC A 1 -4.279 -2.372 -0.074 1.00 0.00 C
 ATOM 6 O3' DC A 1 -2.942 -2.860 -0.244 1.00 0.00 O
 ATOM 7 C2' DC A 1 -4.232 -0.924 0.389 1.00 0.00 C
 ATOM 8 C1' DC A 1 -3.916 -0.209 -0.925 1.00 0.00 C
 ATOM 9 N1 DC A 1 -4.376 1.173 -0.980 1.00 0.00 N
 ATOM 10 C2 DC A 1 -3.476 2.205 -1.312 1.00 0.00 C
 ATOM 11 O2 DC A 1 -2.305 1.919 -1.592 1.00 0.00 O
 ATOM 12 N3 DC A 1 -3.924 3.484 -1.289 1.00 0.00 N
 ATOM 13 C4 DC A 1 -5.208 3.735 -1.054 1.00 0.00 C
 ATOM 14 N4 DC A 1 -5.593 5.017 -1.030 1.00 0.00 N
 ATOM 15 C5 DC A 1 -6.176 2.698 -0.842 1.00 0.00 C
 ATOM 16 C6 DC A 1 -5.705 1.431 -0.812 1.00 0.00 C
 ATOM 17 P DA A 2 -2.271 -3.653 1.007 1.00 0.00 P
 ATOM 18 OP1 DA A 2 -3.148 -4.800 1.432 1.00 0.00 O
 ATOM 19 OP2 DA A 2 -1.902 -2.705 2.129 1.00 0.00 O
 ATOM 20 O5' DA A 2 -0.903 -4.159 0.317 1.00 0.00 O
 ATOM 21 C5' DA A 2 -0.593 -4.167 -1.075 1.00 0.00 C
 ATOM 22 C4' DA A 2 0.519 -3.173 -1.352 1.00 0.00 C
 ATOM 23 O4' DA A 2 -0.003 -1.835 -1.306 1.00 0.00 O
 ATOM 24 C3' DA A 2 1.660 -3.220 -0.316 1.00 0.00 C
 ATOM 25 O3' DA A 2 2.868 -3.022 -1.054 1.00 0.00 O
 ATOM 26 C2' DA A 2 1.319 -2.043 0.592 1.00 0.00 C
 ATOM 27 C1' DA A 2 0.797 -1.057 -0.440 1.00 0.00 C
 ATOM 28 N9 DA A 2 -0.015 0.017 0.106 1.00 0.00 N
 ATOM 29 C8 DA A 2 -0.916 -0.049 1.117 1.00 0.00 C
 ATOM 30 N7 DA A 2 -1.466 1.135 1.433 1.00 0.00 N
 ATOM 31 C5 DA A 2 -0.882 1.993 0.595 1.00 0.00 C
 ATOM 32 C6 DA A 2 -1.036 3.401 0.375 1.00 0.00 C
 ATOM 33 N6 DA A 2 -1.895 4.106 1.073 1.00 0.00 N
 ATOM 34 N1 DA A 2 -0.273 4.004 -0.569 1.00 0.00 N
 ATOM 35 C2 DA A 2 0.556 3.257 -1.267 1.00 0.00 C
 ATOM 36 N3 DA A 2 0.775 1.907 -1.191 1.00 0.00 N

ATOM 37 C4 DA A 2 0.049 1.343 -0.272 1.00 0.00 C
 ATOM 38 P DA A 3 4.304 -2.733 -0.332 1.00 0.00 P
 ATOM 39 O¹ DA A 3 5.354 -3.538 -1.046 1.00 0.00 O
 ATOM 40 O² DA A 3 4.217 -2.874 1.161 1.00 0.00 O
 ATOM 41 O⁵ DA A 3 4.489 -1.142 -0.658 1.00 0.00 O
 ATOM 42 C⁵ DA A 3 4.605 -0.768 -2.033 1.00 0.00 C
 ATOM 43 C⁴ DA A 3 5.027 0.683 -2.119 1.00 0.00 C
 ATOM 44 O⁴ DA A 3 4.011 1.515 -1.552 1.00 0.00 O
 ATOM 45 C³ DA A 3 6.320 1.021 -1.369 1.00 0.00 C
 ATOM 46 O³ DA A 3 6.934 2.071 -2.095 1.00 0.00 O
 ATOM 47 C² DA A 3 5.802 1.506 -0.020 1.00 0.00 C
 ATOM 48 C¹ DA A 3 4.503 2.206 -0.419 1.00 0.00 C
 ATOM 49 N⁹ DA A 3 3.490 2.182 0.620 1.00 0.00 N
 ATOM 50 C⁸ DA A 3 2.990 1.069 1.253 1.00 0.00 C
 ATOM 51 N⁷ DA A 3 2.015 1.331 2.088 1.00 0.00 N
 ATOM 52 C⁵ DA A 3 1.860 2.703 1.996 1.00 0.00 C
 ATOM 53 C⁶ DA A 3 0.988 3.617 2.625 1.00 0.00 C
 ATOM 54 N⁶ DA A 3 0.028 3.221 3.485 1.00 0.00 N
 ATOM 55 N¹ DA A 3 1.077 4.917 2.301 1.00 0.00 N
 ATOM 56 C² DA A 3 1.992 5.291 1.396 1.00 0.00 C
 ATOM 57 N³ DA A 3 2.870 4.538 0.738 1.00 0.00 N
 ATOM 58 C⁴ DA A 3 2.761 3.249 1.081 1.00 0.00 N
 END
 COMPND ACC / M06-2X / PO- MODEL / NEUTRAL
 ATOM 1 O⁵ DA A 1 -7.345 -0.356 0.238 1.00 0.00 O
 ATOM 2 C⁵ DA A 1 -7.088 -1.316 -0.770 1.00 0.00 C
 ATOM 3 C⁴ DA A 1 -5.591 -1.521 -0.882 1.00 0.00 C
 ATOM 4 O⁴ DA A 1 -4.979 -0.326 -1.384 1.00 0.00 O
 ATOM 5 C³ DA A 1 -4.893 -1.817 0.455 1.00 0.00 C
 ATOM 6 O³ DA A 1 -3.776 -2.649 0.159 1.00 0.00 O
 ATOM 7 C² DA A 1 -4.423 -0.433 0.884 1.00 0.00 C
 ATOM 8 C¹ DA A 1 -3.985 0.111 -0.467 1.00 0.00 C
 ATOM 9 N⁹ DA A 1 -3.863 1.550 -0.533 1.00 0.00 N
 ATOM 10 C⁸ DA A 1 -4.436 2.509 0.270 1.00 0.00 C
 ATOM 11 N⁷ DA A 1 -4.003 3.722 0.039 1.00 0.00 N
 ATOM 12 C⁵ DA A 1 -3.089 3.557 -0.984 1.00 0.00 C
 ATOM 13 C⁶ DA A 1 -2.215 4.447 -1.636 1.00 0.00 C
 ATOM 14 N⁶ DA A 1 -2.114 5.748 -1.266 1.00 0.00 N
 ATOM 15 N¹ DA A 1 -1.414 3.976 -2.600 1.00 0.00 N
 ATOM 16 C² DA A 1 -1.435 2.660 -2.857 1.00 0.00 C
 ATOM 17 N³ DA A 1 -2.170 1.711 -2.284 1.00 0.00 N
 ATOM 18 C⁴ DA A 1 -2.984 2.220 -1.352 1.00 0.00 C
 ATOM 19 P DC A 2 -2.907 -3.279 1.407 1.00 0.00 P
 ATOM 20 O² DC A 2 -2.163 -2.194 2.139 1.00 0.00 O
 ATOM 21 O¹ DC A 2 -3.772 -4.217 2.191 1.00 0.00 O
 ATOM 22 O⁵ DC A 2 -1.864 -4.171 0.520 1.00 0.00 O
 ATOM 23 C⁵ DC A 2 -1.548 -3.963 -0.848 1.00 0.00 C
 ATOM 24 C⁴ DC A 2 -0.401 -2.999 -1.074 1.00 0.00 C
 ATOM 25 O⁴ DC A 2 -0.834 -1.645 -0.910 1.00 0.00 O
 ATOM 26 C³ DC A 2 0.810 -3.199 -0.128 1.00 0.00 C
 ATOM 27 O³ DC A 2 1.955 -3.386 -0.958 1.00 0.00 O
 ATOM 28 C² DC A 2 0.841 -1.887 0.651 1.00 0.00 C
 ATOM 29 C¹ DC A 2 0.286 -0.933 -0.399 1.00 0.00 C
 ATOM 30 N¹ DC A 2 -0.134 0.372 0.073 1.00 0.00 N
 ATOM 31 C² DC A 2 0.426 1.534 -0.522 1.00 0.00 C
 ATOM 32 O² DC A 2 1.206 1.417 -1.461 1.00 0.00 O
 ATOM 33 N³ DC A 2 0.071 2.746 -0.003 1.00 0.00 N
 ATOM 34 C⁴ DC A 2 -0.794 2.822 0.992 1.00 0.00 C
 ATOM 35 N⁴ DC A 2 -1.108 4.054 1.460 1.00 0.00 N
 ATOM 36 C⁵ DC A 2 -1.377 1.666 1.611 1.00 0.00 C
 ATOM 37 C⁶ DC A 2 -1.016 0.458 1.105 1.00 0.00 C
 ATOM 38 P DC A 3 3.451 -3.493 -0.278 1.00 0.00 P
 ATOM 39 O¹ DC A 3 4.264 -4.375 -1.171 1.00 0.00 O
 ATOM 40 O² DC A 3 3.333 -3.710 1.195 1.00 0.00 O
 ATOM 41 O⁵ DC A 3 3.962 -1.930 -0.469 1.00 0.00 O
 ATOM 42 C⁵ DC A 3 4.229 -1.489 -1.795 1.00 0.00 C
 ATOM 43 C⁴ DC A 3 5.240 -0.361 -1.771 1.00 0.00 C
 ATOM 44 O⁴ DC A 3 4.649 0.825 -1.233 1.00 0.00 O
 ATOM 45 C³ DC A 3 6.477 -0.646 -0.912 1.00 0.00 C
 ATOM 46 O³ DC A 3 7.566 0.012 -1.539 1.00 0.00 O
 ATOM 47 C² DC A 3 6.115 0.015 0.414 1.00 0.00 C
 ATOM 48 C¹ DC A 3 5.311 1.232 -0.049 1.00 0.00 C
 ATOM 49 N¹ DC A 3 4.298 1.675 0.906 1.00 0.00 N
 ATOM 50 C² DC A 3 4.373 2.969 1.470 1.00 0.00 C
 ATOM 51 O² DC A 3 5.332 3.688 1.202 1.00 0.00 O
 ATOM 52 N³ DC A 3 3.359 3.359 2.295 1.00 0.00 N
 ATOM 53 C⁴ DC A 3 2.372 2.526 2.579 1.00 0.00 C
 ATOM 54 N⁴ DC A 3 1.410 2.954 3.422 1.00 0.00 N
 ATOM 55 C⁵ DC A 3 2.291 1.194 2.054 1.00 0.00 C
 ATOM 56 C⁶ DC A 3 3.269 0.832 1.192 1.00 0.00 C
 END
 COMPND ACC / M06-2X / PO- MODEL / SINGLY IONIZED
 ATOM 1 O⁵ DA A 1 -7.125 -0.479 0.519 1.00 0.00 O
 ATOM 2 C⁵ DA A 1 -7.010 -1.611 -0.327 1.00 0.00 C
 ATOM 3 C⁴ DA A 1 -5.543 -1.918 -0.560 1.00 0.00 C
 ATOM 4 O⁴ DA A 1 -4.924 -0.838 -1.279 1.00 0.00 O
 ATOM 5 C³ DA A 1 -4.697 -2.083 0.705 1.00 0.00 C
 ATOM 6 O³ DA A 1 -3.590 -2.901 0.345 1.00 0.00 O
 ATOM 7 C² DA A 1 -4.229 -0.658 0.969 1.00 0.00 C
 ATOM 8 C¹ DA A 1 -3.989 -0.178 -0.460 1.00 0.00 C
 ATOM 9 N⁹ DA A 1 -4.181 1.269 -0.610 1.00 0.00 N
 ATOM 10 C⁸ DA A 1 -5.242 1.998 -0.210 1.00 0.00 C
 ATOM 11 N⁷ DA A 1 -5.130 3.316 -0.464 1.00 0.00 N
 ATOM 12 C⁵ DA A 1 -3.945 3.424 -1.066 1.00 0.00 C
 ATOM 13 C⁶ DA A 1 -3.218 4.540 -1.603 1.00 0.00 C
 ATOM 14 N⁶ DA A 1 -3.677 5.766 -1.527 1.00 0.00 N
 ATOM 15 N¹ DA A 1 -2.025 4.310 -2.210 1.00 0.00 N
 ATOM 16 C² DA A 1 -1.581 3.076 -2.277 1.00 0.00 C
 ATOM 17 N³ DA A 1 -2.176 1.922 -1.811 1.00 0.00 N
 ATOM 18 C⁴ DA A 1 -3.306 2.155 -1.212 1.00 0.00 C
 ATOM 19 P DC A 2 -2.583 -3.437 1.535 1.00 0.00 P
 ATOM 20 O² DC A 2 -1.894 -2.271 2.193 1.00 0.00 O
 ATOM 21 O¹ DC A 2 -3.304 -4.427 2.392 1.00 0.00 O
 ATOM 22 O⁵ DC A 2 -1.527 -4.248 0.592 1.00 0.00 O
 ATOM 23 C⁵ DC A 2 -1.340 -4.077 -0.807 1.00 0.00 C
 ATOM 24 C⁴ DC A 2 -0.206 -3.132 -1.153 1.00 0.00 C
 ATOM 25 O⁴ DC A 2 -0.629 -1.766 -1.024 1.00 0.00 O
 ATOM 26 C³ DC A 2 1.061 -3.290 -0.273 1.00 0.00 C
 ATOM 27 O³ DC A 2 2.171 -3.389 -1.167 1.00 0.00 O
 ATOM 28 C² DC A 2 1.036 -2.002 0.548 1.00 0.00 C
 ATOM 29 C¹ DC A 2 0.483 -1.048 -0.500 1.00 0.00 C
 ATOM 30 N¹ DC A 2 0.039 0.249 -0.027 1.00 0.00 N
 ATOM 31 C² DC A 2 0.530 1.422 -0.651 1.00 0.00 C
 ATOM 32 O² DC A 2 1.296 1.338 -1.604 1.00 0.00 O
 ATOM 33 N³ DC A 2 0.104 2.628 -0.164 1.00 0.00 N
 ATOM 34 C⁴ DC A 2 -0.812 2.678 0.790 1.00 0.00 C
 ATOM 35 N⁴ DC A 2 -1.321 3.893 1.086 1.00 0.00 N

ATOM 36 C5 DC A 2 -1.330 1.509 1.436 1.00 0.00 C
 ATOM 37 C6 DC A 2 -0.870 0.315 0.982 1.00 0.00 C
 ATOM 38 P DC A 3 3.705 -3.051 -0.675 1.00 0.00 P
 ATOM 39 OP1 DC A 3 4.614 -3.636 -1.707 1.00 0.00 O
 ATOM 40 OP2 DC A 3 3.853 -3.320 0.789 1.00 0.00 O
 ATOM 41 O5' DC A 3 3.701 -1.407 -0.829 1.00 0.00 O
 ATOM 42 C5' DC A 3 4.054 -0.795 -2.065 1.00 0.00 C
 ATOM 43 C4' DC A 3 5.224 0.147 -1.846 1.00 0.00 C
 ATOM 44 O4' DC A 3 4.801 1.298 -1.109 1.00 0.00 O
 ATOM 45 C3' DC A 3 6.366 -0.475 -1.035 1.00 0.00 C
 ATOM 46 O3' DC A 3 7.568 0.119 -1.502 1.00 0.00 O
 ATOM 47 C2' DC A 3 6.041 -0.032 0.390 1.00 0.00 C
 ATOM 48 C1' DC A 3 5.426 1.352 0.162 1.00 0.00 C
 ATOM 49 N1 DC A 3 4.416 1.739 1.146 1.00 0.00 N
 ATOM 50 C2 DC A 3 4.555 2.950 1.866 1.00 0.00 C
 ATOM 51 O2 DC A 3 5.595 3.596 1.763 1.00 0.00 O
 ATOM 52 N3 DC A 3 3.510 3.347 2.649 1.00 0.00 N
 ATOM 53 C4 DC A 3 2.428 2.594 2.749 1.00 0.00 C
 ATOM 54 N4 DC A 3 1.407 3.051 3.496 1.00 0.00 N
 ATOM 55 C5 DC A 3 2.284 1.334 2.080 1.00 0.00 C
 ATOM 56 C6 DC A 3 3.294 0.975 1.257 1.00 0.00 C
 END

COMPND AAC / M06-2X / PO- MODEL / NEUTRAL

ATOM 1 O5' DA A 1 -7.349 -0.395 0.191 1.00 0.00 O
 ATOM 2 C5' DA A 1 -6.992 -1.252 -0.877 1.00 0.00 C
 ATOM 3 C4' DA A 1 -5.484 -1.388 -0.903 1.00 0.00 C
 ATOM 4 O4' DA A 1 -4.898 -0.137 -1.290 1.00 0.00 O
 ATOM 5 C3' DA A 1 -4.867 -1.737 0.464 1.00 0.00 C
 ATOM 6 O3' DA A 1 -3.733 -2.562 0.213 1.00 0.00 O
 ATOM 7 C2' DA A 1 -4.416 -0.371 0.973 1.00 0.00 C
 ATOM 8 C1' DA A 1 -3.904 0.208 -0.336 1.00 0.00 C
 ATOM 9 N9 DA A 1 -3.705 1.658 -0.348 1.00 0.00 N
 ATOM 10 C8 DA A 1 -4.064 2.583 0.584 1.00 0.00 C
 ATOM 11 N7 DA A 1 -3.620 3.785 0.317 1.00 0.00 N
 ATOM 12 C5 DA A 1 -2.921 3.626 -0.865 1.00 0.00 C
 ATOM 13 C6 DA A 1 -2.148 4.506 -1.647 1.00 0.00 C
 ATOM 14 N6 DA A 1 -1.970 5.804 -1.309 1.00 0.00 N
 ATOM 15 N1 DA A 1 -1.523 4.022 -2.730 1.00 0.00 N
 ATOM 16 C2 DA A 1 -1.638 2.714 -3.005 1.00 0.00 C
 ATOM 17 N3 DA A 1 -2.325 1.782 -2.350 1.00 0.00 N
 ATOM 18 C4 DA A 1 -2.950 2.301 -1.286 1.00 0.00 C
 ATOM 19 P DA A 2 -3.046 -3.340 1.489 1.00 0.00 P
 ATOM 20 OP1 DA A 2 -4.031 -4.316 2.053 1.00 0.00 O
 ATOM 21 OP2 DA A 2 -2.378 -2.354 2.411 1.00 0.00 O
 ATOM 22 O5' DA A 2 -1.864 -4.129 0.684 1.00 0.00 O
 ATOM 23 C5' DA A 2 -1.701 -4.209 -0.724 1.00 0.00 C
 ATOM 24 C4' DA A 2 -0.498 -3.399 -1.169 1.00 0.00 C
 ATOM 25 O4' DA A 2 -0.804 -1.996 -1.113 1.00 0.00 O
 ATOM 26 C3' DA A 2 0.751 -3.608 -0.283 1.00 0.00 C
 ATOM 27 O3' DA A 2 1.875 -3.673 -1.163 1.00 0.00 O
 ATOM 28 C2' DA A 2 0.728 -2.352 0.582 1.00 0.00 C
 ATOM 29 C1' DA A 2 0.264 -1.338 -0.449 1.00 0.00 C
 ATOM 30 N9 DA A 2 -0.190 -0.081 0.094 1.00 0.00 N
 ATOM 31 C8 DA A 2 -1.016 0.115 1.178 1.00 0.00 C
 ATOM 32 N7 DA A 2 -1.140 1.379 1.513 1.00 0.00 N
 ATOM 33 C5 DA A 2 -0.338 2.049 0.607 1.00 0.00 C
 ATOM 34 C6 DA A 2 -0.001 3.407 0.449 1.00 0.00 C
 ATOM 35 N6 DA A 2 -0.531 4.379 1.237 1.00 0.00 N
 ATOM 36 N1 DA A 2 0.899 3.740 -0.487 1.00 0.00 N
 ATOM 37 C2 DA A 2 1.408 2.768 -1.260 1.00 0.00 C
 ATOM 38 N3 DA A 2 1.141 1.466 -1.241 1.00 0.00 N
 ATOM 39 C4 DA A 2 0.266 1.161 -0.278 1.00 0.00 C
 ATOM 40 P DC A 3 3.409 -3.364 -0.650 1.00 0.00 P
 ATOM 41 OP1 DC A 3 4.319 -3.958 -1.676 1.00 0.00 O
 ATOM 42 OP2 DC A 3 3.537 -3.641 0.814 1.00 0.00 O
 ATOM 43 O5' DC A 3 3.438 -1.720 -0.789 1.00 0.00 O
 ATOM 44 C5' DC A 3 3.834 -1.096 -2.005 1.00 0.00 C
 ATOM 45 C4' DC A 3 5.033 -0.201 -1.745 1.00 0.00 C
 ATOM 46 O4' DC A 3 4.639 0.926 -0.959 1.00 0.00 O
 ATOM 47 C3' DC A 3 6.157 -0.889 -0.964 1.00 0.00 C
 ATOM 48 O3' DC A 3 7.373 -0.307 -1.407 1.00 0.00 O
 ATOM 49 C2' DC A 3 5.847 -0.503 0.483 1.00 0.00 C
 ATOM 50 C1' DC A 3 5.250 0.898 0.322 1.00 0.00 C
 ATOM 51 N1 DC A 3 4.240 1.260 1.315 1.00 0.00 N
 ATOM 52 C2 DC A 3 4.273 2.550 1.903 1.00 0.00 C
 ATOM 53 O2 DC A 3 5.255 3.266 1.731 1.00 0.00 O
 ATOM 54 N3 DC A 3 3.187 2.946 2.631 1.00 0.00 N
 ATOM 55 C4 DC A 3 2.163 2.128 2.803 1.00 0.00 C
 ATOM 56 N4 DC A 3 1.097 2.578 3.505 1.00 0.00 N
 ATOM 57 C5 DC A 3 2.135 0.792 2.285 1.00 0.00 C
 ATOM 58 C6 DC A 3 3.173 0.433 1.499 1.00 0.00 C
 END

COMPND AAC / M06-2X / PO- MODEL / SINGLY IONIZED

ATOM 1 O5' DA A 1 -7.282 0.442 -0.256 1.00 0.00 O
 ATOM 2 C5' DA A 1 -7.004 -0.491 -1.283 1.00 0.00 C
 ATOM 3 C4' DA A 1 -5.549 -0.901 -1.192 1.00 0.00 C
 ATOM 4 O4' DA A 1 -4.712 0.211 -1.548 1.00 0.00 O
 ATOM 5 C3' DA A 1 -5.099 -1.324 0.219 1.00 0.00 C
 ATOM 6 O3' DA A 1 -4.098 -2.321 0.037 1.00 0.00 O
 ATOM 7 C2' DA A 1 -4.475 -0.036 0.748 1.00 0.00 C
 ATOM 8 C1' DA A 1 -3.753 0.394 -0.518 1.00 0.00 C
 ATOM 9 N9 DA A 1 -3.279 1.760 -0.529 1.00 0.00 N
 ATOM 10 C8 DA A 1 -3.599 2.806 0.307 1.00 0.00 C
 ATOM 11 N7 DA A 1 -2.922 3.902 0.066 1.00 0.00 N
 ATOM 12 C5 DA A 1 -2.108 3.561 -0.991 1.00 0.00 C
 ATOM 13 C6 DA A 1 -1.122 4.274 -1.710 1.00 0.00 C
 ATOM 14 N6 DA A 1 -0.825 5.552 -1.423 1.00 0.00 N
 ATOM 15 N1 DA A 1 -0.440 3.630 -2.671 1.00 0.00 N
 ATOM 16 C2 DA A 1 -0.735 2.344 -2.916 1.00 0.00 C
 ATOM 17 N3 DA A 1 -1.646 1.569 -2.330 1.00 0.00 N
 ATOM 18 C4 DA A 1 -2.304 2.233 -1.372 1.00 0.00 C
 ATOM 19 P DA A 2 -3.403 -3.058 1.329 1.00 0.00 P
 ATOM 20 OP1 DA A 2 -4.411 -3.905 2.037 1.00 0.00 O
 ATOM 21 OP2 DA A 2 -2.599 -2.068 2.134 1.00 0.00 O
 ATOM 22 O5' DA A 2 -2.379 -4.031 0.504 1.00 0.00 O
 ATOM 23 C5' DA A 2 -2.059 -3.920 -0.875 1.00 0.00 C
 ATOM 24 C4' DA A 2 -0.795 -3.124 -1.126 1.00 0.00 C
 ATOM 25 O4' DA A 2 -1.037 -1.720 -0.905 1.00 0.00 O
 ATOM 26 C3' DA A 2 0.405 -3.511 -0.225 1.00 0.00 C
 ATOM 27 O3' DA A 2 1.541 -3.644 -1.082 1.00 0.00 O
 ATOM 28 C2' DA A 2 0.481 -2.321 0.723 1.00 0.00 C
 ATOM 29 C1' DA A 2 0.104 -2.125 -0.244 1.00 0.00 C
 ATOM 30 N9 DA A 2 -0.204 0.065 0.376 1.00 0.00 N
 ATOM 31 C8 DA A 2 -1.089 0.328 1.362 1.00 0.00 C
 ATOM 32 N7 DA A 2 -1.091 1.620 1.755 1.00 0.00 N
 ATOM 33 C5 DA A 2 -0.145 2.189 1.002 1.00 0.00 C

ATOM 34 C6 DA A 2 0.373 3.525 0.909 1.00 0.00 C
 ATOM 35 N6 DA A 2 -0.157 4.517 1.588 1.00 0.00 N
 ATOM 36 N1 DA A 2 1.411 3.769 0.071 1.00 0.00 N
 ATOM 37 C2 DA A 2 1.879 2.779 -0.655 1.00 0.00 C
 ATOM 38 N3 DA A 2 1.441 1.477 -0.711 1.00 0.00 N
 ATOM 39 C4 DA A 2 0.465 1.248 0.120 1.00 0.00 C
 ATOM 40 P DC A 3 3.091 -3.314 -0.637 1.00 0.00 P
 ATOM 41 OP1 DC A 3 3.954 -3.928 -1.690 1.00 0.00 O
 ATOM 42 OP2 DC A 3 3.295 -3.559 0.820 1.00 0.00 O
 ATOM 43 O5' DC A 3 3.113 -1.674 -0.790 1.00 0.00 O
 ATOM 44 C5' DC A 3 3.443 -1.068 -2.034 1.00 0.00 C
 ATOM 45 C4' DC A 3 4.712 -0.254 -1.874 1.00 0.00 C
 ATOM 46 O4' DC A 3 4.509 0.832 -0.969 1.00 0.00 O
 ATOM 47 C3' DC A 3 5.887 -1.039 -1.294 1.00 0.00 C
 ATOM 48 O3' DC A 3 7.066 -0.409 -1.770 1.00 0.00 O
 ATOM 49 C2' DC A 3 5.714 -0.846 0.217 1.00 0.00 C
 ATOM 50 C1' DC A 3 5.048 0.534 0.306 1.00 0.00 C
 ATOM 51 N1 DC A 3 4.002 0.687 1.332 1.00 0.00 N
 ATOM 52 C2 DC A 3 3.675 2.032 1.660 1.00 0.00 C
 ATOM 53 O2 DC A 3 4.292 2.940 1.110 1.00 0.00 O
 ATOM 54 N3 DC A 3 2.678 2.260 2.561 1.00 0.00 N
 ATOM 55 C4 DC A 3 2.047 1.238 3.128 1.00 0.00 C
 ATOM 56 N4 DC A 3 1.113 1.523 4.050 1.00 0.00 N
 ATOM 57 C5 DC A 3 2.317 -0.123 2.783 1.00 0.00 C
 ATOM 58 C6 DC A 3 3.298 -0.342 1.870 1.00 0.00 C
 END

COMPND CCA / M06-2X / PO- MODEL / NEUTRAL
 ATOM 1 O5' DC A 1 -7.342 -0.870 0.004 1.00 0.00 O
 ATOM 2 C5' DC A 1 -7.024 -1.924 -0.888 1.00 0.00 C
 ATOM 3 C4' DC A 1 -5.516 -2.029 -1.013 1.00 0.00 C
 ATOM 4 O4' DC A 1 -5.005 -0.856 -1.652 1.00 0.00 O
 ATOM 5 C3' DC A 1 -4.772 -2.129 0.326 1.00 0.00 C
 ATOM 6 O3' DC A 1 -3.577 -2.871 0.082 1.00 0.00 O
 ATOM 7 C2' DC A 1 -4.439 -0.673 0.619 1.00 0.00 C
 ATOM 8 C1' DC A 1 -4.098 -0.180 -0.784 1.00 0.00 C
 ATOM 9 N1 DC A 1 -4.246 1.252 -0.979 1.00 0.00 N
 ATOM 10 C2 DC A 1 -3.180 1.987 -1.559 1.00 0.00 C
 ATOM 11 O2 DC A 1 -2.197 1.389 -1.982 1.00 0.00 O
 ATOM 12 N3 DC A 1 -3.306 3.345 -1.606 1.00 0.00 N
 ATOM 13 C4 DC A 1 -4.428 3.925 -1.217 1.00 0.00 C
 ATOM 14 N4 DC A 1 -4.473 5.269 -1.236 1.00 0.00 N
 ATOM 15 C5 DC A 1 -5.582 3.190 -0.776 1.00 0.00 C
 ATOM 16 C6 DC A 1 -5.430 1.850 -0.673 1.00 0.00 C
 ATOM 17 P DC A 2 -2.699 -3.370 1.382 1.00 0.00 P
 ATOM 18 OP1 DC A 2 -3.437 -4.460 2.096 1.00 0.00 O
 ATOM 19 OP2 DC A 2 -2.216 -2.184 2.172 1.00 0.00 O
 ATOM 20 O5' DC A 2 -1.449 -4.045 0.576 1.00 0.00 O
 ATOM 21 C5' DC A 2 -1.102 -3.858 -0.789 1.00 0.00 C
 ATOM 22 C4' DC A 2 0.042 -2.878 -0.983 1.00 0.00 C
 ATOM 23 O4' DC A 2 -0.439 -1.531 -0.952 1.00 0.00 O
 ATOM 24 C3' DC A 2 1.145 -2.971 0.096 1.00 0.00 C
 ATOM 25 O3' DC A 2 2.399 -2.899 -0.586 1.00 0.00 O
 ATOM 26 C2' DC A 2 0.886 -1.718 0.926 1.00 0.00 C
 ATOM 27 C1' DC A 2 0.496 -0.763 -0.193 1.00 0.00 C
 ATOM 28 N1 DC A 2 -0.116 0.496 0.175 1.00 0.00 N
 ATOM 29 C2 DC A 2 0.337 1.687 -0.448 1.00 0.00 C
 ATOM 30 Q2 DC A 2 1.240 1.633 -1.277 1.00 0.00 O
 ATOM 31 N3 DC A 2 -0.242 2.863 -0.068 1.00 0.00 N
 ATOM 32 C4 DC A 2 -1.242 2.866 0.787 1.00 0.00 C
 ATOM 33 N4 DC A 2 -1.742 4.076 1.166 1.00 0.00 N
 ATOM 34 C5 DC A 2 -1.770 1.674 1.379 1.00 0.00 C
 ATOM 35 C6 DC A 2 -1.163 0.509 1.037 1.00 0.00 C
 ATOM 36 P DA A 3 3.798 -3.229 0.223 1.00 0.00 P
 ATOM 37 OP1 DA A 3 4.359 -4.512 -0.303 1.00 0.00 O
 ATOM 38 OP2 DA A 3 3.619 -2.974 1.686 1.00 0.00 O
 ATOM 39 O5' DA A 3 4.701 -1.971 -0.336 1.00 0.00 O
 ATOM 40 C5' DA A 3 4.915 -1.854 -1.733 1.00 0.00 C
 ATOM 41 C4' DA A 3 5.583 -0.521 -2.009 1.00 0.00 C
 ATOM 42 O4' DA A 3 4.701 0.545 -1.643 1.00 0.00 O
 ATOM 43 C3' DA A 3 6.877 -0.275 -1.229 1.00 0.00 C
 ATOM 44 O3' DA A 3 7.675 0.573 -2.040 1.00 0.00 O
 ATOM 45 C2' DA A 3 6.380 0.455 0.014 1.00 0.00 C
 ATOM 46 C1' DA A 3 5.217 1.276 -0.545 1.00 0.00 C
 ATOM 47 N9 DA A 3 4.164 1.515 0.422 1.00 0.00 N
 ATOM 48 C8 DA A 3 3.489 0.575 1.169 1.00 0.00 C
 ATOM 49 N7 DA A 3 2.545 1.085 1.921 1.00 0.00 N
 ATOM 50 C5 DA A 3 2.592 2.436 1.648 1.00 0.00 C
 ATOM 51 C6 DA A 3 1.817 3.523 2.090 1.00 0.00 C
 ATOM 52 N6 DA A 3 0.835 3.375 3.018 1.00 0.00 N
 ATOM 53 N1 DA A 3 2.085 4.745 1.608 1.00 0.00 N
 ATOM 54 C2 DA A 3 3.073 4.870 0.709 1.00 0.00 C
 ATOM 55 N3 DA A 3 3.865 3.930 0.202 1.00 0.00 N
 ATOM 56 C4 DA A 3 3.576 2.723 0.706 1.00 0.00 C
 END

COMPND CCA / M06-2X / PO- MODEL / SINGLY IONIZED
 ATOM 1 O5' DC A 1 -7.286 -0.330 -0.186 1.00 0.00 O
 ATOM 2 C5' DC A 1 -7.026 -1.451 -1.014 1.00 0.00 C
 ATOM 3 C4' DC A 1 -5.532 -1.709 -1.042 1.00 0.00 C
 ATOM 4 O4' DC A 1 -4.868 -0.630 -1.706 1.00 0.00 O
 ATOM 5 C3' DC A 1 -4.880 -1.809 0.344 1.00 0.00 C
 ATOM 6 O3' DC A 1 -3.763 -2.684 0.211 1.00 0.00 O
 ATOM 7 C2' DC A 1 -4.411 -0.380 0.582 1.00 0.00 C
 ATOM 8 C1' DC A 1 -3.947 0.003 -0.822 1.00 0.00 C
 ATOM 9 N1 DC A 1 -3.954 1.433 -1.088 1.00 0.00 N
 ATOM 10 C2 DC A 1 -2.796 2.055 -1.621 1.00 0.00 C
 ATOM 11 O2 DC A 1 -1.841 1.365 -1.964 1.00 0.00 O
 ATOM 12 N3 DC A 1 -2.801 3.416 -1.716 1.00 0.00 N
 ATOM 13 C4 DC A 1 -3.893 4.102 -1.422 1.00 0.00 C
 ATOM 14 N4 DC A 1 -3.822 5.443 -1.494 1.00 0.00 N
 ATOM 15 C5 DC A 1 -5.131 3.482 -1.038 1.00 0.00 C
 ATOM 16 C6 DC A 1 -5.100 2.140 -0.881 1.00 0.00 C
 ATOM 17 P DC A 2 -2.985 -3.179 1.574 1.00 0.00 P
 ATOM 18 OP2 DC A 2 -2.325 -2.013 2.262 1.00 0.00 O
 ATOM 19 OP1 DC A 2 -3.889 -4.068 2.370 1.00 0.00 O
 ATOM 20 O5' DC A 2 -1.850 -4.116 0.862 1.00 0.00 O
 ATOM 21 C5' DC A 2 -1.448 -4.051 -0.498 1.00 0.00 C
 ATOM 22 C4' DC A 2 -0.267 -3.134 -0.740 1.00 0.00 C
 ATOM 23 O4' DC A 2 -0.679 -1.763 -0.713 1.00 0.00 O
 ATOM 24 C3' DC A 2 0.894 -3.275 0.286 1.00 0.00 C
 ATOM 25 O3' DC A 2 2.083 -3.532 -0.470 1.00 0.00 O
 ATOM 26 C2' DC A 2 0.853 -1.924 0.990 1.00 0.00 C
 ATOM 27 C1' DC A 2 0.413 -1.037 -0.163 1.00 0.00 C
 ATOM 28 N1 DC A 2 -0.013 0.303 0.190 1.00 0.00 N
 ATOM 29 C2 DC A 2 0.723 1.400 -0.304 1.00 0.00 C
 ATOM 30 Q2 DC A 2 1.742 1.201 -0.965 1.00 0.00 O
 ATOM 31 N3 DC A 2 0.302 2.654 0.030 1.00 0.00 N

ATOM 32 C4 DC A 2 -0.777 2.816 0.780 1.00 0.00 C
 ATOM 33 N4 DC A 2 -1.107 4.085 1.117 1.00 0.00 N
 ATOM 34 C5 DC A 2 -1.524 1.722 1.317 1.00 0.00 C
 ATOM 35 C6 DC A 2 -1.090 0.476 0.995 1.00 0.00 C
 ATOM 36 P DA A 3 3.585 -2.975 -0.136 1.00 0.00 P
 ATOM 37 OP1 DA A 3 4.506 -3.753 -1.019 1.00 0.00 O
 ATOM 38 OP2 DA A 3 3.815 -2.830 1.342 1.00 0.00 O
 ATOM 39 O5' DA A 3 3.484 -1.419 -0.667 1.00 0.00 O
 ATOM 40 C5' DA A 3 3.685 -1.104 -2.038 1.00 0.00 C
 ATOM 41 C4' DA A 3 4.959 -0.296 -2.197 1.00 0.00 C
 ATOM 42 O4' DA A 3 4.840 0.997 -1.587 1.00 0.00 O
 ATOM 43 C3' DA A 3 6.187 -0.917 -1.532 1.00 0.00 C
 ATOM 44 O3' DA A 3 7.316 -0.430 -2.238 1.00 0.00 O
 ATOM 45 C2' DA A 3 6.130 -0.346 -0.106 1.00 0.00 C
 ATOM 46 C1' DA A 3 5.381 0.982 -0.298 1.00 0.00 C
 ATOM 47 N9 DA A 3 4.305 1.202 0.696 1.00 0.00 N
 ATOM 48 C8 DA A 3 3.639 0.337 1.485 1.00 0.00 C
 ATOM 49 N7 DA A 3 2.661 0.910 2.227 1.00 0.00 N
 ATOM 50 C5 DA A 3 2.698 2.190 1.871 1.00 0.00 C
 ATOM 51 C6 DA A 3 1.948 3.340 2.284 1.00 0.00 C
 ATOM 52 N6 DA A 3 1.025 3.264 3.215 1.00 0.00 N
 ATOM 53 N1 DA A 3 2.222 4.539 1.708 1.00 0.00 N
 ATOM 54 C2 DA A 3 3.173 4.590 0.805 1.00 0.00 C
 ATOM 55 N3 DA A 3 3.983 3.582 0.341 1.00 0.00 N
 ATOM 56 C4 DA A 3 3.706 2.434 0.886 1.00 0.00 C
 END

COMPND CAA / M06-2X / PO- MODEL / NEUTRAL
 ATOM 1 O5' DC A 1 -7.289 -0.728 0.077 1.00 0.00 O
 ATOM 2 C5' DC A 1 -7.020 -1.785 -0.828 1.00 0.00 C
 ATOM 3 C4' DC A 1 -5.518 -1.942 -0.971 1.00 0.00 C
 ATOM 4 O4' DC A 1 -4.974 -0.783 -1.609 1.00 0.00 O
 ATOM 5 C3' DC A 1 -4.767 -2.075 0.360 1.00 0.00 C
 ATOM 6 O3' DC A 1 -3.599 -2.857 0.110 1.00 0.00 O
 ATOM 7 C2' DC A 1 -4.379 -0.634 0.658 1.00 0.00 C
 ATOM 8 C1' DC A 1 -4.038 -0.144 -0.746 1.00 0.00 C
 ATOM 9 N1 DC A 1 -4.136 1.294 -0.928 1.00 0.00 N
 ATOM 10 C2 DC A 1 -3.077 1.985 -1.570 1.00 0.00 C
 ATOM 11 O2 DC A 1 -2.156 1.349 -2.070 1.00 0.00 O
 ATOM 12 N3 DC A 1 -3.137 3.348 -1.587 1.00 0.00 N
 ATOM 13 C4 DC A 1 -4.214 3.973 -1.143 1.00 0.00 C
 ATOM 14 N4 DC A 1 -4.196 5.319 -1.140 1.00 0.00 N
 ATOM 15 C5 DC A 1 -5.382 3.284 -0.669 1.00 0.00 C
 ATOM 16 C6 DC A 1 -5.283 1.939 -0.577 1.00 0.00 C
 ATOM 17 P DA A 2 -2.818 -3.471 1.422 1.00 0.00 P
 ATOM 18 OP1 DA A 2 -3.672 -4.525 2.057 1.00 0.00 O
 ATOM 19 OP2 DA A 2 -2.281 -2.356 2.278 1.00 0.00 O
 ATOM 20 O5' DA A 2 -1.567 -4.187 0.655 1.00 0.00 O
 ATOM 21 C5' DA A 2 -1.261 -4.158 -0.731 1.00 0.00 C
 ATOM 22 C4' DA A 2 -0.046 -3.294 -1.024 1.00 0.00 C
 ATOM 23 O4' DA A 2 -0.415 -1.908 -1.056 1.00 0.00 O
 ATOM 24 C3' DA A 2 1.073 -3.416 0.031 1.00 0.00 C
 ATOM 25 O3' DA A 2 2.306 -3.408 -0.687 1.00 0.00 O
 ATOM 26 C2' DA A 2 0.869 -2.150 0.852 1.00 0.00 C
 ATOM 27 C1' DA A 2 0.506 -0.176 -0.256 1.00 0.00 C
 ATOM 28 N9 DA A 2 -0.104 0.058 0.166 1.00 0.00 N
 ATOM 29 C8 DA A 2 -1.057 0.230 1.140 1.00 0.00 C
 ATOM 30 N7 DA A 2 -1.393 1.486 1.325 1.00 0.00 N
 ATOM 31 C5 DA A 2 -0.605 2.180 0.426 1.00 0.00 C
 ATOM 32 C6 DA A 2 -0.476 3.544 0.127 1.00 0.00 C
 ATOM 33 N6 DA A 2 -1.183 4.499 0.794 1.00 0.00 N
 ATOM 34 N1 DA A 2 0.421 3.921 -0.796 1.00 0.00 N
 ATOM 35 C2 DA A 2 1.148 2.970 -1.400 1.00 0.00 C
 ATOM 36 N3 DA A 2 1.105 1.652 -1.226 1.00 0.00 N
 ATOM 37 C4 DA A 2 0.204 1.311 -0.301 1.00 0.00 C
 ATOM 38 P DA A 3 3.756 -3.513 0.092 1.00 0.00 P
 ATOM 39 OP1 DA A 3 4.530 -4.619 -0.551 1.00 0.00 O
 ATOM 40 OP2 DA A 3 3.562 -3.414 1.570 1.00 0.00 O
 ATOM 41 O5' DA A 3 4.403 -2.068 -0.374 1.00 0.00 O
 ATOM 42 C5' DA A 3 4.617 -1.865 -1.762 1.00 0.00 C
 ATOM 43 C4' DA A 3 5.322 -0.539 -1.966 1.00 0.00 C
 ATOM 44 O4' DA A 3 4.479 0.533 -1.531 1.00 0.00 O
 ATOM 45 C3' DA A 3 6.630 -0.383 -1.185 1.00 0.00 C
 ATOM 46 O3' DA A 3 7.456 0.476 -1.955 1.00 0.00 O
 ATOM 47 C2' DA A 3 6.168 0.296 0.099 1.00 0.00 C
 ATOM 48 C1' DA A 3 5.036 1.191 -0.408 1.00 0.00 C
 ATOM 49 N9 DA A 3 3.997 1.422 0.574 1.00 0.00 N
 ATOM 50 C8 DA A 3 3.248 0.466 1.222 1.00 0.00 C
 ATOM 51 N7 DA A 3 2.308 0.963 1.987 1.00 0.00 N
 ATOM 52 C5 DA A 3 2.438 2.330 1.828 1.00 0.00 C
 ATOM 53 C6 DA A 3 1.711 3.426 2.330 1.00 0.00 C
 ATOM 54 N6 DA A 3 0.650 3.271 3.167 1.00 0.00 N
 ATOM 55 N1 DA A 3 2.093 4.664 1.986 1.00 0.00 N
 ATOM 56 C2 DA A 3 3.122 4.803 1.139 1.00 0.00 C
 ATOM 57 N3 DA A 3 3.861 3.859 0.564 1.00 0.00 N
 ATOM 58 C4 DA A 3 3.475 2.635 0.948 1.00 0.00 C
 END

COMPND CAA / M06-2X / PO- MODEL / SINGLY IONIZED
 ATOM 1 O5' DC A 1 -7.072 -0.991 -0.518 1.00 0.00 O
 ATOM 2 C5' DC A 1 -6.518 -2.012 -1.332 1.00 0.00 C
 ATOM 3 C4' DC A 1 -5.009 -1.979 -1.191 1.00 0.00 C
 ATOM 4 O4' DC A 1 -4.504 -0.766 -1.761 1.00 0.00 O
 ATOM 5 C3' DC A 1 -4.512 -1.995 0.261 1.00 0.00 C
 ATOM 6 O3' DC A 1 -3.235 -2.631 0.278 1.00 0.00 O
 ATOM 7 C2' DC A 1 -4.340 -0.515 0.569 1.00 0.00 C
 ATOM 8 C1' DC A 1 -3.827 -0.001 -0.776 1.00 0.00 C
 ATOM 9 N1 DC A 1 -4.100 1.410 -1.024 1.00 0.00 N
 ATOM 10 C2 DC A 1 -3.056 2.281 -1.395 1.00 0.00 C
 ATOM 11 O2 DC A 1 -1.920 1.822 -1.573 1.00 0.00 O
 ATOM 12 N3 DC A 1 -3.336 3.601 -1.529 1.00 0.00 N
 ATOM 13 C4 DC A 1 -4.590 4.028 -1.427 1.00 0.00 C
 ATOM 14 N4 DC A 1 -4.807 5.344 -1.560 1.00 0.00 N
 ATOM 15 C5 DC A 1 -5.696 3.144 -1.197 1.00 0.00 C
 ATOM 16 C6 DC A 1 -5.393 1.842 -0.997 1.00 0.00 C
 ATOM 17 P DA A 2 -2.826 -3.383 1.688 1.00 0.00 P
 ATOM 18 OP1 DA A 2 -3.735 -4.551 1.895 1.00 0.00 O
 ATOM 19 OP2 DA A 2 -2.627 -2.350 2.762 1.00 0.00 O
 ATOM 20 O5' DA A 2 -1.322 -3.852 1.258 1.00 0.00 O
 ATOM 21 C5' DA A 2 -1.056 -4.469 0.008 1.00 0.00 C
 ATOM 22 C4' DA A 2 0.113 -3.757 -0.652 1.00 0.00 C
 ATOM 23 O4' DA A 2 -0.249 -2.384 -0.879 1.00 0.00 O
 ATOM 24 C3' DA A 2 1.408 -3.711 0.182 1.00 0.00 C
 ATOM 25 O3' DA A 2 2.487 -3.695 -0.756 1.00 0.00 O
 ATOM 26 C2' DA A 2 1.243 -2.393 0.924 1.00 0.00 C
 ATOM 27 C1' DA A 2 0.653 -1.546 -0.188 1.00 0.00 C
 ATOM 28 N9 DA A 2 -0.034 -0.355 0.288 1.00 0.00 N
 ATOM 29 C8 DA A 2 -0.966 -0.232 1.262 1.00 0.00 C

ATOM 30 N7 DA A 2 -1.320 1.045 1.511 1.00 0.00 N
 ATOM 31 C5 DA A 2 -0.566 1.758 0.674 1.00 0.00 C
 ATOM 32 C6 DA A 2 -0.477 3.160 0.379 1.00 0.00 C
 ATOM 33 N6 DA A 2 -1.252 4.033 0.978 1.00 0.00 N
 ATOM 34 N1 DA A 2 0.434 3.581 -0.533 1.00 0.00 N
 ATOM 35 C2 DA A 2 1.169 2.675 -1.143 1.00 0.00 C
 ATOM 36 N3 DA A 2 1.136 1.312 -1.018 1.00 0.00 N
 ATOM 37 C4 DA A 2 0.277 0.923 -0.122 1.00 0.00 C
 ATOM 38 P DA A 3 3.955 -2.997 -0.474 1.00 0.00 P
 ATOM 39 OP1 DA A 3 4.890 -3.589 -1.477 1.00 0.00 O
 ATOM 40 OP2 DA A 3 4.247 -2.958 0.996 1.00 0.00 O
 ATOM 41 O5' DA A 3 3.642 -1.431 -0.876 1.00 0.00 O
 ATOM 42 C5' DA A 3 3.796 -0.963 -2.208 1.00 0.00 C
 ATOM 43 C4' DA A 3 4.837 0.141 -2.235 1.00 0.00 C
 ATOM 44 O4' DA A 3 4.382 1.297 -1.525 1.00 0.00 O
 ATOM 45 C3' DA A 3 6.156 -0.247 -1.566 1.00 0.00 C
 ATOM 46 O3' DA A 3 7.170 0.516 -2.198 1.00 0.00 O
 ATOM 47 C2' DA A 3 5.945 0.190 -0.114 1.00 0.00 C
 ATOM 48 C1' DA A 3 4.985 1.379 -0.250 1.00 0.00 C
 ATOM 49 N9 DA A 3 3.958 1.431 0.785 1.00 0.00 N
 ATOM 50 C8 DA A 3 3.317 0.406 1.444 1.00 0.00 C
 ATOM 51 N7 DA A 3 2.340 0.808 2.222 1.00 0.00 N
 ATOM 52 C5 DA A 3 2.332 2.181 2.065 1.00 0.00 C
 ATOM 53 C6 DA A 3 1.550 3.210 2.631 1.00 0.00 C
 ATOM 54 N6 DA A 3 0.528 2.957 3.474 1.00 0.00 N
 ATOM 55 N1 DA A 3 1.795 4.479 2.267 1.00 0.00 N
 ATOM 56 C2 DA A 3 2.774 4.712 1.381 1.00 0.00 C
 ATOM 57 N3 DA A 3 3.583 3.841 0.783 1.00 0.00 N
 ATOM 58 C4 DA A 3 3.322 2.586 1.172 1.00 0.00 C
 END

COMPND ACC / D-B3LYP / H MODEL / NEUTRAL

ATOM 1 O5' DA A 1 -6.945 -1.095 0.101 1.00 0.00 O
 ATOM 2 C5' DA A 1 -6.699 -2.134 -0.844 1.00 0.00 C
 ATOM 3 C4' DA A 1 -5.200 -2.316 -1.042 1.00 0.00 C
 ATOM 4 O4' DA A 1 -4.620 -1.144 -1.642 1.00 0.00 O
 ATOM 5 C3' DA A 1 -4.415 -2.534 0.257 1.00 0.00 C
 ATOM 6 O3' DA A 1 -3.215 -3.281 -0.114 1.00 0.00 O
 ATOM 7 C2' DA A 1 -4.058 -1.117 0.663 1.00 0.00 C
 ATOM 8 C1' DA A 1 -3.786 -0.466 -0.696 1.00 0.00 C
 ATOM 9 N9 DA A 1 -4.048 0.958 -0.713 1.00 0.00 N
 ATOM 10 C8 DA A 1 -5.012 1.667 -0.019 1.00 0.00 C
 ATOM 11 N7 DA A 1 -4.897 2.970 -0.140 1.00 0.00 N
 ATOM 12 C5 DA A 1 -3.806 3.132 -0.976 1.00 0.00 C
 ATOM 13 C6 DA A 1 -3.108 4.268 -1.427 1.00 0.00 C
 ATOM 14 N6 DA A 1 -3.398 5.516 -0.962 1.00 0.00 N
 ATOM 15 N1 DA A 1 -2.076 4.106 -2.275 1.00 0.00 N
 ATOM 16 C2 DA A 1 -1.707 2.854 -2.591 1.00 0.00 C
 ATOM 17 N3 DA A 1 -2.230 1.697 -2.171 1.00 0.00 N
 ATOM 18 C4 DA A 1 -3.275 1.899 -1.360 1.00 0.00 C
 ATOM 19 P DC A 2 -2.125 -3.735 0.975 1.00 0.00 P
 ATOM 20 OP1 DC A 2 -2.664 -5.092 1.645 1.00 0.00 O
 ATOM 21 OP2 DC A 2 -1.781 -2.757 2.038 1.00 0.00 O
 ATOM 22 O5' DC A 2 -0.940 -4.256 0.037 1.00 0.00 O
 ATOM 23 C5' DC A 2 -0.570 -3.698 -1.249 1.00 0.00 C
 ATOM 24 C4' DC A 2 0.375 -2.518 -1.133 1.00 0.00 C
 ATOM 25 O4' DC A 2 -0.350 -1.328 -0.803 1.00 0.00 O
 ATOM 26 C3' DC A 2 1.477 -2.671 -0.047 1.00 0.00 C
 ATOM 27 O3' DC A 2 2.743 -2.417 -0.718 1.00 0.00 O
 ATOM 28 C2' DC A 2 1.144 -1.561 0.946 1.00 0.00 C
 ATOM 29 C1' DC A 2 0.550 -0.526 -0.008 1.00 0.00 C
 ATOM 30 N1 DC A 2 -0.173 0.592 0.548 1.00 0.00 N
 ATOM 31 C2 DC A 2 0.023 1.893 -0.034 1.00 0.00 C
 ATOM 32 O2 DC A 2 0.863 2.033 -0.921 1.00 0.00 O
 ATOM 33 N3 DC A 2 -0.741 2.915 0.439 1.00 0.00 N
 ATOM 34 C4 DC A 2 -1.642 2.708 1.387 1.00 0.00 C
 ATOM 35 N4 DC A 2 -2.357 3.780 1.816 1.00 0.00 N
 ATOM 36 C5 DC A 2 -1.856 1.425 1.995 1.00 0.00 C
 ATOM 37 C6 DC A 2 -1.098 0.398 1.533 1.00 0.00 C
 ATOM 38 P DC A 3 4.146 -2.641 0.033 1.00 0.00 P
 ATOM 39 OP1 DC A 3 4.298 -4.225 0.263 1.00 0.00 O
 ATOM 40 OP2 DC A 3 4.348 -1.932 1.322 1.00 0.00 O
 ATOM 41 O5' DC A 3 5.185 -2.341 -1.142 1.00 0.00 O
 ATOM 42 C5' DC A 3 4.970 -1.402 -2.229 1.00 0.00 C
 ATOM 43 C4' DC A 3 5.214 0.036 -1.828 1.00 0.00 C
 ATOM 44 O4' DC A 3 4.131 0.514 -1.020 1.00 0.00 O
 ATOM 45 C3' DC A 3 6.523 0.296 -1.029 1.00 0.00 C
 ATOM 46 O3' DC A 3 7.294 1.220 -1.795 1.00 0.00 O
 ATOM 47 C2' DC A 3 6.008 0.906 0.283 1.00 0.00 C
 ATOM 48 C1' DC A 3 4.703 1.544 -0.194 1.00 0.00 C
 ATOM 49 N1 DC A 3 3.748 1.946 0.815 1.00 0.00 N
 ATOM 50 C2 DC A 3 3.200 3.270 0.773 1.00 0.00 C
 ATOM 51 O2 DC A 3 3.631 4.076 -0.054 1.00 0.00 O
 ATOM 52 N3 DC A 3 2.227 3.583 1.677 1.00 0.00 N
 ATOM 53 C4 DC A 3 1.794 2.675 2.542 1.00 0.00 C
 ATOM 54 N4 DC A 3 0.799 3.037 3.391 1.00 0.00 N
 ATOM 55 C5 DC A 3 2.347 1.355 2.632 1.00 0.00 C
 ATOM 56 C6 DC A 3 3.316 1.037 1.736 1.00 0.00 C
 END

COMPND ACC / D-B3LYP / H MODEL / SINGLY IONIZED

ATOM 1 O5' DA A 1 -6.571 -0.378 0.321 1.00 0.00 O
 ATOM 2 C5' DA A 1 -6.680 -1.456 -0.609 1.00 0.00 C
 ATOM 3 C4' DA A 1 -5.288 -1.904 -1.030 1.00 0.00 C
 ATOM 4 O4' DA A 1 -4.649 -0.877 -1.821 1.00 0.00 O
 ATOM 5 C3' DA A 1 -4.346 -2.169 0.152 1.00 0.00 C
 ATOM 6 O3' DA A 1 -3.398 -3.197 -0.262 1.00 0.00 O
 ATOM 7 C2' DA A 1 -3.637 -0.839 0.324 1.00 0.00 C
 ATOM 8 C1' DA A 1 -3.563 -0.313 -1.111 1.00 0.00 C
 ATOM 9 N9 DA A 1 -3.631 1.152 -1.149 1.00 0.00 N
 ATOM 10 C8 DA A 1 -4.548 1.958 -0.548 1.00 0.00 C
 ATOM 11 N7 DA A 1 -4.256 3.266 -0.626 1.00 0.00 N
 ATOM 12 C5 DA A 1 -3.093 3.297 -1.303 1.00 0.00 C
 ATOM 13 C6 DA A 1 -2.234 4.366 -1.719 1.00 0.00 C
 ATOM 14 N6 DA A 1 -2.511 5.630 -1.438 1.00 0.00 N
 ATOM 15 N1 DA A 1 -1.115 4.070 -2.424 1.00 0.00 N
 ATOM 16 C2 DA A 1 -0.861 2.800 -2.694 1.00 0.00 C
 ATOM 17 N3 DA A 1 -1.604 1.688 -2.382 1.00 0.00 N
 ATOM 18 C4 DA A 1 -2.666 1.990 -1.679 1.00 0.00 C
 ATOM 19 P DC A 2 -2.333 -3.788 0.795 1.00 0.00 P
 ATOM 20 OP1 DC A 2 -2.924 -5.178 1.335 1.00 0.00 O
 ATOM 21 OP2 DC A 2 -1.986 -2.902 1.934 1.00 0.00 O
 ATOM 22 O5' DC A 2 -1.149 -4.277 -0.166 1.00 0.00 O
 ATOM 23 C5' DC A 2 -0.731 -3.609 -1.381 1.00 0.00 C
 ATOM 24 C4' DC A 2 0.249 -2.481 -1.132 1.00 0.00 C
 ATOM 25 O4' DC A 2 -0.438 -1.320 -0.647 1.00 0.00 O
 ATOM 26 C3' DC A 2 1.361 -2.790 -0.091 1.00 0.00 C
 ATOM 27 O3' DC A 2 2.612 -2.476 -0.756 1.00 0.00 O
 ATOM 28 C2' DC A 2 1.063 -1.811 1.045 1.00 0.00 C

ATOM 29 C1' DC A 2 0.489 -0.652 0.238 1.00 0.00 C
 ATOM 30 N1 DC A 2 -0.219 0.407 0.925 1.00 0.00 N
 ATOM 31 C2 DC A 2 -0.206 1.707 0.331 1.00 0.00 C
 ATOM 32 O2 DC A 2 0.567 1.930 -0.606 1.00 0.00 O
 ATOM 33 N3 DC A 2 -1.076 2.642 0.798 1.00 0.00 N
 ATOM 34 C4 DC A 2 -1.851 2.371 1.855 1.00 0.00 C
 ATOM 35 N4 DC A 2 -2.718 3.317 2.244 1.00 0.00 N
 ATOM 36 C5 DC A 2 -1.779 1.127 2.560 1.00 0.00 C
 ATOM 37 C6 DC A 2 -0.979 0.165 2.033 1.00 0.00 C
 ATOM 38 P DC A 3 4.051 -2.696 -0.078 1.00 0.00 P
 ATOM 39 OP1 DC A 3 4.272 -4.284 0.040 1.00 0.00 O
 ATOM 40 OP2 DC A 3 4.283 -2.065 1.246 1.00 0.00 O
 ATOM 41 O5' DC A 3 5.020 -2.269 -1.273 1.00 0.00 O
 ATOM 42 C5' DC A 3 4.685 -1.320 -2.319 1.00 0.00 C
 ATOM 43 C4' DC A 3 4.851 0.118 -1.880 1.00 0.00 C
 ATOM 44 O4' DC A 3 3.800 0.487 -0.975 1.00 0.00 O
 ATOM 45 C3' DC A 3 6.195 0.438 -1.167 1.00 0.00 C
 ATOM 46 O3' DC A 3 6.832 1.460 -1.931 1.00 0.00 O
 ATOM 47 C2' DC A 3 5.739 0.935 0.212 1.00 0.00 C
 ATOM 48 C1' DC A 3 4.367 1.510 -0.138 1.00 0.00 C
 ATOM 49 N1 DC A 3 3.471 1.801 0.959 1.00 0.00 N
 ATOM 50 C2 DC A 3 2.878 3.105 1.049 1.00 0.00 C
 ATOM 51 O2 DC A 3 3.185 3.968 0.225 1.00 0.00 O
 ATOM 52 N3 DC A 3 2.005 3.331 2.073 1.00 0.00 N
 ATOM 53 C4 DC A 3 1.722 2.370 2.944 1.00 0.00 C
 ATOM 54 N4 DC A 3 0.847 2.660 3.932 1.00 0.00 N
 ATOM 55 C5 DC A 3 2.307 1.061 2.885 1.00 0.00 C
 ATOM 56 C6 DC A 3 3.175 0.827 1.868 1.00 0.00 C
 END

COMPND AAC / D-B3LYP / H MODEL / NEUTRAL
 ATOM 1 O5' DA A 1 0.000 0.000 0.000 1.00 0.00 O
 ATOM 2 C5' DA A 1 0.000 0.000 1.426 1.00 0.00 C
 ATOM 3 C4' DA A 1 1.435 0.000 1.936 1.00 0.00 C
 ATOM 4 O4' DA A 1 2.072 -1.260 1.660 1.00 0.00 O
 ATOM 5 C3' DA A 1 2.312 1.071 1.276 1.00 0.00 C
 ATOM 6 O3' DA A 1 3.319 1.469 2.260 1.00 0.00 O
 ATOM 7 C2' DA A 1 2.961 0.320 0.128 1.00 0.00 C
 ATOM 8 C1' DA A 1 3.105 -1.097 0.687 1.00 0.00 C
 ATOM 9 N9 DA A 1 2.999 -2.118 -0.337 1.00 0.00 N
 ATOM 10 C8 DA A 1 2.041 -2.225 -1.328 1.00 0.00 C
 ATOM 11 N7 DA A 1 2.246 -3.229 -2.151 1.00 0.00 N
 ATOM 12 C5 DA A 1 3.402 -3.823 -1.672 1.00 0.00 C
 ATOM 13 C6 DA A 1 4.174 -4.911 -2.124 1.00 0.00 C
 ATOM 14 N6 DA A 1 3.832 -5.633 -3.221 1.00 0.00 N
 ATOM 15 N1 DA A 1 5.280 -5.254 -1.432 1.00 0.00 N
 ATOM 16 C2 DA A 1 5.599 -4.531 -0.343 1.00 0.00 C
 ATOM 17 N3 DA A 1 4.969 -3.474 0.175 1.00 0.00 N
 ATOM 18 C4 DA A 1 3.880 -3.158 -0.539 1.00 0.00 C
 ATOM 19 P DA A 2 4.296 2.709 1.956 1.00 0.00 P
 ATOM 20 OP1 DA A 2 3.732 3.967 2.780 1.00 0.00 O
 ATOM 21 OP2 DA A 2 4.462 3.072 0.527 1.00 0.00 O
 ATOM 22 O5' DA A 2 5.612 2.329 2.789 1.00 0.00 O
 ATOM 23 C5' DA A 2 6.084 0.985 3.047 1.00 0.00 C
 ATOM 24 C4' DA A 2 6.957 0.447 1.931 1.00 0.00 C
 ATOM 25 O4' DA A 2 6.140 0.078 0.811 1.00 0.00 O
 ATOM 26 C3' DA A 2 8.014 1.444 1.387 1.00 0.00 C
 ATOM 27 O3' DA A 2 9.260 0.702 1.334 1.00 0.00 O
 ATOM 28 C2' DA A 2 7.503 1.758 -0.016 1.00 0.00 C
 ATOM 29 C1' DA A 2 6.884 0.411 -0.375 1.00 0.00 C
 ATOM 30 N9 DA A 2 6.015 0.395 -1.515 1.00 0.00 N
 ATOM 31 C8 DA A 2 5.246 1.419 -2.042 1.00 0.00 C
 ATOM 32 N7 DA A 2 4.587 1.073 -3.123 1.00 0.00 N
 ATOM 33 C5 DA A 2 4.942 -0.252 -3.326 1.00 0.00 C
 ATOM 34 C6 DA A 2 4.664 -1.166 -4.361 1.00 0.00 C
 ATOM 35 N6 DA A 2 3.935 -0.810 -5.452 1.00 0.00 N
 ATOM 36 N1 DA A 2 5.201 -2.397 -4.303 1.00 0.00 N
 ATOM 37 C2 DA A 2 6.041 -2.677 -3.292 1.00 0.00 C
 ATOM 38 N3 DA A 2 6.423 -1.890 -2.282 1.00 0.00 N
 ATOM 39 C4 DA A 2 5.833 -0.691 -2.348 1.00 0.00 C
 ATOM 40 P DC A 3 10.644 1.342 0.843 1.00 0.00 P
 ATOM 41 OP1 DC A 3 11.057 2.350 2.029 1.00 0.00 O
 ATOM 42 OP2 DC A 3 10.671 1.990 -0.497 1.00 0.00 O
 ATOM 43 O5' DC A 3 11.641 0.119 1.073 1.00 0.00 O
 ATOM 44 C5' DC A 3 11.269 -1.282 0.994 1.00 0.00 C
 ATOM 45 C4' DC A 3 11.238 -1.803 -0.426 1.00 0.00 C
 ATOM 46 O4' DC A 3 10.082 -1.310 -1.121 1.00 0.00 O
 ATOM 47 C3' DC A 3 12.478 -1.431 -1.289 1.00 0.00 C
 ATOM 48 O3' DC A 3 13.064 -2.660 -1.714 1.00 0.00 O
 ATOM 49 C2' DC A 3 11.859 -0.645 -2.452 1.00 0.00 C
 ATOM 50 C1' DC A 3 10.469 -1.275 -2.506 1.00 0.00 C
 ATOM 51 N1 DA A 3 9.465 -0.582 -3.285 1.00 0.00 N
 ATOM 52 C2 DC A 3 8.801 -1.280 -4.346 1.00 0.00 C
 ATOM 53 Q2 DC A 3 9.100 -2.453 -4.580 1.00 0.00 O
 ATOM 54 N3 DC A 3 7.861 -0.598 -5.059 1.00 0.00 N
 ATOM 55 C4 DC A 3 7.559 0.658 -4.754 1.00 0.00 C
 ATOM 56 N4 DC A 3 6.602 1.273 -5.500 1.00 0.00 N
 ATOM 57 C5 DC A 3 8.211 1.384 -3.703 1.00 0.00 C
 ATOM 58 C6 DC A 3 9.160 0.718 -2.996 1.00 0.00 C
 END

COMPND AAC / D-B3LYP / H MODEL / SINGLY IONIZED
 ATOM 1 O5' DA A 1 -6.362 -1.479 -0.024 1.00 0.00 O
 ATOM 2 C5' DA A 1 -6.067 -2.473 -1.001 1.00 0.00 C
 ATOM 3 C4' DA A 1 -4.559 -2.646 -1.129 1.00 0.00 C
 ATOM 4 O4' DA A 1 -3.935 -1.434 -1.589 1.00 0.00 O
 ATOM 5 C3' DA A 1 -3.833 -2.975 0.177 1.00 0.00 C
 ATOM 6 O3' DA A 1 -2.589 -3.629 -0.219 1.00 0.00 O
 ATOM 7 C2' DA A 1 -3.559 -1.595 0.756 1.00 0.00 C
 ATOM 8 C1' DA A 1 -3.274 -0.779 -0.510 1.00 0.00 C
 ATOM 9 N9 DA A 1 -3.722 0.610 -0.439 1.00 0.00 N
 ATOM 10 C8 DA A 1 -4.718 1.187 0.304 1.00 0.00 C
 ATOM 11 N7 DA A 1 -4.931 2.466 0.015 1.00 0.00 N
 ATOM 12 C5 DA A 1 -4.047 2.730 -0.990 1.00 0.00 C
 ATOM 13 C6 DA A 1 -3.783 3.897 -1.758 1.00 0.00 C
 ATOM 14 N6 DA A 1 -4.424 5.047 -1.539 1.00 0.00 N
 ATOM 15 N1 DA A 1 -2.855 3.827 -2.741 1.00 0.00 N
 ATOM 16 C2 DA A 1 -2.210 2.674 -2.922 1.00 0.00 C
 ATOM 17 N3 DA A 1 -2.355 1.504 -2.255 1.00 0.00 N
 ATOM 18 C4 DA A 1 -3.279 1.596 -1.307 1.00 0.00 C
 ATOM 19 P DA A 2 -1.501 -4.152 0.837 1.00 0.00 P
 ATOM 20 OP1 DA A 2 -2.009 -5.574 1.381 1.00 0.00 O
 ATOM 21 OP2 DA A 2 -1.193 -3.264 1.987 1.00 0.00 O
 ATOM 22 O5' DA A 2 -0.276 -4.535 -0.118 1.00 0.00 O
 ATOM 23 C5' DA A 2 0.008 -3.960 -1.417 1.00 0.00 C
 ATOM 24 C4' DA A 2 0.861 -2.710 -1.326 1.00 0.00 C
 ATOM 25 O4' DA A 2 0.058 -1.604 -0.873 1.00 0.00 O
 ATOM 26 C3' DA A 2 2.056 -2.803 -0.341 1.00 0.00 C

ATOM 27 O3' DA A 2 3.190 -2.244 -1.056 1.00 0.00 O
 ATOM 28 C2' DA A 2 1.628 -1.898 0.809 1.00 0.00 C
 ATOM 29 C1' DA A 2 0.870 -0.835 0.024 1.00 0.00 C
 ATOM 30 N9 DA A 2 0.006 0.037 0.782 1.00 0.00 N
 ATOM 31 C8 DA A 2 -0.730 -0.243 1.903 1.00 0.00 C
 ATOM 32 N7 DA A 2 -1.526 0.752 2.281 1.00 0.00 N
 ATOM 33 C5 DA A 2 -1.290 1.728 1.365 1.00 0.00 C
 ATOM 34 C6 DA A 2 -1.812 3.042 1.197 1.00 0.00 C
 ATOM 35 N6 DA A 2 -2.753 3.527 1.999 1.00 0.00 N
 ATOM 36 N1 DA A 2 -1.346 3.804 0.176 1.00 0.00 N
 ATOM 37 C2 DA A 2 -0.423 3.287 -0.627 1.00 0.00 C
 ATOM 38 N3 DA A 2 0.123 2.041 -0.599 1.00 0.00 N
 ATOM 39 C4 DA A 2 -0.338 1.324 0.407 1.00 0.00 C
 ATOM 40 P DC A 3 4.668 -2.095 -0.445 1.00 0.00 P
 ATOM 41 OP1 DC A 3 5.345 -3.551 -0.496 1.00 0.00 O
 ATOM 42 OP2 DC A 3 4.789 -1.542 0.927 1.00 0.00 O
 ATOM 43 O5' DC A 3 5.403 -1.309 -1.627 1.00 0.00 O
 ATOM 44 C5' DC A 3 4.754 -0.372 -2.526 1.00 0.00 C
 ATOM 45 C4' DC A 3 4.584 1.004 -1.922 1.00 0.00 C
 ATOM 46 O4' DC A 3 3.570 0.984 -0.906 1.00 0.00 O
 ATOM 47 C3' DC A 3 5.860 1.607 -1.270 1.00 0.00 C
 ATOM 48 O3' DC A 3 6.140 2.819 -1.966 1.00 0.00 O
 ATOM 49 C2' DC A 3 5.423 1.851 0.183 1.00 0.00 C
 ATOM 50 C1' DC A 3 3.920 2.048 -0.005 1.00 0.00 C
 ATOM 51 N1 DC A 3 3.086 1.981 1.178 1.00 0.00 N
 ATOM 52 C2 DC A 3 2.146 3.031 1.429 1.00 0.00 C
 ATOM 53 O2 DC A 3 2.099 3.998 0.663 1.00 0.00 O
 ATOM 54 N3 DC A 3 1.321 2.905 2.506 1.00 0.00 N
 ATOM 55 C4 DC A 3 1.433 1.858 3.325 1.00 0.00 C
 ATOM 56 N4 DC A 3 0.604 1.808 4.384 1.00 0.00 N
 ATOM 57 C5 DC A 3 2.386 0.808 3.116 1.00 0.00 C
 ATOM 58 C6 DC A 3 3.184 0.912 2.023 1.00 0.00 C
 END

COMPND CCA / D-B3LYP / H MODEL / NEUTRAL

ATOM 1 O5' DC A 1 -6.945 -1.204 0.390 1.00 0.00 O
 ATOM 2 C5' DC A 1 -6.738 -2.253 -0.555 1.00 0.00 C
 ATOM 3 C4' DC A 1 -5.255 -2.356 -0.884 1.00 0.00 C
 ATOM 4 O4' DC A 1 -4.829 -1.211 -1.639 1.00 0.00 O
 ATOM 5 C3' DC A 1 -4.356 -2.400 0.360 1.00 0.00 C
 ATOM 6 O3' DC A 1 -3.188 -3.207 0.012 1.00 0.00 O
 ATOM 7 C2' DC A 1 -3.962 -0.946 0.547 1.00 0.00 C
 ATOM 8 C1' DC A 1 -3.857 -0.457 -0.901 1.00 0.00 C
 ATOM 9 N1 DC A 1 -4.113 0.967 -1.076 1.00 0.00 N
 ATOM 10 C2 DC A 1 -3.258 1.734 -1.931 1.00 0.00 C
 ATOM 11 O2 DC A 1 -2.380 1.162 -2.578 1.00 0.00 O
 ATOM 12 N3 DC A 1 -3.460 3.083 -1.972 1.00 0.00 N
 ATOM 13 C4 DC A 1 -4.469 3.631 -1.312 1.00 0.00 C
 ATOM 14 N4 DC A 1 -4.532 4.986 -1.278 1.00 0.00 N
 ATOM 15 C5 DC A 1 -5.458 2.863 -0.610 1.00 0.00 C
 ATOM 16 C6 DC A 1 -5.233 1.528 -0.526 1.00 0.00 C
 ATOM 17 P DC A 2 -2.130 -3.616 1.150 1.00 0.00 P
 ATOM 18 OP1 DC A 2 -2.734 -4.896 1.912 1.00 0.00 O
 ATOM 19 OP2 DC A 2 -1.775 -2.570 2.142 1.00 0.00 O
 ATOM 20 O5' DC A 2 -0.949 -4.257 0.283 1.00 0.00 O
 ATOM 21 C5' DC A 2 -0.543 -3.832 -1.043 1.00 0.00 C
 ATOM 22 C4' DC A 2 0.416 -2.658 -1.019 1.00 0.00 C
 ATOM 23 O4' DC A 2 -0.304 -1.442 -0.796 1.00 0.00 O
 ATOM 24 C3' DC A 2 1.505 -2.728 0.088 1.00 0.00 C
 ATOM 25 O3' DC A 2 2.780 -2.527 -0.585 1.00 0.00 O
 ATOM 26 C2' DC A 2 1.162 -1.547 0.993 1.00 0.00 C
 ATOM 27 C1' DC A 2 0.584 -0.585 -0.044 1.00 0.00 C
 ATOM 28 N1 DC A 2 -0.153 0.571 0.404 1.00 0.00 N
 ATOM 29 C2 DC A 2 -0.092 1.759 -0.408 1.00 0.00 C
 ATOM 30 O2 DC A 2 0.693 1.798 -1.353 1.00 0.00 O
 ATOM 31 N3 DC A 2 -0.913 2.794 -0.076 1.00 0.00 N
 ATOM 32 C4 DC A 2 -1.724 2.707 0.966 1.00 0.00 C
 ATOM 33 N4 DC A 2 -2.512 3.779 1.242 1.00 0.00 N
 ATOM 34 C5 DC A 2 -1.770 1.561 1.827 1.00 0.00 C
 ATOM 35 C6 DC A 2 -0.979 0.511 1.488 1.00 0.00 C
 ATOM 36 P DA A 3 4.173 -2.694 0.197 1.00 0.00 P
 ATOM 37 OP1 DA A 3 4.320 -4.258 0.544 1.00 0.00 O
 ATOM 38 OP2 DA A 3 4.362 -1.894 1.433 1.00 0.00 O
 ATOM 39 O5' DA A 3 5.230 -2.476 -0.982 1.00 0.00 O
 ATOM 40 C5' DA A 3 5.014 -1.669 -2.168 1.00 0.00 C
 ATOM 41 C4' DA A 3 5.243 -0.193 -1.924 1.00 0.00 C
 ATOM 42 O4' DA A 3 4.151 0.347 -1.164 1.00 0.00 O
 ATOM 43 C3' DA A 3 6.545 0.157 -1.152 1.00 0.00 C
 ATOM 44 O3' DA A 3 7.263 1.086 -1.962 1.00 0.00 O
 ATOM 45 C2' DA A 3 6.017 0.799 0.138 1.00 0.00 C
 ATOM 46 C1' DA A 3 4.706 1.399 -0.362 1.00 0.00 C
 ATOM 47 N9 DA A 3 3.752 1.790 0.639 1.00 0.00 N
 ATOM 48 C8 DA A 3 3.307 1.048 1.718 1.00 0.00 C
 ATOM 49 N7 DA A 3 2.344 1.631 2.394 1.00 0.00 N
 ATOM 50 C5 DA A 3 2.143 2.831 1.729 1.00 0.00 C
 ATOM 51 C6 DA A 3 1.242 3.895 1.928 1.00 0.00 C
 ATOM 52 N6 DA A 3 0.335 3.898 2.945 1.00 0.00 N
 ATOM 53 N1 DA A 3 1.299 4.956 1.100 1.00 0.00 N
 ATOM 54 C2 DA A 3 2.196 4.935 0.100 1.00 0.00 C
 ATOM 55 N3 DA A 3 3.080 3.981 -0.215 1.00 0.00 N
 ATOM 56 C4 DA A 3 3.003 2.950 0.634 1.00 0.00 C
 END

COMPND CCA / D-B3LYP / H MODEL / SINGLY IONIZED

ATOM 1 O5' DC A 1 -6.867 -1.394 0.467 1.00 0.00 O
 ATOM 2 C5' DC A 1 -6.637 -2.474 -0.436 1.00 0.00 C
 ATOM 3 C4' DC A 1 -5.159 -2.535 -0.793 1.00 0.00 C
 ATOM 4 O4' DC A 1 -4.794 -1.415 -1.617 1.00 0.00 O
 ATOM 5 C3' DC A 1 -4.233 -2.479 0.430 1.00 0.00 C
 ATOM 6 O3' DC A 1 -3.044 -3.257 0.090 1.00 0.00 O
 ATOM 7 C2' DC A 1 -3.891 -1.002 0.529 1.00 0.00 C
 ATOM 8 C1' DC A 1 -3.839 -0.585 -0.944 1.00 0.00 C
 ATOM 9 N1 DC A 1 -4.169 0.817 -1.173 1.00 0.00 N
 ATOM 10 C2 DC A 1 -3.320 1.620 -1.999 1.00 0.00 C
 ATOM 11 O2 DC A 1 -2.365 1.097 -2.576 1.00 0.00 O
 ATOM 12 N3 DC A 1 -3.606 2.951 -2.083 1.00 0.00 N
 ATOM 13 C4 DC A 1 -4.679 3.447 -1.482 1.00 0.00 C
 ATOM 14 N4 DC A 1 -4.823 4.796 -1.476 1.00 0.00 N
 ATOM 15 C5 DC A 1 -5.648 2.633 -0.805 1.00 0.00 C
 ATOM 16 C6 DC A 1 -5.342 1.318 -0.678 1.00 0.00 C
 ATOM 17 P DC A 2 -1.906 -3.548 1.185 1.00 0.00 P
 ATOM 18 OP1 DC A 2 -2.361 -4.842 2.018 1.00 0.00 O
 ATOM 19 OP2 DC A 2 -1.581 -2.442 2.120 1.00 0.00 O
 ATOM 20 O5' DC A 2 -0.729 -4.124 0.267 1.00 0.00 O
 ATOM 21 C5' DC A 2 -0.408 -3.664 -1.069 1.00 0.00 C
 ATOM 22 C4' DC A 2 0.517 -2.463 -1.067 1.00 0.00 C
 ATOM 23 O4' DC A 2 -0.228 -1.266 -0.793 1.00 0.00 O
 ATOM 24 C3' DC A 2 1.648 -2.520 -0.006 1.00 0.00 C

ATOM 25 O3' DC A 2 2.890 -2.297 -0.722 1.00 0.00 O
 ATOM 26 C2' DC A 2 1.320 -1.353 0.927 1.00 0.00 C
 ATOM 27 C1' DC A 2 0.661 -0.403 -0.070 1.00 0.00 C
 ATOM 28 N1 DC A 2 -0.125 0.715 0.440 1.00 0.00 N
 ATOM 29 C2 DC A 2 -0.221 1.907 -0.390 1.00 0.00 C
 ATOM 30 O2 DC A 2 0.584 2.045 -1.306 1.00 0.00 O
 ATOM 31 N3 DC A 2 -1.172 2.823 -0.088 1.00 0.00 N
 ATOM 32 C4 DC A 2 -1.975 2.658 0.956 1.00 0.00 C
 ATOM 33 N4 DC A 2 -2.916 3.587 1.187 1.00 0.00 N
 ATOM 34 C5 DC A 2 -1.834 1.544 1.855 1.00 0.00 C
 ATOM 35 C6 DC A 2 -0.934 0.568 1.512 1.00 0.00 C
 ATOM 36 P DA A 3 4.311 -2.507 -0.001 1.00 0.00 P
 ATOM 37 OP1 DA A 3 4.476 -4.085 0.246 1.00 0.00 O
 ATOM 38 OP2 DA A 3 4.528 -1.782 1.276 1.00 0.00 O
 ATOM 39 O5' DA A 3 5.324 -2.206 -1.198 1.00 0.00 O
 ATOM 40 C5' DA A 3 5.050 -1.337 -2.327 1.00 0.00 C
 ATOM 41 C4' DA A 3 5.239 0.126 -1.991 1.00 0.00 C
 ATOM 42 O4' DA A 3 4.167 0.561 -1.132 1.00 0.00 O
 ATOM 43 C3' DA A 3 6.565 0.472 -1.261 1.00 0.00 C
 ATOM 44 O3' DA A 3 7.181 1.517 -2.008 1.00 0.00 O
 ATOM 45 C2' DA A 3 6.088 0.952 0.119 1.00 0.00 C
 ATOM 46 C1' DA A 3 4.732 1.537 -0.256 1.00 0.00 C
 ATOM 47 N9 DA A 3 3.800 1.774 0.824 1.00 0.00 N
 ATOM 48 C8 DA A 3 3.566 1.019 1.943 1.00 0.00 C
 ATOM 49 N7 DA A 3 2.533 1.439 2.668 1.00 0.00 N
 ATOM 50 C5 DA A 3 2.070 2.521 1.983 1.00 0.00 C
 ATOM 51 C6 DA A 3 0.994 3.423 2.196 1.00 0.00 C
 ATOM 52 N6 DA A 3 0.120 3.261 3.192 1.00 0.00 N
 ATOM 53 N1 DA A 3 0.810 4.455 1.333 1.00 0.00 N
 ATOM 54 C2 DA A 3 1.623 4.551 0.289 1.00 0.00 C
 ATOM 55 N3 DA A 3 2.664 3.742 -0.045 1.00 0.00 N
 ATOM 56 C4 DA A 3 2.840 2.767 0.824 1.00 0.00 C
 END

COMPND CAA / D-B3LYP / H MODEL / NEUTRAL

ATOM 1 O5' DC A 1 -6.236 -0.662 0.768 1.00 0.00 O
 ATOM 2 C5' DC A 1 -6.430 -1.860 0.018 1.00 0.00 C
 ATOM 3 C4' DC A 1 -5.099 -2.314 -0.565 1.00 0.00 C
 ATOM 4 O4' DC A 1 -4.686 -1.441 -1.628 1.00 0.00 O
 ATOM 5 C3' DC A 1 -3.958 -2.303 0.461 1.00 0.00 C
 ATOM 6 O3' DC A 1 -3.054 -3.398 0.102 1.00 0.00 O
 ATOM 7 C2' DC A 1 -3.283 -0.963 0.223 1.00 0.00 C
 ATOM 8 C1' DC A 1 -3.473 -0.756 -1.282 1.00 0.00 C
 ATOM 9 N1 DC A 1 -3.586 0.640 -1.691 1.00 0.00 N
 ATOM 10 C2 DC A 1 -2.790 1.130 -2.772 1.00 0.00 C
 ATOM 11 O2 DC A 1 -2.000 0.371 -3.339 1.00 0.00 O
 ATOM 12 N3 DC A 1 -2.939 2.441 -3.122 1.00 0.00 N
 ATOM 13 C4 DC A 1 -3.842 3.203 -2.512 1.00 0.00 C
 ATOM 14 N4 DC A 1 -3.932 4.494 -2.896 1.00 0.00 N
 ATOM 15 C5 DC A 1 -4.729 2.708 -1.499 1.00 0.00 C
 ATOM 16 C6 DC A 1 -4.562 1.416 -1.127 1.00 0.00 C
 ATOM 17 P DA A 2 -1.953 -3.890 1.164 1.00 0.00 P
 ATOM 18 OP1 DA A 2 -2.497 -5.256 1.813 1.00 0.00 O
 ATOM 19 OP2 DA A 2 -1.613 -2.918 2.231 1.00 0.00 O
 ATOM 20 O5' DA A 2 -0.767 -4.425 0.227 1.00 0.00 O
 ATOM 21 C5' DA A 2 -0.428 -3.914 -1.084 1.00 0.00 C
 ATOM 22 C4' DA A 2 0.532 -2.742 -1.022 1.00 0.00 C
 ATOM 23 O4' DA A 2 -0.176 -1.563 -0.613 1.00 0.00 O
 ATOM 24 C3' DA A 2 1.702 -2.911 -0.021 1.00 0.00 C
 ATOM 25 O3' DA A 2 2.894 -2.491 -0.738 1.00 0.00 O
 ATOM 26 C2' DA A 2 1.351 -1.933 1.097 1.00 0.00 C
 ATOM 27 C1' DA A 2 0.672 -0.833 0.291 1.00 0.00 C
 ATOM 28 N9 DA A 2 -0.141 0.087 1.037 1.00 0.00 N
 ATOM 29 C8 DA A 2 -0.684 -0.046 2.300 1.00 0.00 C
 ATOM 30 N7 DA A 2 -1.473 0.946 2.639 1.00 0.00 N
 ATOM 31 C5 DA A 2 -1.460 1.781 1.532 1.00 0.00 C
 ATOM 32 C6 DA A 2 -2.058 3.027 1.267 1.00 0.00 C
 ATOM 33 N6 DA A 2 -2.806 3.678 2.199 1.00 0.00 N
 ATOM 34 N1 DA A 2 -1.826 3.618 0.081 1.00 0.00 N
 ATOM 35 C2 DA A 2 -1.002 3.007 -0.786 1.00 0.00 C
 ATOM 36 N3 DA A 2 -0.358 1.842 -0.648 1.00 0.00 N
 ATOM 37 C4 DA A 2 -0.632 1.274 0.531 1.00 0.00 C
 ATOM 38 P DA A 3 4.364 -2.537 -0.095 1.00 0.00 P
 ATOM 39 OP1 DA A 3 4.738 -4.089 0.096 1.00 0.00 O
 ATOM 40 OP2 DA A 3 4.571 -1.818 1.187 1.00 0.00 O
 ATOM 41 O5' DA A 3 5.259 -2.083 -1.339 1.00 0.00 O
 ATOM 42 C5' DA A 3 4.802 -1.255 -2.441 1.00 0.00 C
 ATOM 43 C4' DA A 3 4.837 0.221 -2.111 1.00 0.00 C
 ATOM 44 O4' DA A 3 3.773 0.548 -1.201 1.00 0.00 O
 ATOM 45 C3' DA A 3 6.157 0.714 -1.456 1.00 0.00 C
 ATOM 46 O3' DA A 3 6.621 1.808 -2.245 1.00 0.00 O
 ATOM 47 C2' DA A 3 5.698 1.160 -0.062 1.00 0.00 C
 ATOM 48 C1' DA A 3 4.273 1.607 -0.374 1.00 0.00 C
 ATOM 49 N9 DA A 3 3.394 1.798 0.747 1.00 0.00 N
 ATOM 50 C8 DA A 3 3.163 0.945 1.811 1.00 0.00 C
 ATOM 51 N7 DA A 3 2.260 1.393 2.653 1.00 0.00 N
 ATOM 52 C5 DA A 3 1.876 2.615 2.123 1.00 0.00 C
 ATOM 53 C6 DA A 3 0.930 3.584 2.520 1.00 0.00 C
 ATOM 54 N6 DA A 3 0.158 3.447 3.633 1.00 0.00 N
 ATOM 55 N1 DA A 3 0.797 4.699 1.773 1.00 0.00 N
 ATOM 56 C2 DA A 3 1.547 4.825 0.667 1.00 0.00 C
 ATOM 57 N3 DA A 3 2.447 3.972 0.166 1.00 0.00 N
 ATOM 58 C4 DA A 3 2.567 2.885 0.937 1.00 0.00 C
 END

COMPND CAA / D-B3LYP / H MODEL / SINGLY IONIZED

ATOM 1 O5' DC A 1 6.234 0.592 0.562 1.00 0.00 O
 ATOM 2 C5' DC A 1 6.392 1.838 -0.116 1.00 0.00 C
 ATOM 3 C4' DC A 1 5.038 2.319 -0.617 1.00 0.00 C
 ATOM 4 O4' DC A 1 4.576 1.498 -1.704 1.00 0.00 O
 ATOM 5 C3' DC A 1 3.938 2.261 0.452 1.00 0.00 C
 ATOM 6 O3' DC A 1 3.019 3.362 0.162 1.00 0.00 O
 ATOM 7 C2' DC A 1 3.264 0.926 0.181 1.00 0.00 C
 ATOM 8 C1' DC A 1 3.377 0.803 -1.341 1.00 0.00 C
 ATOM 9 N1 DC A 1 3.459 -0.572 -1.821 1.00 0.00 N
 ATOM 10 C2 DC A 1 2.546 -1.033 -2.821 1.00 0.00 C
 ATOM 11 O2 DC A 1 1.708 -0.258 -3.284 1.00 0.00 O
 ATOM 12 N3 DC A 1 2.641 -2.341 -3.204 1.00 0.00 N
 ATOM 13 C4 DC A 1 3.611 -3.117 -2.724 1.00 0.00 C
 ATOM 14 N4 DC A 1 3.631 -4.404 -3.120 1.00 0.00 N
 ATOM 15 C5 DC A 1 4.611 -2.644 -1.810 1.00 0.00 C
 ATOM 16 C6 DC A 1 4.487 -1.362 -1.387 1.00 0.00 C
 ATOM 17 P DA A 2 1.893 3.783 1.227 1.00 0.00 P
 ATOM 18 OP1 DA A 2 2.387 5.132 1.940 1.00 0.00 O
 ATOM 19 OP2 DA A 2 1.559 2.757 2.245 1.00 0.00 O
 ATOM 20 O5' DA A 2 0.709 4.320 0.287 1.00 0.00 O
 ATOM 21 C5' DA A 2 0.403 3.818 -1.038 1.00 0.00 C
 ATOM 22 C4' DA A 2 -0.552 2.643 -1.002 1.00 0.00 C

ATOM 23 O4' DA A 2 0.140 1.464 -0.557 1.00 0.00 O
 ATOM 24 C3' DA A 2 -1.761 2.812 -0.049 1.00 0.00 C
 ATOM 25 O3' DA A 2 -2.912 2.372 -0.812 1.00 0.00 O
 ATOM 26 C2' DA A 2 -1.448 1.848 1.093 1.00 0.00 C
 ATOM 27 C1' DA A 2 -0.738 0.746 0.316 1.00 0.00 C
 ATOM 28 N9 DA A 2 0.068 -0.174 1.091 1.00 0.00 N
 ATOM 29 C8 DA A 2 0.651 -0.000 2.317 1.00 0.00 C
 ATOM 30 N7 DA A 2 1.517 -0.960 2.639 1.00 0.00 N
 ATOM 31 C5 DA A 2 1.496 -1.798 1.571 1.00 0.00 C
 ATOM 32 C6 DA A 2 2.221 -2.987 1.260 1.00 0.00 C
 ATOM 33 N6 DA A 2 3.127 -3.491 2.090 1.00 0.00 N
 ATOM 34 N1 DA A 2 1.975 -3.605 0.081 1.00 0.00 N
 ATOM 35 C2 DA A 2 1.069 -3.076 -0.738 1.00 0.00 C
 ATOM 36 N3 DA A 2 0.340 -1.936 -0.574 1.00 0.00 N
 ATOM 37 C4 DA A 2 0.595 -1.352 0.580 1.00 0.00 C
 ATOM 38 P DA A 3 -4.422 2.417 -0.272 1.00 0.00 P
 ATOM 39 OP1 DA A 3 -4.848 3.964 -0.204 1.00 0.00 O
 ATOM 40 OP2 DA A 3 -4.696 1.772 1.037 1.00 0.00 O
 ATOM 41 O5' DA A 3 -5.215 1.846 -1.536 1.00 0.00 O
 ATOM 42 C5' DA A 3 -4.643 1.026 -2.587 1.00 0.00 C
 ATOM 43 C4' DA A 3 -4.605 -0.440 -2.216 1.00 0.00 C
 ATOM 44 O4' DA A 3 -3.606 -0.665 -1.201 1.00 0.00 O
 ATOM 45 C3' DA A 3 -5.940 -1.007 -1.661 1.00 0.00 C
 ATOM 46 O3' DA A 3 -6.226 -2.185 -2.409 1.00 0.00 O
 ATOM 47 C2' DA A 3 -5.594 -1.324 -0.200 1.00 0.00 C
 ATOM 48 C1' DA A 3 -4.122 -1.688 -0.347 1.00 0.00 C
 ATOM 49 N9 DA A 3 -3.346 -1.721 0.871 1.00 0.00 N
 ATOM 50 C8 DA A 3 -3.376 -0.838 1.924 1.00 0.00 C
 ATOM 51 N7 DA A 3 -2.511 -1.131 2.885 1.00 0.00 N
 ATOM 52 C5 DA A 3 -1.889 -2.263 2.436 1.00 0.00 C
 ATOM 53 C6 DA A 3 -0.862 -3.076 2.981 1.00 0.00 C
 ATOM 54 N6 DA A 3 -0.307 -2.816 4.164 1.00 0.00 N
 ATOM 55 N1 DA A 3 -0.419 -4.140 2.260 1.00 0.00 N
 ATOM 56 C2 DA A 3 -0.991 -4.389 1.083 1.00 0.00 C
 ATOM 57 N3 DA A 3 -1.979 -3.701 0.465 1.00 0.00 N
 ATOM 58 C4 DA A 3 -2.378 -2.656 1.180 1.00 0.00 C

END

COMPND ACC / D-B3LYP / NA MODEL / NEUTRAL

ATOM 1 O5' DA A 1 7.165 0.633 0.116 1.00 0.00 O
 ATOM 2 C5' DA A 1 6.985 1.644 -0.873 1.00 0.00 C
 ATOM 3 C4' DA A 1 5.501 1.892 -1.110 1.00 0.00 C
 ATOM 4 O4' DA A 1 4.874 0.718 -1.658 1.00 0.00 O
 ATOM 5 C3' DA A 1 4.691 2.229 0.150 1.00 0.00 C
 ATOM 6 O3' DA A 1 3.547 2.987 -0.296 1.00 0.00 O
 ATOM 7 C2' DA A 1 4.273 0.847 0.631 1.00 0.00 C
 ATOM 8 C1' DA A 1 3.968 0.158 -0.701 1.00 0.00 C
 ATOM 9 N9 DA A 1 4.109 -1.284 -0.664 1.00 0.00 N
 ATOM 10 C8 DA A 1 4.941 -2.053 0.125 1.00 0.00 C
 ATOM 11 N7 DA A 1 4.717 -3.344 0.032 1.00 0.00 N
 ATOM 12 C5 DA A 1 3.686 -3.436 -0.888 1.00 0.00 C
 ATOM 13 C6 DA A 1 2.932 -4.522 -1.373 1.00 0.00 C
 ATOM 14 N6 DA A 1 3.084 -5.780 -0.875 1.00 0.00 N
 ATOM 15 N1 DA A 1 1.982 -4.294 -2.298 1.00 0.00 N
 ATOM 16 C2 DA A 1 1.746 -3.023 -2.664 1.00 0.00 C
 ATOM 17 N3 DA A 1 2.335 -1.904 -2.230 1.00 0.00 N
 ATOM 18 C4 DA A 1 3.301 -2.173 -1.342 1.00 0.00 C
 ATOM 19 P DC A 2 2.448 3.587 0.763 1.00 0.00 P
 ATOM 20 OP1 DC A 2 3.005 4.773 1.523 1.00 0.00 O
 ATOM 21 OP2 DC A 2 1.857 2.497 1.646 1.00 0.00 O
 ATOM 22 O5' DC A 2 1.362 4.156 -0.310 1.00 0.00 O
 ATOM 23 C5' DC A 2 1.051 3.580 -1.588 1.00 0.00 C
 ATOM 24 C4' DC A 2 0.000 2.481 -1.552 1.00 0.00 C
 ATOM 25 O4' DC A 2 0.598 1.218 -1.217 1.00 0.00 O
 ATOM 26 C3' DC A 2 -1.162 2.706 -0.546 1.00 0.00 C
 ATOM 27 O3' DC A 2 -2.384 2.709 -1.312 1.00 0.00 O
 ATOM 28 C2' DC A 2 -1.039 1.506 0.394 1.00 0.00 C
 ATOM 29 C1' DC A 2 -0.411 0.466 -0.532 1.00 0.00 C
 ATOM 30 N1 DC A 2 0.203 -0.685 0.103 1.00 0.00 N
 ATOM 31 C2 DC A 2 -0.175 -2.002 -0.311 1.00 0.00 C
 ATOM 32 O2 DC A 2 -1.039 -2.143 -1.177 1.00 0.00 O
 ATOM 33 N3 DC A 2 0.456 -3.055 0.285 1.00 0.00 N
 ATOM 34 C4 DC A 2 1.398 -2.853 1.193 1.00 0.00 C
 ATOM 35 N4 DC A 2 1.995 -3.950 1.737 1.00 0.00 N
 ATOM 36 C5 DC A 2 1.776 -1.547 1.648 1.00 0.00 C
 ATOM 37 C6 DC A 2 1.149 -0.491 1.063 1.00 0.00 C
 ATOM 38 P DC A 3 -3.837 2.872 -0.554 1.00 0.00 P
 ATOM 39 OP1 DC A 3 -4.713 3.742 -1.427 1.00 0.00 O
 ATOM 40 OP2 DC A 3 -3.662 3.262 0.896 1.00 0.00 O
 ATOM 41 O5' DC A 3 -4.389 1.315 -0.571 1.00 0.00 O
 ATOM 42 C5' DC A 3 -4.651 0.739 -1.864 1.00 0.00 C
 ATOM 43 C4' DC A 3 -5.356 -0.596 -1.717 1.00 0.00 C
 ATOM 44 O4' DC A 3 -4.466 -1.567 -1.141 1.00 0.00 O
 ATOM 45 C3' DC A 3 -6.610 -0.570 -0.825 1.00 0.00 C
 ATOM 46 O3' DC A 3 -7.538 -1.494 -1.394 1.00 0.00 O
 ATOM 47 C2' DC A 3 -6.069 -1.065 0.516 1.00 0.00 C
 ATOM 48 C1' DC A 3 -5.011 -2.083 0.075 1.00 0.00 C
 ATOM 49 N1 DC A 3 -3.917 -2.272 1.026 1.00 0.00 N
 ATOM 50 C2 DC A 3 -3.593 -3.580 1.497 1.00 0.00 C
 ATOM 51 O2 DC A 3 -4.295 -4.537 1.160 1.00 0.00 O
 ATOM 52 N3 DC A 3 -2.506 -3.707 2.315 1.00 0.00 N
 ATOM 53 C4 DC A 3 -1.790 -2.641 2.653 1.00 0.00 C
 ATOM 54 N4 DC A 3 -0.734 -2.825 3.488 1.00 0.00 N
 ATOM 55 C5 DC A 3 -2.095 -1.316 2.198 1.00 0.00 C
 ATOM 56 C6 DC A 3 -3.154 -1.194 1.361 1.00 0.00 C

END

COMPND ACC / D-B3LYP / NA MODEL / SINGLY IONIZED

ATOM 1 O5' DA A 1 -7.099 -1.048 0.057 1.00 0.00 O
 ATOM 2 C5' DA A 1 -6.812 -2.024 -0.945 1.00 0.00 C
 ATOM 3 C4' DA A 1 -5.308 -2.158 -1.134 1.00 0.00 C
 ATOM 4 O4' DA A 1 -4.755 -0.953 -1.711 1.00 0.00 O
 ATOM 5 C3' DA A 1 -4.507 -2.386 0.155 1.00 0.00 C
 ATOM 6 O3' DA A 1 -3.307 -3.081 -0.228 1.00 0.00 O
 ATOM 7 C2' DA A 1 -4.187 -0.960 0.591 1.00 0.00 C
 ATOM 8 C1' DA A 1 -3.944 -0.282 -0.763 1.00 0.00 C
 ATOM 9 N9 DA A 1 -4.304 1.139 -0.751 1.00 0.00 N
 ATOM 10 C8 DA A 1 -5.469 1.689 -0.317 1.00 0.00 C
 ATOM 11 N7 DA A 1 -5.501 3.029 -0.396 1.00 0.00 N
 ATOM 12 C5 DA A 1 -4.299 3.350 -0.913 1.00 0.00 C
 ATOM 13 C6 DA A 1 -3.690 4.598 -1.266 1.00 0.00 C
 ATOM 14 N6 DA A 1 -4.306 5.753 -1.071 1.00 0.00 N
 ATOM 15 N1 DA A 1 -2.450 4.590 -1.820 1.00 0.00 N
 ATOM 16 C2 DA A 1 -1.854 3.426 -2.014 1.00 0.00 C
 ATOM 17 N3 DA A 1 -2.334 2.163 -1.745 1.00 0.00 N
 ATOM 18 C4 DA A 1 -3.517 2.191 -1.181 1.00 0.00 C
 ATOM 19 P DC A 2 -2.278 -3.640 0.922 1.00 0.00 P
 ATOM 20 OP1 DC A 2 -2.909 -4.764 1.716 1.00 0.00 O
 ATOM 21 OP2 DC A 2 -1.732 -2.501 1.776 1.00 0.00 O

ATOM 22 O5' DC A 2 -1.130 -4.268 -0.045 1.00 0.00 O
 ATOM 23 C5' DC A 2 -0.841 -3.874 -1.397 1.00 0.00 C
 ATOM 24 C4' DC A 2 0.186 -2.763 -1.509 1.00 0.00 C
 ATOM 25 O4' DC A 2 -0.420 -1.481 -1.249 1.00 0.00 O
 ATOM 26 C3' DC A 2 1.397 -2.887 -0.543 1.00 0.00 C
 ATOM 27 O3' DC A 2 2.583 -2.807 -1.360 1.00 0.00 O
 ATOM 28 C2' DC A 2 1.212 -1.681 0.380 1.00 0.00 C
 ATOM 29 C1' DC A 2 0.577 -0.693 -0.592 1.00 0.00 C
 ATOM 30 N1 DC A 2 -0.050 0.481 -0.011 1.00 0.00 N
 ATOM 31 C2 DC A 2 0.306 1.772 -0.495 1.00 0.00 C
 ATOM 32 O2 DC A 2 1.104 1.889 -1.421 1.00 0.00 O
 ATOM 33 N3 DC A 2 -0.282 2.861 0.098 1.00 0.00 N
 ATOM 34 C4 DC A 2 -1.239 2.694 1.013 1.00 0.00 C
 ATOM 35 N4 DC A 2 -1.888 3.804 1.414 1.00 0.00 N
 ATOM 36 C5 DC A 2 -1.619 1.409 1.507 1.00 0.00 C
 ATOM 37 C6 DC A 2 -0.989 0.330 0.962 1.00 0.00 C
 ATOM 38 P DC A 3 4.073 -2.683 -0.672 1.00 0.00 P
 ATOM 39 OP1 DC A 3 5.056 -3.347 -1.608 1.00 0.00 O
 ATOM 40 OP2 DC A 3 4.052 -3.117 0.775 1.00 0.00 O
 ATOM 41 O5' DC A 3 4.303 -1.047 -0.673 1.00 0.00 O
 ATOM 42 C5' DC A 3 4.468 -0.396 -1.947 1.00 0.00 C
 ATOM 43 C4' DC A 3 5.174 0.937 -1.773 1.00 0.00 C
 ATOM 44 O4' DC A 3 4.310 1.885 -1.123 1.00 0.00 O
 ATOM 45 C3' DC A 3 6.459 0.875 -0.931 1.00 0.00 C
 ATOM 46 O3' DC A 3 7.359 1.836 -1.482 1.00 0.00 O
 ATOM 47 C2' DC A 3 5.964 1.291 0.455 1.00 0.00 C
 ATOM 48 C1' DC A 3 4.879 2.319 0.111 1.00 0.00 C
 ATOM 49 N1 DC A 3 3.805 2.418 1.101 1.00 0.00 N
 ATOM 50 C2 DC A 3 3.494 3.676 1.704 1.00 0.00 C
 ATOM 51 O2 DC A 3 4.198 4.657 1.452 1.00 0.00 O
 ATOM 52 N3 DC A 3 2.416 3.727 2.541 1.00 0.00 N
 ATOM 53 C4 DC A 3 1.705 2.634 2.792 1.00 0.00 C
 ATOM 54 N4 DC A 3 0.656 2.744 3.639 1.00 0.00 N
 ATOM 55 C5 DC A 3 2.007 1.354 2.218 1.00 0.00 C
 ATOM 56 C6 DC A 3 3.054 1.310 1.356 1.00 0.00 C
 END

COMPND AAC / D-B3LYP / NA MODEL / NEUTRAL
 ATOM 1 O5' DA A 1 6.694 2.349 -0.633 1.00 0.00 O
 ATOM 2 C5' DA A 1 6.012 2.782 -1.806 1.00 0.00 C
 ATOM 3 C4' DA A 1 4.537 2.434 -1.689 1.00 0.00 C
 ATOM 4 O4' DA A 1 4.368 1.002 -1.751 1.00 0.00 O
 ATOM 5 C3' DA A 1 3.898 2.882 -0.362 1.00 0.00 C
 ATOM 6 O3' DA A 1 2.536 3.253 -0.644 1.00 0.00 O
 ATOM 7 C2' DA A 1 3.947 1.608 0.477 1.00 0.00 C
 ATOM 8 C1' DA A 1 3.663 0.558 -0.593 1.00 0.00 C
 ATOM 9 N9 DA A 1 4.115 -0.777 -0.251 1.00 0.00 N
 ATOM 10 C8 DA A 1 4.754 -1.204 0.895 1.00 0.00 C
 ATOM 11 N7 DA A 1 4.950 -2.502 0.943 1.00 0.00 N
 ATOM 12 C5 DA A 1 4.409 -2.963 -0.245 1.00 0.00 C
 ATOM 13 C6 DA A 1 4.229 -4.257 -0.769 1.00 0.00 C
 ATOM 14 N6 DA A 1 4.580 -5.369 -0.068 1.00 0.00 N
 ATOM 15 N1 DA A 1 3.626 -4.395 -1.964 1.00 0.00 N
 ATOM 16 C2 DA A 1 3.165 -3.289 -2.573 1.00 0.00 C
 ATOM 17 N3 DA A 1 3.229 -2.017 -2.166 1.00 0.00 N
 ATOM 18 C4 DA A 1 3.872 -1.912 -0.994 1.00 0.00 C
 ATOM 19 P DA A 2 1.697 4.055 0.513 1.00 0.00 P
 ATOM 20 OP1 DA A 2 2.399 5.351 0.870 1.00 0.00 O
 ATOM 21 OP2 DA A 2 1.391 3.161 1.708 1.00 0.00 O
 ATOM 22 O5' DA A 2 0.323 4.367 -0.303 1.00 0.00 O
 ATOM 23 C5' DA A 2 -0.024 3.930 -1.627 1.00 0.00 C
 ATOM 24 C4' DA A 2 -0.884 2.676 -1.617 1.00 0.00 C
 ATOM 25 O4' DA A 2 -0.075 1.515 -1.350 1.00 0.00 O
 ATOM 26 C3' DA A 2 -1.993 2.690 -0.539 1.00 0.00 C
 ATOM 27 O3' DA A 2 -3.169 2.114 -1.142 1.00 0.00 O
 ATOM 28 C2' DA A 2 -1.422 1.766 0.532 1.00 0.00 C
 ATOM 29 C1' DA A 2 -0.752 0.727 -0.361 1.00 0.00 C
 ATOM 30 N9 DA A 2 0.189 -0.145 0.291 1.00 0.00 N
 ATOM 31 C8 DA A 2 0.986 0.122 1.388 1.00 0.00 C
 ATOM 32 N7 DA A 2 1.647 -0.930 1.822 1.00 0.00 N
 ATOM 33 C5 DA A 2 1.262 -1.950 0.966 1.00 0.00 C
 ATOM 34 C6 DA A 2 1.546 -3.328 0.927 1.00 0.00 C
 ATOM 35 N6 DA A 2 2.318 -3.940 1.877 1.00 0.00 N
 ATOM 36 N1 DA A 2 0.989 -4.078 -0.039 1.00 0.00 N
 ATOM 37 C2 DA A 2 0.152 -3.486 -0.905 1.00 0.00 C
 ATOM 38 N3 DA A 2 -0.230 -2.206 -0.953 1.00 0.00 N
 ATOM 39 C4 DA A 2 0.359 -1.485 0.008 1.00 0.00 C
 ATOM 40 P DC A 3 -4.596 2.135 -0.334 1.00 0.00 P
 ATOM 41 OP1 DC A 3 -5.080 3.557 -0.131 1.00 0.00 O
 ATOM 42 OP2 DC A 3 -4.535 1.306 0.942 1.00 0.00 O
 ATOM 43 O5' DC A 3 -5.533 1.432 -1.466 1.00 0.00 O
 ATOM 44 C5' DC A 3 -5.095 0.485 -2.454 1.00 0.00 C
 ATOM 45 C4' DC A 3 -5.059 -0.953 -1.972 1.00 0.00 C
 ATOM 46 O4' DC A 3 -3.884 -1.198 -1.184 1.00 0.00 O
 ATOM 47 C3' DC A 3 -6.278 -1.399 -1.118 1.00 0.00 C
 ATOM 48 O3' DC A 3 -6.868 -2.515 -1.787 1.00 0.00 O
 ATOM 49 C2' DC A 3 -5.633 -1.791 0.216 1.00 0.00 C
 ATOM 50 C1' DC A 3 -4.247 -2.229 -0.256 1.00 0.00 C
 ATOM 51 N1 DC A 3 -3.224 -2.346 0.768 1.00 0.00 N
 ATOM 52 C2 DC A 3 -2.551 -3.595 0.951 1.00 0.00 C
 ATOM 53 O2 DC A 3 -2.864 -4.562 0.252 1.00 0.00 O
 ATOM 54 N3 DC A 3 -1.583 -3.655 1.910 1.00 0.00 N
 ATOM 55 C4 DC A 3 -1.268 -2.581 2.622 1.00 0.00 C
 ATOM 56 N4 DC A 3 -0.280 -2.709 3.552 1.00 0.00 N
 ATOM 57 C5 DC A 3 -1.935 -1.322 2.468 1.00 0.00 C
 ATOM 58 C6 DC A 3 -2.908 -1.254 1.521 1.00 0.00 C
 END

COMPND AAC / D-B3LYP / NA MODEL / SINGLY IONIZED
 ATOM 1 O5' DA A 1 6.440 1.524 -0.259 1.00 0.00 O
 ATOM 2 C5' DA A 1 6.103 2.379 -1.349 1.00 0.00 C
 ATOM 3 C4' DA A 1 4.591 2.461 -1.506 1.00 0.00 C
 ATOM 4 O4' DA A 1 4.038 1.169 -1.824 1.00 0.00 O
 ATOM 5 C3' DA A 1 3.823 2.910 -0.259 1.00 0.00 C
 ATOM 6 O3' DA A 1 2.564 3.432 -0.726 1.00 0.00 O
 ATOM 7 C2' DA A 1 3.615 1.593 0.480 1.00 0.00 C
 ATOM 8 C1' DA A 1 3.360 0.636 -0.689 1.00 0.00 C
 ATOM 9 N9 DA A 1 3.824 -0.731 -0.457 1.00 0.00 N
 ATOM 10 C8 DA A 1 4.820 -1.197 0.359 1.00 0.00 C
 ATOM 11 N7 DA A 1 5.060 -2.497 0.225 1.00 0.00 N
 ATOM 12 C5 DA A 1 4.192 -2.894 -0.751 1.00 0.00 C
 ATOM 13 C6 DA A 1 3.958 -4.147 -1.378 1.00 0.00 C
 ATOM 14 N6 DA A 1 4.616 -5.252 -1.017 1.00 0.00 N
 ATOM 15 N1 DA A 1 3.043 -4.212 -2.372 1.00 0.00 N
 ATOM 16 C2 DA A 1 2.380 -3.099 -2.695 1.00 0.00 C
 ATOM 17 N3 DA A 1 2.495 -1.858 -2.171 1.00 0.00 N
 ATOM 18 C4 DA A 1 3.409 -1.819 -1.208 1.00 0.00 C
 ATOM 19 P DA A 2 1.523 4.168 0.305 1.00 0.00 P

ATOM 20 OP1 DA A 2 2.007 5.559 0.658 1.00 0.00 O
 ATOM 21 OP2 DA A 2 1.213 3.290 1.510 1.00 0.00 O
 ATOM 22 O5' DA A 2 0.220 4.286 -0.668 1.00 0.00 O
 ATOM 23 C5' DA A 2 0.001 3.613 -1.917 1.00 0.00 C
 ATOM 24 C4' DA A 2 -0.849 2.361 -1.779 1.00 0.00 C
 ATOM 25 O4' DA A 2 -0.059 1.269 -1.261 1.00 0.00 O
 ATOM 26 C3' DA A 2 -2.057 2.505 -0.823 1.00 0.00 C
 ATOM 27 O3' DA A 2 -3.145 1.797 -1.440 1.00 0.00 O
 ATOM 28 C2' DA A 2 -1.573 1.779 0.431 1.00 0.00 C
 ATOM 29 C1' DA A 2 -0.825 0.630 -0.234 1.00 0.00 C
 ATOM 30 N9 DA A 2 0.082 -0.119 0.609 1.00 0.00 N
 ATOM 31 C8 DA A 2 0.807 0.320 1.680 1.00 0.00 C
 ATOM 32 N7 DA A 2 1.617 -0.606 2.193 1.00 0.00 N
 ATOM 33 C5 DA A 2 1.399 -1.697 1.417 1.00 0.00 C
 ATOM 34 C6 DA A 2 1.943 -3.013 1.424 1.00 0.00 C
 ATOM 35 N6 DA A 2 2.888 -3.370 2.288 1.00 0.00 N
 ATOM 36 N1 DA A 2 1.498 -3.910 0.510 1.00 0.00 N
 ATOM 37 C2 DA A 2 0.571 -3.519 -0.356 1.00 0.00 C
 ATOM 38 N3 DA A 2 0.000 -2.291 -0.492 1.00 0.00 N
 ATOM 39 C4 DA A 2 0.443 -1.438 0.410 1.00 0.00 C
 ATOM 40 P DC A 3 -4.655 1.797 -0.805 1.00 0.00 P
 ATOM 41 OP1 DC A 3 -5.301 3.162 -0.938 1.00 0.00 O
 ATOM 42 OP2 DC A 3 -4.680 1.232 0.609 1.00 0.00 O
 ATOM 43 O5' DC A 3 -5.364 0.782 -1.866 1.00 0.00 O
 ATOM 44 C5' DC A 3 -4.692 -0.219 -2.649 1.00 0.00 C
 ATOM 45 C4' DC A 3 -4.507 -1.545 -1.936 1.00 0.00 C
 ATOM 46 O4' DC A 3 -3.451 -1.462 -0.965 1.00 0.00 O
 ATOM 47 C3' DC A 3 -5.762 -2.075 -1.191 1.00 0.00 C
 ATOM 48 O3' DC A 3 -6.035 -3.375 -1.714 1.00 0.00 O
 ATOM 49 C2' DC A 3 -5.299 -2.121 0.270 1.00 0.00 C
 ATOM 50 C1' DC A 3 -3.804 -2.376 0.080 1.00 0.00 C
 ATOM 51 N1 DC A 3 -2.947 -2.151 1.232 1.00 0.00 N
 ATOM 52 C2 DC A 3 -2.003 -3.154 1.610 1.00 0.00 C
 ATOM 53 O2 DC A 3 -1.947 -4.209 0.971 1.00 0.00 O
 ATOM 54 N3 DC A 3 -1.180 -2.892 2.665 1.00 0.00 N
 ATOM 55 C4 DC A 3 -1.286 -1.743 3.336 1.00 0.00 C
 ATOM 56 N4 DC A 3 -0.466 -1.564 4.390 1.00 0.00 N
 ATOM 57 C5 DC A 3 -2.220 -0.720 2.976 1.00 0.00 C
 ATOM 58 C6 DC A 3 -3.024 -0.968 1.907 1.00 0.00 C
 END

COMPND CCA / D-B3LYP / NA MODEL / NEUTRAL

ATOM 1 O5' DC A 1 -7.309 -0.916 -0.182 1.00 0.00 O
 ATOM 2 C5' DC A 1 -6.975 -1.875 -1.184 1.00 0.00 C
 ATOM 3 C4' DC A 1 -5.464 -1.965 -1.340 1.00 0.00 C
 ATOM 4 O4' DC A 1 -4.947 -0.750 -1.910 1.00 0.00 O
 ATOM 5 C3' DC A 1 -4.699 -2.166 -0.022 1.00 0.00 C
 ATOM 6 O3' DC A 1 -3.497 -2.894 -0.351 1.00 0.00 O
 ATOM 7 C2' DC A 1 -4.373 -0.734 0.384 1.00 0.00 C
 ATOM 8 C1' DC A 1 -4.066 -0.103 -0.977 1.00 0.00 C
 ATOM 9 N1 DC A 1 -4.275 1.337 -1.046 1.00 0.00 N
 ATOM 10 C2 DC A 1 -3.279 2.163 -1.657 1.00 0.00 C
 ATOM 11 O2 DC A 1 -2.309 1.636 -2.204 1.00 0.00 O
 ATOM 12 N3 DC A 1 -3.456 3.515 -1.591 1.00 0.00 N
 ATOM 13 C4 DC A 1 -4.563 4.023 -1.064 1.00 0.00 C
 ATOM 14 N4 DC A 1 -4.644 5.369 -0.957 1.00 0.00 N
 ATOM 15 C5 DC A 1 -5.653 3.213 -0.597 1.00 0.00 C
 ATOM 16 C6 DC A 1 -5.456 1.870 -0.614 1.00 0.00 C
 ATOM 17 P DC A 2 -2.520 -3.452 0.841 1.00 0.00 P
 ATOM 18 OP1 DC A 2 -3.157 -4.615 1.574 1.00 0.00 O
 ATOM 19 OP2 DC A 2 -2.054 -2.323 1.751 1.00 0.00 O
 ATOM 20 O5' DC A 2 -1.302 -4.032 -0.071 1.00 0.00 O
 ATOM 21 C5' DC A 2 -0.925 -3.602 -1.390 1.00 0.00 C
 ATOM 22 C4' DC A 2 0.143 -2.519 -1.411 1.00 0.00 C
 ATOM 23 O4' DC A 2 -0.445 -1.223 -1.217 1.00 0.00 O
 ATOM 24 C3' DC A 2 1.246 -2.672 -0.330 1.00 0.00 C
 ATOM 25 O3' DC A 2 2.513 -2.535 -1.009 1.00 0.00 O
 ATOM 26 C2' DC A 2 0.968 -1.494 0.599 1.00 0.00 C
 ATOM 27 C1' DC A 2 0.476 -0.464 -0.413 1.00 0.00 C
 ATOM 28 N1 DC A 2 -0.200 0.710 0.100 1.00 0.00 N
 ATOM 29 C2 DC A 2 0.147 2.000 -0.411 1.00 0.00 C
 ATOM 30 O2 DC A 2 1.056 2.103 -1.236 1.00 0.00 O
 ATOM 31 N3 DC A 2 -0.540 3.078 0.069 1.00 0.00 N
 ATOM 32 C4 DC A 2 -1.525 2.913 0.936 1.00 0.00 C
 ATOM 33 N4 DC A 2 -2.143 4.034 1.411 1.00 0.00 N
 ATOM 34 C5 DC A 2 -1.920 1.633 1.439 1.00 0.00 C
 ATOM 35 C6 DC A 2 -1.225 0.557 0.981 1.00 0.00 C
 ATOM 36 P DA A 3 3.925 -2.833 -0.219 1.00 0.00 P
 ATOM 37 OP1 DA A 3 4.573 -4.071 -0.805 1.00 0.00 O
 ATOM 38 OP2 DA A 3 3.742 -2.789 1.283 1.00 0.00 O
 ATOM 39 O5' DA A 3 4.770 -1.487 -0.629 1.00 0.00 O
 ATOM 40 C5' DA A 3 4.948 -1.155 -2.013 1.00 0.00 C
 ATOM 41 C4' DA A 3 5.529 0.244 -2.119 1.00 0.00 C
 ATOM 42 O4' DA A 3 4.584 1.207 -1.619 1.00 0.00 O
 ATOM 43 C3' DA A 3 6.825 0.469 -1.326 1.00 0.00 C
 ATOM 44 O3' DA A 3 7.582 1.441 -2.046 1.00 0.00 O
 ATOM 45 C2' DC A 3 6.308 1.026 -0.000 1.00 0.00 C
 ATOM 46 C1' DA A 3 5.087 1.841 -0.444 1.00 0.00 C
 ATOM 47 N9 DA A 3 4.039 1.910 0.556 1.00 0.00 N
 ATOM 48 C8 DA A 3 3.444 0.856 1.221 1.00 0.00 C
 ATOM 49 N7 DA A 3 2.450 1.219 2.000 1.00 0.00 N
 ATOM 50 C5 DA A 3 2.376 2.591 1.834 1.00 0.00 C
 ATOM 51 C6 DA A 3 1.496 3.566 2.341 1.00 0.00 C
 ATOM 52 N6 DA A 3 0.496 3.262 3.216 1.00 0.00 N
 ATOM 53 N1 DA A 3 1.670 4.850 1.970 1.00 0.00 N
 ATOM 54 C2 DC A 3 2.661 5.135 1.108 1.00 0.00 C
 ATOM 55 N3 DA A 3 3.539 4.304 0.534 1.00 0.00 N
 ATOM 56 C4 DA A 3 3.345 3.040 0.932 1.00 0.00 C
 END

COMPND CCA / D-B3LYP / NA MODEL / SINGLY IONIZED

ATOM 1 O5' DC A 1 -7.286 -0.567 -0.118 1.00 0.00 O
 ATOM 2 C5' DC A 1 -7.029 -1.636 -1.028 1.00 0.00 C
 ATOM 3 C4' DC A 1 -5.530 -1.858 -1.164 1.00 0.00 C
 ATOM 4 O4' DC A 1 -4.916 -0.740 -1.832 1.00 0.00 O
 ATOM 5 C3' DC A 1 -4.781 -2.005 0.170 1.00 0.00 C
 ATOM 6 O3' DC A 1 -3.639 -2.846 -0.099 1.00 0.00 O
 ATOM 7 C2' DC A 1 -4.347 -0.573 0.452 1.00 0.00 C
 ATOM 8 C1' DC A 1 -3.980 -0.096 -0.956 1.00 0.00 C
 ATOM 9 N1 DC A 1 -4.067 1.347 -1.151 1.00 0.00 N
 ATOM 10 C2 DC A 1 -2.987 2.043 -1.780 1.00 0.00 C
 ATOM 11 O2 DC A 1 -2.013 1.403 -2.187 1.00 0.00 O
 ATOM 12 N3 DC A 1 -3.081 3.398 -1.880 1.00 0.00 N
 ATOM 13 C4 DC A 1 -4.182 4.028 -1.483 1.00 0.00 C
 ATOM 14 N4 DC A 1 -4.194 5.373 -1.556 1.00 0.00 N
 ATOM 15 C5 DC A 1 -5.341 3.338 -0.986 1.00 0.00 C
 ATOM 16 C6 DC A 1 -5.231 1.993 -0.845 1.00 0.00 C
 ATOM 17 P DC A 2 -2.643 -3.339 1.103 1.00 0.00 P

ATOM 18 OP2 DC A 2 -2.057 -2.155 1.865 1.00 0.00 O
 ATOM 19 OP1 DC A 2 -3.308 -4.372 1.988 1.00 0.00 O
 ATOM 20 O5' DC A 2 -1.508 -4.086 0.204 1.00 0.00 O
 ATOM 21 C5' DC A 2 -1.186 -3.812 -1.170 1.00 0.00 C
 ATOM 22 C4' DC A 2 -0.074 -2.794 -1.350 1.00 0.00 C
 ATOM 23 O4' DC A 2 -0.587 -1.454 -1.196 1.00 0.00 O
 ATOM 24 C3' DC A 2 1.114 -2.936 -0.357 1.00 0.00 C
 ATOM 25 O3' DC A 2 2.321 -2.997 -1.154 1.00 0.00 O
 ATOM 26 C2' DC A 2 0.966 -1.680 0.500 1.00 0.00 C
 ATOM 27 C1' DC A 2 0.442 -0.703 -0.548 1.00 0.00 C
 ATOM 28 N1 DC A 2 -0.115 0.547 -0.043 1.00 0.00 N
 ATOM 29 C2 DC A 2 0.349 1.805 -0.586 1.00 0.00 C
 ATOM 30 O2 DC A 2 1.251 1.794 -1.417 1.00 0.00 O
 ATOM 31 N3 DC A 2 -0.214 2.953 -0.120 1.00 0.00 N
 ATOM 32 C4 DC A 2 -1.240 2.908 0.721 1.00 0.00 C
 ATOM 33 N4 DC A 2 -1.787 4.068 1.126 1.00 0.00 N
 ATOM 34 C5 DC A 2 -1.719 1.674 1.276 1.00 0.00 C
 ATOM 35 C6 DC A 2 -1.117 0.511 0.850 1.00 0.00 C
 ATOM 36 P DA A 3 3.777 -2.409 -0.690 1.00 0.00 P
 ATOM 37 OP1 DA A 3 4.803 -2.955 -1.657 1.00 0.00 O
 ATOM 38 OP2 DA A 3 4.030 -2.599 0.795 1.00 0.00 O
 ATOM 39 O5' DA A 3 3.587 -0.790 -0.890 1.00 0.00 O
 ATOM 40 C5' DA A 3 3.888 -0.145 -2.140 1.00 0.00 C
 ATOM 41 C4' DA A 3 5.087 0.776 -1.972 1.00 0.00 C
 ATOM 42 O4' DA A 3 4.783 1.885 -1.105 1.00 0.00 O
 ATOM 43 C3' DA A 3 6.317 0.114 -1.343 1.00 0.00 C
 ATOM 44 O3' DA A 3 7.459 0.810 -1.838 1.00 0.00 O
 ATOM 45 C2' DA A 3 6.106 0.353 0.163 1.00 0.00 C
 ATOM 46 C1' DA A 3 5.232 1.622 0.208 1.00 0.00 C
 ATOM 47 N9 DA A 3 4.066 1.518 1.113 1.00 0.00 N
 ATOM 48 C8 DA A 3 3.501 0.431 1.725 1.00 0.00 C
 ATOM 49 N7 DA A 3 2.380 0.705 2.391 1.00 0.00 N
 ATOM 50 C5 2.195 2.036 2.186 1.00 0.00 C
 ATOM 51 C6 1.202 2.951 2.615 1.00 0.00 C
 ATOM 52 N6 DA A 3 0.146 2.554 3.335 1.00 0.00 N
 ATOM 53 N1 DA A 3 1.306 4.256 2.265 1.00 0.00 N
 ATOM 54 C2 DA A 3 2.340 4.619 1.514 1.00 0.00 C
 ATOM 55 N3 DA A 3 3.349 3.849 1.029 1.00 0.00 N
 ATOM 56 C4 DA A 3 3.233 2.583 1.393 1.00 0.00 C

END

COMPND CAA / D-B3LYP / NA MODEL / NEUTRAL

ATOM 1 O5' DC A 1 -7.176 -0.880 -0.067 1.00 0.00 O
 ATOM 2 C5' DC A 1 -6.902 -1.870 -1.058 1.00 0.00 C
 ATOM 3 C4' DC A 1 -5.399 -2.009 -1.254 1.00 0.00 C
 ATOM 4 O4' DC A 1 -4.864 -0.815 -1.851 1.00 0.00 O
 ATOM 5 C3' DC A 1 -4.609 -2.214 0.048 1.00 0.00 C
 ATOM 6 O3' DC A 1 -3.430 -2.979 -0.288 1.00 0.00 O
 ATOM 7 C2' DC A 1 -4.232 -0.788 0.428 1.00 0.00 C
 ATOM 8 C1' DC A 1 -3.942 -0.185 -0.950 1.00 0.00 C
 ATOM 9 N1 DC A 1 -4.096 1.261 -1.045 1.00 0.00 N
 ATOM 10 C2 DC A 1 -3.145 2.012 -1.807 1.00 0.00 C
 ATOM 11 O2 DC A 1 -2.264 1.419 -2.430 1.00 0.00 O
 ATOM 12 N3 DC A 1 -3.260 3.372 -1.793 1.00 0.00 N
 ATOM 13 C4 DC A 1 -4.274 3.955 -1.163 1.00 0.00 C
 ATOM 14 N4 DC A 1 -4.272 5.310 -1.106 1.00 0.00 N
 ATOM 15 C5 DC A 1 -5.340 3.219 -0.553 1.00 0.00 C
 ATOM 16 C6 DC A 1 -5.205 1.868 -0.528 1.00 0.00 C
 ATOM 17 P DA A 2 -2.551 -3.648 0.923 1.00 0.00 P
 ATOM 18 OP1 DA A 2 -3.313 -4.786 1.573 1.00 0.00 O
 ATOM 19 OP2 DA A 2 -2.053 -2.593 1.902 1.00 0.00 O
 ATOM 20 O5' DA A 2 -1.309 -4.256 0.064 1.00 0.00 O
 ATOM 21 C5' DA A 2 -0.959 -3.984 -1.302 1.00 0.00 C
 ATOM 22 C4' DA A 2 0.178 -2.983 -1.439 1.00 0.00 C
 ATOM 23 O4' DA A 2 -0.318 -1.637 -1.307 1.00 0.00 O
 ATOM 24 C3' DA A 2 1.302 -3.144 -0.386 1.00 0.00 C
 ATOM 25 O3' DA A 2 2.547 -3.036 -1.106 1.00 0.00 O
 ATOM 26 C2' DA A 2 1.041 -1.967 0.549 1.00 0.00 C
 ATOM 27 C1' DA A 2 0.573 -0.912 -0.446 1.00 0.00 C
 ATOM 28 N9 DA A 2 -0.126 0.210 0.117 1.00 0.00 N
 ATOM 29 C8 DA A 2 -1.057 0.187 1.131 1.00 0.00 C
 ATOM 30 N7 DA A 2 -1.563 1.371 1.407 1.00 0.00 N
 ATOM 31 C5 DA A 2 -0.915 2.223 0.528 1.00 0.00 C
 ATOM 32 C6 DA A 2 -1.005 3.608 0.312 1.00 0.00 C
 ATOM 33 N6 DA A 2 -1.862 4.392 1.031 1.00 0.00 N
 ATOM 34 N1 DA A 2 -0.184 4.176 -0.590 1.00 0.00 N
 ATOM 35 C2 DA A 2 0.664 3.381 -1.264 1.00 0.00 C
 ATOM 36 N3 DA A 2 0.811 2.053 -1.186 1.00 0.00 N
 ATOM 37 C4 DA A 2 -0.011 1.526 -0.274 1.00 0.00 C
 ATOM 38 P DA A 3 4.003 -3.003 -0.340 1.00 0.00 P
 ATOM 39 OP1 DA A 3 4.938 -3.933 -1.081 1.00 0.00 O
 ATOM 40 OP2 DA A 3 3.860 -3.184 1.154 1.00 0.00 O
 ATOM 41 O5' DA A 3 4.447 -1.436 -0.592 1.00 0.00 O
 ATOM 42 C5' DA A 3 4.642 -0.016 -1.953 1.00 0.00 C
 ATOM 43 C4' DA A 3 5.237 0.380 -1.984 1.00 0.00 C
 ATOM 44 O4' DA A 3 4.298 1.339 -1.467 1.00 0.00 O
 ATOM 45 C3' DA A 3 6.519 0.550 -1.157 1.00 0.00 C
 ATOM 46 O3' DA A 3 7.306 1.540 -1.818 1.00 0.00 O
 ATOM 47 C2' DA A 3 5.979 1.057 0.180 1.00 0.00 C
 ATOM 48 C1' DA A 3 4.785 1.914 -0.257 1.00 0.00 C
 ATOM 49 N9 DA A 3 3.714 1.957 0.719 1.00 0.00 N
 ATOM 50 C8 DA A 3 3.066 0.882 1.294 1.00 0.00 C
 ATOM 51 N7 DA A 3 2.067 1.223 2.075 1.00 0.00 N
 ATOM 52 C5 DA A 3 2.043 2.607 2.003 1.00 0.00 C
 ATOM 53 C6 DA A 3 1.189 3.583 2.554 1.00 0.00 C
 ATOM 54 N6 DA A 3 0.126 3.268 3.349 1.00 0.00 N
 ATOM 55 N1 DA A 3 1.438 4.882 2.293 1.00 0.00 N
 ATOM 56 C2 DA A 3 2.461 5.187 1.478 1.00 0.00 C
 ATOM 57 N3 DA A 3 3.310 4.362 0.857 1.00 0.00 N
 ATOM 58 C4 DA A 3 3.053 3.081 1.159 1.00 0.00 C

END

COMPND CAA / D-B3LYP / NA MODEL / SINGLY IONIZED

ATOM 1 O5' DC A 1 -7.152 0.834 0.489 1.00 0.00 O
 ATOM 2 C5' DC A 1 -6.710 1.786 1.457 1.00 0.00 C
 ATOM 3 C4' DC A 1 -5.192 1.879 1.432 1.00 0.00 C
 ATOM 4 O4' DC A 1 -4.606 0.670 1.955 1.00 0.00 O
 ATOM 5 C3' DC A 1 -4.595 2.061 0.029 1.00 0.00 C
 ATOM 6 O3' DC A 1 -3.380 2.821 0.178 1.00 0.00 O
 ATOM 7 C2' DC A 1 -4.283 0.626 -0.387 1.00 0.00 C
 ATOM 8 C1' DC A 1 -3.850 0.003 0.946 1.00 0.00 O
 ATOM 9 N1 DC A 1 -4.113 -1.437 1.045 1.00 0.00 N
 ATOM 10 C2 DC A 1 -3.092 -2.334 1.452 1.00 0.00 C
 ATOM 11 O2 DC A 1 -1.979 -1.880 1.777 1.00 0.00 O
 ATOM 12 N3 DC A 1 -3.357 -3.663 1.460 1.00 0.00 N
 ATOM 13 C4 DC A 1 -4.588 -4.104 1.185 1.00 0.00 C
 ATOM 14 N4 DC A 1 -4.783 -5.432 1.177 1.00 0.00 N
 ATOM 15 C5 DC A 1 -5.680 -3.214 0.921 1.00 0.00 C

ATOM 16 C6 DC A 1 -5.390 -1.887 0.861 1.00 0.00 C
 ATOM 17 P2 DA A 2 -2.715 3.493 -1.160 1.00 0.00 P
 ATOM 18 OP2 DA A 2 -2.143 2.430 -2.091 1.00 0.00 O
 ATOM 19 OP1 DA A 2 -3.698 4.433 -1.826 1.00 0.00 O
 ATOM 20 O5' DA A 2 -1.496 4.335 -0.484 1.00 0.00 O
 ATOM 21 C5' DA A 2 -1.110 4.343 0.899 1.00 0.00 C
 ATOM 22 C4' DA A 2 0.040 3.390 1.177 1.00 0.00 C
 ATOM 23 O4' DA A 2 -0.437 2.026 1.134 1.00 0.00 O
 ATOM 24 C3' DA A 2 1.212 3.482 0.160 1.00 0.00 C
 ATOM 25 O3' DA A 2 2.441 3.483 0.926 1.00 0.00 O
 ATOM 26 C2' DA A 2 0.984 2.236 -0.691 1.00 0.00 C
 ATOM 27 C1' DA A 2 0.516 1.273 0.390 1.00 0.00 C
 ATOM 28 N9 DA A 2 -0.088 0.051 -0.087 1.00 0.00 N
 ATOM 29 C8 DA A 2 -1.006 -0.110 -1.082 1.00 0.00 C
 ATOM 30 N7 DA A 2 -1.332 -1.390 -1.310 1.00 0.00 N
 ATOM 31 C5 DA A 2 -0.568 -2.078 -0.434 1.00 0.00 C
 ATOM 32 C6 DA A 2 -0.454 -3.466 -0.118 1.00 0.00 C
 ATOM 33 N6 DA A 2 -1.249 -4.365 -0.679 1.00 0.00 N
 ATOM 34 N1 DA A 2 0.486 -3.859 0.775 1.00 0.00 N
 ATOM 35 C2 DA A 2 1.221 -2.924 1.364 1.00 0.00 C
 ATOM 36 N3 DA A 2 1.144 -1.569 1.233 1.00 0.00 N
 ATOM 37 C4 DA A 2 0.253 -1.211 0.341 1.00 0.00 C
 ATOM 38 P DA A 3 3.835 2.712 0.536 1.00 0.00 P
 ATOM 39 OP2 DA A 3 4.086 2.698 -0.960 1.00 0.00 O
 ATOM 40 OP1 DA A 3 4.915 3.271 1.433 1.00 0.00 O
 ATOM 41 O5' DA A 3 3.509 1.151 0.925 1.00 0.00 O
 ATOM 42 C5' DA A 3 3.779 0.598 2.222 1.00 0.00 C
 ATOM 43 C4' DA A 3 4.850 -0.477 2.109 1.00 0.00 C
 ATOM 44 O4' DA A 3 4.369 -1.616 1.374 1.00 0.00 O
 ATOM 45 C3' DA A 3 6.117 -0.024 1.372 1.00 0.00 C
 ATOM 46 O3' DA A 3 7.199 -0.788 1.903 1.00 0.00 O
 ATOM 47 C2' DA A 3 5.815 -0.397 -0.087 1.00 0.00 C
 ATOM 48 C1' DA A 3 4.885 -1.615 0.048 1.00 0.00 C
 ATOM 49 N9 DA A 3 3.786 -1.638 -0.917 1.00 0.00 N
 ATOM 50 C8 DA A 3 3.083 -0.595 -1.491 1.00 0.00 C
 ATOM 51 N7 DA A 3 2.054 -0.983 -2.214 1.00 0.00 N
 ATOM 52 C5 DA A 3 2.067 -2.363 -2.103 1.00 0.00 C
 ATOM 53 C6 DA A 3 1.263 -3.385 -2.657 1.00 0.00 C
 ATOM 54 N6 DA A 3 0.209 -3.126 -3.464 1.00 0.00 N
 ATOM 55 N1 DA A 3 1.562 -4.670 -2.371 1.00 0.00 N
 ATOM 56 C2 DA A 3 2.610 -4.921 -1.572 1.00 0.00 C
 ATOM 57 N3 DA A 3 3.448 -4.052 -0.990 1.00 0.00 N
 ATOM 58 C4 DA A 3 3.134 -2.788 -1.300 1.00 0.00 C
 END

COMPND ACC / D-B3LYP / PO- MODEL / NEUTRAL
 ATOM 1 O5' DA A 1 -7.224 -0.279 0.355 1.00 0.00 O
 ATOM 2 C5' DA A 1 -7.076 -1.356 -0.567 1.00 0.00 C
 ATOM 3 C4' DA A 1 -5.601 -1.677 -0.764 1.00 0.00 C
 ATOM 4 O4' DA A 1 -4.928 -0.564 -1.384 1.00 0.00 O
 ATOM 5 C3' DA A 1 -4.822 -1.961 0.531 1.00 0.00 C
 ATOM 6 O3' DA A 1 -3.740 -2.832 0.180 1.00 0.00 O
 ATOM 7 C2' DA A 1 -4.312 -0.571 0.898 1.00 0.00 C
 ATOM 8 C1' DA A 1 -3.949 -0.034 -0.484 1.00 0.00 C
 ATOM 9 N9 DA A 1 -3.925 1.413 -0.575 1.00 0.00 N
 ATOM 10 C8 DA A 1 -4.587 2.342 0.203 1.00 0.00 C
 ATOM 11 N7 DA A 1 -4.220 3.584 -0.022 1.00 0.00 N
 ATOM 12 C5 DA A 1 -3.265 3.471 -1.019 1.00 0.00 C
 ATOM 13 C6 DA A 1 -2.423 4.410 -1.645 1.00 0.00 C
 ATOM 14 N6 DA A 1 -2.389 5.713 -1.254 1.00 0.00 N
 ATOM 15 N1 DA A 1 -1.575 3.989 -2.600 1.00 0.00 N
 ATOM 16 C2 DA A 1 -1.520 2.673 -2.864 1.00 0.00 C
 ATOM 17 N3 DA A 1 -2.212 1.677 -2.301 1.00 0.00 N
 ATOM 18 C4 DA A 1 -3.071 2.136 -1.381 1.00 0.00 C
 ATOM 19 P DC A 2 -2.733 -3.422 1.361 1.00 0.00 P
 ATOM 20 OP2 DC A 2 -2.002 -2.291 2.054 1.00 0.00 O
 ATOM 21 OP1 DC A 2 -3.468 -4.432 2.199 1.00 0.00 O
 ATOM 22 O5' DC A 2 -1.709 -4.253 0.374 1.00 0.00 O
 ATOM 23 C5' DC A 2 -1.410 -3.939 -0.988 1.00 0.00 C
 ATOM 24 C4' DC A 2 -0.260 -2.964 -1.188 1.00 0.00 C
 ATOM 25 O4' DC A 2 -0.705 -1.604 -1.030 1.00 0.00 O
 ATOM 26 C3' DC A 2 0.946 -3.156 -0.229 1.00 0.00 C
 ATOM 27 O3' DC A 2 2.114 -3.314 -1.049 1.00 0.00 O
 ATOM 28 C2' DC A 2 0.941 -1.857 0.579 1.00 0.00 C
 ATOM 29 C1' DC A 2 0.389 -0.880 -0.455 1.00 0.00 C
 ATOM 30 N1 DC A 2 -0.091 0.395 0.049 1.00 0.00 N
 ATOM 31 C2 DC A 2 0.414 1.609 -0.513 1.00 0.00 C
 ATOM 32 O2 DC A 2 1.228 1.561 -1.437 1.00 0.00 O
 ATOM 33 N3 DC A 2 -0.036 2.786 0.014 1.00 0.00 N
 ATOM 34 C4 DC A 2 -0.950 2.789 0.974 1.00 0.00 C
 ATOM 35 N4 DC A 2 -1.370 3.996 1.445 1.00 0.00 N
 ATOM 36 C5 DC A 2 -1.472 1.587 1.555 1.00 0.00 C
 ATOM 37 C6 DC A 2 -1.013 0.410 1.051 1.00 0.00 C
 ATOM 38 P DC A 3 3.613 -3.430 -0.340 1.00 0.00 P
 ATOM 39 OP1 DC A 3 4.451 -4.272 -1.261 1.00 0.00 O
 ATOM 40 OP2 DC A 3 3.479 -3.701 1.131 1.00 0.00 O
 ATOM 41 O5' DC A 3 4.110 -1.838 -0.459 1.00 0.00 O
 ATOM 42 C5' DC A 3 4.354 -1.319 -1.769 1.00 0.00 C
 ATOM 43 C4' DC A 3 5.305 -0.135 -1.711 1.00 0.00 C
 ATOM 44 O4' DC A 3 4.651 1.024 -1.160 1.00 0.00 O
 ATOM 45 C3' DC A 3 6.556 -0.367 -0.851 1.00 0.00 C
 ATOM 46 O3' DC A 3 7.621 0.367 -1.456 1.00 0.00 O
 ATOM 47 C2' DC A 3 6.144 0.235 0.494 1.00 0.00 C
 ATOM 48 C1' DC A 3 5.263 1.416 0.067 1.00 0.00 C
 ATOM 49 N1 DC A 3 4.205 1.752 1.022 1.00 0.00 N
 ATOM 50 C2 DC A 3 4.122 3.064 1.576 1.00 0.00 C
 ATOM 51 O2 DC A 3 5.009 3.886 1.326 1.00 0.00 O
 ATOM 52 N3 DC A 3 3.044 3.350 2.367 1.00 0.00 N
 ATOM 53 C4 DC A 3 2.135 2.418 2.627 1.00 0.00 C
 ATOM 54 N4 DC A 3 1.095 2.751 3.434 1.00 0.00 N
 ATOM 55 C5 DC A 3 2.215 1.081 2.117 1.00 0.00 C
 ATOM 56 C6 DC A 3 3.254 0.811 1.288 1.00 0.00 C
 END

COMPND ACC / D-B3LYP / PO- MODEL / SINGLY IONIZED
 ATOM 1 O5' DA A 1 -7.135 0.513 -0.429 1.00 0.00 O
 ATOM 2 C5' DA A 1 -6.977 1.615 0.463 1.00 0.00 C
 ATOM 3 C4' DA A 1 -5.502 1.922 0.668 1.00 0.00 C
 ATOM 4 O4' DA A 1 -4.854 0.840 1.378 1.00 0.00 O
 ATOM 5 C3' DA A 1 -4.684 2.106 -0.618 1.00 0.00 C
 ATOM 6 O3' DA A 1 -3.581 2.956 -0.289 1.00 0.00 O
 ATOM 7 C2' DA A 1 -4.210 0.682 -0.899 1.00 0.00 C
 ATOM 8 C1' DA A 1 -3.944 0.173 0.521 1.00 0.00 C
 ATOM 9 N9 DA A 1 -4.147 -1.275 0.645 1.00 0.00 N
 ATOM 10 C8 DA A 1 -5.219 -1.995 0.221 1.00 0.00 C
 ATOM 11 N7 DA A 1 -5.104 -3.317 0.427 1.00 0.00 N
 ATOM 12 C5 DA A 1 -3.900 -3.447 1.018 1.00 0.00 C
 ATOM 13 C6 DA A 1 -3.173 -4.578 1.510 1.00 0.00 C

ATOM 14 N6 DA A 1 -3.642 -5.810 1.397 1.00 0.00
 ATOM 15 N1 DA A 1 -1.971 -4.376 2.110 1.00 0.00 N
 ATOM 16 C2 DA A 1 -1.522 -3.136 2.217 1.00 0.00 C
 ATOM 17 N3 DA A 1 -2.126 -1.968 1.809 1.00 0.00 N
 ATOM 18 C4 DA A 1 -3.269 -2.184 1.204 1.00 0.00 C
 ATOM 19 P DC A 2 -2.575 3.488 -1.500 1.00 0.00 P
 ATOM 20 OP1 DC A 2 -3.298 4.484 -2.364 1.00 0.00 O
 ATOM 21 OP2 DC A 2 -1.887 2.311 -2.163 1.00 0.00 O
 ATOM 22 O5' DC A 2 -1.506 4.314 -0.563 1.00 0.00 O
 ATOM 23 C5' DC A 2 -1.271 4.129 0.838 1.00 0.00 C
 ATOM 24 C4' DC A 2 -0.125 3.184 1.162 1.00 0.00 C
 ATOM 25 O4' DC A 2 -0.557 1.807 1.077 1.00 0.00 O
 ATOM 26 C3' DC A 2 1.121 3.322 0.241 1.00 0.00 C
 ATOM 27 O3' DC A 2 2.269 3.433 1.102 1.00 0.00 O
 ATOM 28 C2' DC A 2 1.051 2.029 -0.575 1.00 0.00 C
 ATOM 29 C1' DC A 2 0.525 1.077 0.491 1.00 0.00 C
 ATOM 30 N1 DC A 2 0.038 -0.211 0.019 1.00 0.00 N
 ATOM 31 C2 DC A 2 0.522 -1.406 0.623 1.00 0.00 C
 ATOM 32 O2 DC A 2 1.294 -1.357 1.578 1.00 0.00 O
 ATOM 33 N3 DC A 2 0.081 -2.603 0.113 1.00 0.00 N
 ATOM 34 C4 DC A 2 -0.853 -2.627 -0.840 1.00 0.00 C
 ATOM 35 N4 DC A 2 -1.358 -3.835 -1.159 1.00 0.00 N
 ATOM 36 C5 DC A 2 -1.359 -1.441 -1.452 1.00 0.00 C
 ATOM 37 C6 DC A 2 -0.883 -0.253 -0.980 1.00 0.00 C
 ATOM 38 P DC A 3 3.793 2.975 0.630 1.00 0.00 P
 ATOM 39 OP2 DC A 3 3.972 3.164 -0.852 1.00 0.00 O
 ATOM 40 OP1 DC A 3 4.735 3.548 1.651 1.00 0.00 O
 ATOM 41 O5' DC A 3 3.671 1.330 0.848 1.00 0.00 O
 ATOM 42 C5' DC A 3 4.045 0.703 2.078 1.00 0.00 C
 ATOM 43 C4' DC A 3 5.218 -0.241 1.848 1.00 0.00 C
 ATOM 44 O4' DC A 3 4.796 -1.400 1.103 1.00 0.00 O
 ATOM 45 C3' DC A 3 6.365 0.385 1.044 1.00 0.00 C
 ATOM 46 O3' DC A 3 7.576 -0.223 1.498 1.00 0.00 O
 ATOM 47 C2' DC A 3 6.026 -0.024 -0.393 1.00 0.00 C
 ATOM 48 C1' DC A 3 5.376 -1.404 -0.201 1.00 0.00 C
 ATOM 49 N1 DC A 3 4.327 -1.731 -1.174 1.00 0.00 N
 ATOM 50 C2 DC A 3 4.346 -2.993 -1.846 1.00 0.00 C
 ATOM 51 O2 DC A 3 5.321 -3.738 -1.710 1.00 0.00 O
 ATOM 52 N3 DC A 3 3.265 -3.323 -2.616 1.00 0.00 N
 ATOM 53 C4 DC A 3 2.251 -2.474 -2.746 1.00 0.00 C
 ATOM 54 N4 DC A 3 1.203 -2.849 -3.513 1.00 0.00 N
 ATOM 55 C5 DC A 3 2.220 -1.189 -2.112 1.00 0.00 C
 ATOM 56 C6 DC A 3 3.269 -0.882 -1.307 1.00 0.00 C
 END

COMPND AAC / D-B3LYP / PO- MODEL / NEUTRAL
 ATOM 1 O5' DA A 1 -7.334 -0.465 0.195 1.00 0.00 O
 ATOM 2 C5' DA A 1 -6.960 -1.328 -0.874 1.00 0.00 C
 ATOM 3 C4' DA A 1 -5.448 -1.466 -0.908 1.00 0.00 C
 ATOM 4 O4' DA A 1 -4.854 -0.211 -1.305 1.00 0.00 O
 ATOM 5 C3' DA A 1 -4.819 -1.827 0.453 1.00 0.00 C
 ATOM 6 O3' DA A 1 -3.690 -2.669 0.192 1.00 0.00 O
 ATOM 7 C2' DA A 1 -4.363 -0.462 0.968 1.00 0.00 C
 ATOM 8 C1' DA A 1 -3.866 0.148 -0.336 1.00 0.00 C
 ATOM 9 N9 DA A 1 -3.709 1.586 -0.322 1.00 0.00 N
 ATOM 10 C8 DA A 1 -4.080 2.492 0.648 1.00 0.00 C
 ATOM 11 N7 DA A 1 -3.687 3.723 0.407 1.00 0.00 N
 ATOM 12 C5 DA A 1 -3.012 3.624 -0.799 1.00 0.00 C
 ATOM 13 C6 DA A 1 -2.303 4.561 -1.576 1.00 0.00 C
 ATOM 14 N6 DA A 1 -2.160 5.857 -1.193 1.00 0.00 N
 ATOM 15 N1 DA A 1 -1.695 4.139 -2.702 1.00 0.00 N
 ATOM 16 C2 DA A 1 -1.768 2.834 -3.016 1.00 0.00 C
 ATOM 17 N3 DA A 1 -2.393 1.848 -2.363 1.00 0.00 N
 ATOM 18 C4 DA A 1 -3.002 2.306 -1.261 1.00 0.00 C
 ATOM 19 P DA A 2 -2.967 -3.437 1.471 1.00 0.00 P
 ATOM 20 OP1 DC A 2 -3.945 -4.399 2.091 1.00 0.00 O
 ATOM 21 OP2 DC A 2 -2.258 -2.436 2.360 1.00 0.00 O
 ATOM 22 O5' DA A 2 -1.818 -4.275 0.639 1.00 0.00 O
 ATOM 23 C5' DA A 2 -1.611 -4.290 -0.774 1.00 0.00 C
 ATOM 24 C4' DA A 2 -0.421 -3.439 -1.191 1.00 0.00 C
 ATOM 25 O4' DA A 2 -0.767 -2.035 -1.151 1.00 0.00 O
 ATOM 26 C3' DA A 2 0.828 -3.613 -0.288 1.00 0.00 C
 ATOM 27 O3' DA A 2 1.972 -3.669 -1.161 1.00 0.00 O
 ATOM 28 C2' DA A 2 0.757 -2.355 0.575 1.00 0.00 C
 ATOM 29 C1' DA A 2 0.281 -1.349 -0.462 1.00 0.00 C
 ATOM 30 N9 DA A 2 -0.206 -0.103 0.079 1.00 0.00 N
 ATOM 31 C8 DA A 2 -1.014 0.072 1.183 1.00 0.00 C
 ATOM 32 N7 DA A 2 -1.160 1.338 1.522 1.00 0.00 N
 ATOM 33 C5 DA A 2 -0.388 2.028 0.602 1.00 0.00 C
 ATOM 34 C6 DA A 2 -0.086 3.394 0.442 1.00 0.00 C
 ATOM 35 N6 DA A 2 -0.632 4.359 1.235 1.00 0.00 N
 ATOM 36 N1 DA A 2 0.795 3.752 -0.511 1.00 0.00 N
 ATOM 37 C2 DA A 2 1.312 2.792 -1.297 1.00 0.00 C
 ATOM 38 N3 DA A 2 1.070 1.478 -1.280 1.00 0.00 N
 ATOM 39 C4 DA A 2 0.220 1.151 -0.298 1.00 0.00 C
 ATOM 40 P DC A 3 3.502 -3.239 -0.681 1.00 0.00 P
 ATOM 41 OP1 DC A 3 4.434 -3.793 -1.724 1.00 0.00 O
 ATOM 42 OP2 DC A 3 3.684 -3.478 0.794 1.00 0.00 O
 ATOM 43 O5' DC A 3 3.404 -1.587 -0.845 1.00 0.00 O
 ATOM 44 C5' DC A 3 3.835 -0.916 -2.031 1.00 0.00 C
 ATOM 45 C4' DC A 3 5.069 -0.070 -1.745 1.00 0.00 C
 ATOM 46 O4' DC A 3 4.728 1.064 -0.927 1.00 0.00 O
 ATOM 47 C3' DC A 3 6.173 -0.819 -0.987 1.00 0.00 C
 ATOM 48 O3' DC A 3 7.421 -0.254 -1.400 1.00 0.00 O
 ATOM 49 C2' DC A 3 5.859 -0.489 0.478 1.00 0.00 C
 ATOM 50 C1' DC A 3 5.281 0.932 0.384 1.00 0.00 C
 ATOM 51 N1 DC A 3 4.237 1.258 1.362 1.00 0.00 N
 ATOM 52 C2 DC A 3 4.225 2.560 1.956 1.00 0.00 C
 ATOM 53 O2 DC A 3 5.204 3.299 1.816 1.00 0.00 O
 ATOM 54 N3 DC A 3 3.108 2.938 2.649 1.00 0.00 N
 ATOM 55 C4 DC A 3 2.088 2.098 2.791 1.00 0.00 C
 ATOM 56 N4 DC A 3 0.992 2.531 3.468 1.00 0.00 N
 ATOM 57 C5 DC A 3 2.100 0.766 2.266 1.00 0.00 C
 ATOM 58 C6 DC A 3 3.173 0.416 1.515 1.00 0.00 C
 END

COMPND AAC / D-B3LYP / PO- MODEL / SINGLY IONIZED
 ATOM 1 O5' DA A 1 -7.277 0.884 -0.165 1.00 0.00 O
 ATOM 2 C5' DA A 1 -7.075 -0.129 -1.143 1.00 0.00 C
 ATOM 3 C4' DA A 1 -5.651 -0.650 -1.053 1.00 0.00 C
 ATOM 4 O4' DA A 1 -4.725 0.380 -1.479 1.00 0.00 O
 ATOM 5 C3' DA A 1 -5.209 -1.052 0.371 1.00 0.00 C
 ATOM 6 O3' DA A 1 -4.299 -2.149 0.224 1.00 0.00 O
 ATOM 7 C2' DA A 1 -4.477 0.206 0.838 1.00 0.00 C
 ATOM 8 C1' DA A 1 -3.748 0.542 -0.456 1.00 0.00 C
 ATOM 9 N9 DA A 1 -3.199 1.887 -0.512 1.00 0.00 N
 ATOM 10 C8 DA A 1 -3.359 2.918 0.384 1.00 0.00 C
 ATOM 11 N7 DA A 1 -2.621 3.985 0.120 1.00 0.00 N

ATOM 12 C5 DA A 1 -1.930 3.628 -1.002 1.00 0.00 C
 ATOM 13 C6 DA A 1 -0.953 4.305 -1.783 1.00 0.00 C
 ATOM 14 N6 DA A 1 -0.560 5.546 -1.481 1.00 0.00 N
 ATOM 15 N1 DA A 1 -0.405 3.657 -2.832 1.00 0.00 N
 ATOM 16 C2 DA A 1 -0.818 2.412 -3.099 1.00 0.00 C
 ATOM 17 N3 DA A 1 -1.744 1.668 -2.460 1.00 0.00 N
 ATOM 18 C4 DA A 1 -2.261 2.319 -1.424 1.00 0.00 C
 ATOM 19 P DA A 2 -3.608 -2.865 1.550 1.00 0.00 P
 ATOM 20 OP2 DA A 2 -2.702 -1.886 2.272 1.00 0.00 O
 ATOM 21 OP1 DA A 2 -4.654 -3.600 2.341 1.00 0.00 O
 ATOM 22 O5' DA A 2 -2.681 -3.971 0.753 1.00 0.00 O
 ATOM 23 C5' DA A 2 -2.385 -3.985 -0.645 1.00 0.00 C
 ATOM 24 C4' DA A 2 -1.082 -3.290 -1.000 1.00 0.00 C
 ATOM 25 O4' DA A 2 -1.243 -1.850 -0.935 1.00 0.00 O
 ATOM 26 C3' DA A 2 0.120 -3.645 -0.081 1.00 0.00 C
 ATOM 27 O3' DA A 2 1.242 -3.905 -0.944 1.00 0.00 O
 ATOM 28 C2' DA A 2 0.249 -2.376 0.761 1.00 0.00 C
 ATOM 29 C1' DA A 2 -0.078 -1.340 -0.302 1.00 0.00 C
 ATOM 30 N9 DA A 2 -0.314 -0.002 0.210 1.00 0.00 N
 ATOM 31 C8 DA A 2 -1.066 0.375 1.290 1.00 0.00 C
 ATOM 32 N7 DA A 2 -0.900 1.656 1.631 1.00 0.00 N
 ATOM 33 C5 DA A 2 0.015 2.122 0.736 1.00 0.00 C
 ATOM 34 C6 DA A 2 0.723 3.352 0.617 1.00 0.00 C
 ATOM 35 N6 DA A 2 0.418 4.412 1.360 1.00 0.00 N
 ATOM 36 N1 DA A 2 1.727 3.446 -0.288 1.00 0.00 N
 ATOM 37 C2 DA A 2 1.969 2.402 -1.075 1.00 0.00 C
 ATOM 38 N3 DA A 2 1.340 1.199 -1.098 1.00 0.00 N
 ATOM 39 C4 DA A 2 0.415 1.107 -0.165 1.00 0.00 C
 ATOM 40 P DC A 3 2.825 -3.646 -0.512 1.00 0.00 P
 ATOM 41 OP1 DC A 3 3.654 -4.362 -1.541 1.00 0.00 O
 ATOM 42 OP2 DC A 3 3.013 -3.823 0.970 1.00 0.00 O
 ATOM 43 O5' DC A 3 2.932 -2.003 -0.762 1.00 0.00 O
 ATOM 44 C5' DC A 3 3.332 -1.456 -2.022 1.00 0.00 C
 ATOM 45 C4' DC A 3 4.642 -0.698 -1.871 1.00 0.00 C
 ATOM 46 O4' DC A 3 4.449 0.499 -1.091 1.00 0.00 O
 ATOM 47 C3' DC A 3 5.738 -1.497 -1.152 1.00 0.00 C
 ATOM 48 O3' DC A 3 6.990 -1.067 -1.691 1.00 0.00 O
 ATOM 49 C2' DC A 3 5.571 -1.061 0.308 1.00 0.00 C
 ATOM 50 C1' DC A 3 5.109 0.398 0.169 1.00 0.00 C
 ATOM 51 N1 DC A 3 4.188 0.856 1.218 1.00 0.00 N
 ATOM 52 C2 DC A 3 4.412 2.123 1.857 1.00 0.00 C
 ATOM 53 O2 DC A 3 5.445 2.746 1.609 1.00 0.00 O
 ATOM 54 N3 DC A 3 3.450 2.586 2.709 1.00 0.00 N
 ATOM 55 C4 DC A 3 2.357 1.869 2.943 1.00 0.00 C
 ATOM 56 N4 DC A 3 1.421 2.385 3.771 1.00 0.00 N
 ATOM 57 C5 DC A 3 2.126 0.584 2.350 1.00 0.00 C
 ATOM 58 C6 DC A 3 3.062 0.136 1.471 1.00 0.00 C
 END

COMPND CCA / D-B3LYP / PO- MODEL / NEUTRAL
 ATOM 1 O5' DC A 1 -7.303 -0.945 0.089 1.00 0.00 O
 ATOM 2 C5' DC A 1 -6.980 -0.985 -0.834 1.00 0.00 C
 ATOM 3 C4' DC A 1 -5.470 -2.090 -0.995 1.00 0.00 C
 ATOM 4 O4' DC A 1 -4.961 -0.920 -1.662 1.00 0.00 O
 ATOM 5 C3' DC A 1 -4.693 -2.193 0.328 1.00 0.00 C
 ATOM 6 O3' DC A 1 -3.505 -2.955 0.065 1.00 0.00 O
 ATOM 7 C2' DC A 1 -4.353 -0.734 0.617 1.00 0.00 C
 ATOM 8 C1' DC A 1 -4.067 -0.210 -0.792 1.00 0.00 C
 ATOM 9 N1 DC A 1 -4.272 1.224 -0.968 1.00 0.00 N
 ATOM 10 C2 DC A 1 -3.291 1.994 -1.668 1.00 0.00 C
 ATOM 11 O2 DC A 1 -2.343 1.420 -2.207 1.00 0.00 O
 ATOM 12 N3 DC A 1 -3.455 3.349 -1.696 1.00 0.00 N
 ATOM 13 C4 DC A 1 -4.553 3.902 -1.192 1.00 0.00 C
 ATOM 14 N4 DC A 1 -4.628 5.252 -1.189 1.00 0.00 N
 ATOM 15 C5 DC A 1 -5.637 3.135 -0.648 1.00 0.00 C
 ATOM 16 C6 DC A 1 -5.443 1.794 -0.560 1.00 0.00 C
 ATOM 17 P DC A 2 -2.587 -3.461 1.353 1.00 0.00 P
 ATOM 18 OP1 DC A 2 -3.310 -4.555 2.093 1.00 0.00 O
 ATOM 19 OP2 DC A 2 -2.074 -2.271 2.136 1.00 0.00 O
 ATOM 20 O5' DC A 2 -1.362 -4.161 0.505 1.00 0.00 O
 ATOM 21 C5' DC A 2 -0.991 -3.912 -0.855 1.00 0.00 C
 ATOM 22 C4' DC A 2 0.145 -2.912 -1.024 1.00 0.00 C
 ATOM 23 O4' DC A 2 -0.358 -1.563 -1.003 1.00 0.00 O
 ATOM 24 C3' DC A 2 1.247 -2.991 0.062 1.00 0.00 C
 ATOM 25 O3' DC A 2 2.510 -2.957 -0.623 1.00 0.00 O
 ATOM 26 C2' DC A 2 0.990 -1.726 0.879 1.00 0.00 C
 ATOM 27 C1' DC A 2 0.562 -0.772 -0.230 1.00 0.00 C
 ATOM 28 N1 DC A 2 -0.097 0.462 0.160 1.00 0.00 N
 ATOM 29 C2 DC A 2 0.272 1.687 -0.478 1.00 0.00 C
 ATOM 30 O2 DC A 2 1.169 1.689 -1.323 1.00 0.00 O
 ATOM 31 N3 DC A 2 -0.383 2.824 -0.098 1.00 0.00 N
 ATOM 32 C4 DC A 2 -1.360 2.769 0.791 1.00 0.00 C
 ATOM 33 N4 DC A 2 -1.946 3.948 1.161 1.00 0.00 N
 ATOM 34 C5 DC A 2 -1.778 1.554 1.417 1.00 0.00 C
 ATOM 35 C6 DC A 2 -1.118 0.420 1.059 1.00 0.00 C
 ATOM 36 P DA A 3 3.933 -3.190 0.203 1.00 0.00 P
 ATOM 37 OP1 DA A 3 4.589 -4.430 -0.340 1.00 0.00 O
 ATOM 38 OP2 DA A 3 3.737 -2.960 1.677 1.00 0.00 O
 ATOM 39 O5' DA A 3 4.756 -1.853 -0.340 1.00 0.00 O
 ATOM 40 C5' DA A 3 4.953 -1.689 -1.742 1.00 0.00 C
 ATOM 41 C4' DA A 3 5.586 -0.332 -2.004 1.00 0.00 C
 ATOM 42 O4' DA A 3 4.678 0.719 -1.621 1.00 0.00 O
 ATOM 43 C3' DA A 3 6.885 -0.067 -1.231 1.00 0.00 C
 ATOM 44 O3' DA A 3 7.676 0.805 -2.039 1.00 0.00 O
 ATOM 45 C2' DA A 3 6.381 0.632 0.031 1.00 0.00 C
 ATOM 46 C1' DA A 3 5.182 1.432 -0.494 1.00 0.00 C
 ATOM 47 N9 DA A 3 4.129 1.608 0.490 1.00 0.00 N
 ATOM 48 C8 DA A 3 3.507 0.632 1.246 1.00 0.00 C
 ATOM 49 N7 DA A 3 2.529 1.091 1.994 1.00 0.00 N
 ATOM 50 C5 DA A 3 2.495 2.445 1.711 1.00 0.00 C
 ATOM 51 C6 DA A 3 1.648 3.486 2.130 1.00 0.00 C
 ATOM 52 N6 DA A 3 0.646 3.286 3.040 1.00 0.00 N
 ATOM 53 N1 DA A 3 1.855 4.728 1.654 1.00 0.00 N
 ATOM 54 C2 DA A 3 2.847 4.908 0.766 1.00 0.00 C
 ATOM 55 N3 DA A 3 3.698 4.005 0.263 1.00 0.00 N
 ATOM 56 C4 DA A 3 3.470 2.785 0.769 1.00 0.00 C
 END

COMPND CCA / D-B3LYP / PO- MODEL / SINGLY IONIZED
 ATOM 1 O5' DC A 1 -7.234 -0.406 0.138 1.00 0.00 O
 ATOM 2 C5' DC A 1 -7.023 -1.552 -0.686 1.00 0.00 C
 ATOM 3 C4' DC A 1 -5.535 -1.838 -0.817 1.00 0.00 C
 ATOM 4 O4' DC A 1 -4.891 -0.794 -1.573 1.00 0.00 O
 ATOM 5 C3' DC A 1 -4.772 -1.916 0.517 1.00 0.00 C
 ATOM 6 O3' DC A 1 -3.672 -2.811 0.308 1.00 0.00 O
 ATOM 7 C2' DC A 1 -4.287 -0.481 0.687 1.00 0.00 C
 ATOM 8 C1' DC A 1 -3.920 -0.126 -0.756 1.00 0.00 C
 ATOM 9 N1 DC A 1 -3.951 1.301 -1.061 1.00 0.00 N

ATOM 10 C2 DC A 1 -2.861 1.902 -1.765 1.00 0.00 C
 ATOM 11 O2 DC A 1 -1.929 1.191 -2.153 1.00 0.00 O
 ATOM 12 N3 DC A 1 -2.897 3.250 -1.958 1.00 0.00 N
 ATOM 13 C4 DC A 1 -3.961 3.954 -1.584 1.00 0.00 C
 ATOM 14 N4 DC A 1 -3.916 5.292 -1.751 1.00 0.00 N
 ATOM 15 C5 DC A 1 -5.137 3.354 -1.017 1.00 0.00 C
 ATOM 16 C6 DC A 1 -5.080 2.018 -0.781 1.00 0.00 C
 ATOM 17 P DC A 2 -2.698 -3.257 1.572 1.00 0.00 P
 ATOM 18 OP2 DC A 2 -2.054 -2.037 2.203 1.00 0.00 O
 ATOM 19 OP1 DC A 2 -3.419 -4.230 2.463 1.00 0.00 O
 ATOM 20 O5' DC A 2 -1.579 -4.100 0.707 1.00 0.00 O
 ATOM 21 C5' DC A 2 -1.309 -3.984 -0.693 1.00 0.00 C
 ATOM 22 C4' DC A 2 -0.143 -3.068 -1.028 1.00 0.00 C
 ATOM 23 O4' DC A 2 -0.561 -1.683 -1.012 1.00 0.00 O
 ATOM 24 C3' DC A 2 1.072 -3.180 -0.067 1.00 0.00 C
 ATOM 25 O3' DC A 2 2.247 -3.319 -0.889 1.00 0.00 O
 ATOM 26 C2' DC A 2 0.978 -1.866 0.707 1.00 0.00 C
 ATOM 27 C1' DC A 2 0.499 -0.941 -0.405 1.00 0.00 C
 ATOM 28 N1 DC A 2 -0.008 0.363 0.016 1.00 0.00 N
 ATOM 29 C2 DC A 2 0.494 1.564 -0.616 1.00 0.00 C
 ATOM 30 O2 DC A 2 1.374 1.463 -1.466 1.00 0.00 O
 ATOM 31 N3 DC A 2 -0.011 2.763 -0.215 1.00 0.00 N
 ATOM 32 C4 DC A 2 -1.021 2.814 0.644 1.00 0.00 C
 ATOM 33 N4 DC A 2 -1.512 4.020 0.986 1.00 0.00 N
 ATOM 34 C5 DC A 2 -1.541 1.639 1.281 1.00 0.00 C
 ATOM 35 C6 DC A 2 -0.997 0.425 0.920 1.00 0.00 C
 ATOM 36 P DA A 3 3.759 -2.803 -0.458 1.00 0.00 P
 ATOM 37 OP1 DA A 3 4.702 -3.437 -1.442 1.00 0.00 O
 ATOM 38 OP2 DA A 3 3.964 -2.884 1.036 1.00 0.00 O
 ATOM 39 O5' DA A 3 3.630 -1.179 -0.764 1.00 0.00 O
 ATOM 40 C5' DA A 3 3.924 -0.642 -2.057 1.00 0.00 C
 ATOM 41 C4' DA A 3 5.168 0.229 -1.986 1.00 0.00 C
 ATOM 42 O4' DA A 3 4.942 1.414 -1.194 1.00 0.00 O
 ATOM 43 C3' DA A 3 6.374 -0.452 -1.333 1.00 0.00 C
 ATOM 44 O3' DA A 3 7.545 0.134 -1.901 1.00 0.00 O
 ATOM 45 C2' DA A 3 6.214 -0.090 0.157 1.00 0.00 C
 ATOM 46 C1' DA A 3 5.394 1.215 0.128 1.00 0.00 C
 ATOM 47 N9 DA A 3 4.238 1.218 1.053 1.00 0.00 N
 ATOM 48 C8 DA A 3 3.630 0.195 1.732 1.00 0.00 C
 ATOM 49 N7 DA A 3 2.535 0.562 2.395 1.00 0.00 N
 ATOM 50 C5 DA A 3 2.410 1.885 2.115 1.00 0.00 C
 ATOM 51 C6 DA A 3 1.463 2.867 2.496 1.00 0.00 C
 ATOM 52 N6 DA A 3 0.396 2.558 3.244 1.00 0.00 N
 ATOM 53 N1 DA A 3 1.621 4.144 2.070 1.00 0.00 N
 ATOM 54 C2 DA A 3 2.663 4.414 1.291 1.00 0.00 C
 ATOM 55 N3 DA A 3 3.630 3.572 0.842 1.00 0.00 N
 ATOM 56 C4 DA A 3 3.461 2.335 1.280 1.00 0.00 C
 END

COMPND CAA / D-B3LYP / PO- MODEL / NEUTRAL
 ATOM 1 O5' DC A 1 -7.203 -0.623 0.231 1.00 0.00 O
 ATOM 2 C5' DC A 1 -7.004 -1.710 -0.672 1.00 0.00 C
 ATOM 3 C4' DC A 1 -5.515 -1.952 -0.879 1.00 0.00 C
 ATOM 4 O4' DC A 1 -4.928 -0.842 -1.582 1.00 0.00 O
 ATOM 5 C3' DC A 1 -4.707 -2.101 0.420 1.00 0.00 C
 ATOM 6 O3' DC A 1 -3.588 -2.954 0.131 1.00 0.00 O
 ATOM 7 C2' DC A 1 -4.243 -0.671 0.676 1.00 0.00 C
 ATOM 8 C1' DC A 1 -3.949 -0.201 -0.751 1.00 0.00 C
 ATOM 9 N1 DC A 1 -4.019 1.240 -0.962 1.00 0.00 N
 ATOM 10 C2 DC A 1 -3.034 1.875 -1.782 1.00 0.00 C
 ATOM 11 O2 DC A 1 -2.194 1.189 -2.365 1.00 0.00 O
 ATOM 12 N3 DC A 1 -3.070 3.237 -1.871 1.00 0.00 N
 ATOM 13 C4 DC A 1 -4.041 3.924 -1.286 1.00 0.00 C
 ATOM 14 N4 DC A 1 -3.959 5.278 -1.326 1.00 0.00 N
 ATOM 15 C5 DC A 1 -5.140 3.301 -0.608 1.00 0.00 C
 ATOM 16 C6 DC A 1 -5.083 1.949 -0.483 1.00 0.00 C
 ATOM 17 P DA A 2 -2.730 -3.583 1.405 1.00 0.00 P
 ATOM 18 OP1 DA A 2 -3.542 -4.659 2.075 1.00 0.00 O
 ATOM 19 OP2 DA A 2 -2.149 -2.473 2.255 1.00 0.00 O
 ATOM 20 O5' DA A 2 -1.518 -4.302 0.553 1.00 0.00 O
 ATOM 21 C5' DA A 2 -1.206 -4.162 -0.835 1.00 0.00 C
 ATOM 22 C4' DA A 2 0.021 -3.295 -1.095 1.00 0.00 C
 ATOM 23 O4' DA A 2 -0.334 -1.895 -1.108 1.00 0.00 O
 ATOM 24 C3' DA A 2 1.146 -3.452 -0.048 1.00 0.00 C
 ATOM 25 O3' DA A 2 2.383 -3.482 -0.775 1.00 0.00 O
 ATOM 26 C2' DA A 2 0.965 -2.195 0.800 1.00 0.00 C
 ATOM 27 C1' DA A 2 0.584 -1.181 -0.269 1.00 0.00 C
 ATOM 28 N9 DA A 2 -0.053 0.019 0.209 1.00 0.00 N
 ATOM 29 C8 DA A 2 -0.983 0.119 1.220 1.00 0.00 C
 ATOM 30 N7 DA A 2 -1.420 1.346 1.412 1.00 0.00 N
 ATOM 31 C5 DA A 2 -0.728 2.097 0.477 1.00 0.00 C
 ATOM 32 C6 DA A 2 -0.742 3.466 0.166 1.00 0.00 C
 ATOM 33 N6 DA A 2 -1.545 4.347 0.837 1.00 0.00 N
 ATOM 34 N1 DA A 2 0.099 3.923 -0.780 1.00 0.00 N
 ATOM 35 C2 DA A 2 0.895 3.036 -1.400 1.00 0.00 C
 ATOM 36 N3 DA A 2 0.974 1.713 -1.227 1.00 0.00 N
 ATOM 37 C4 DA A 2 0.132 1.295 -0.275 1.00 0.00 C
 ATOM 38 P DA A 3 3.853 -3.502 -0.001 1.00 0.00 P
 ATOM 39 OP1 DA A 3 4.713 -4.500 -0.727 1.00 0.00 O
 ATOM 40 OP2 DA A 3 3.677 -3.510 1.491 1.00 0.00 O
 ATOM 41 O5' DA A 3 4.369 -1.964 -0.383 1.00 0.00 O
 ATOM 42 C5' DA A 3 4.568 -1.666 -1.765 1.00 0.00 C
 ATOM 43 C4' DA A 3 5.296 -0.342 -1.915 1.00 0.00 C
 ATOM 44 O4' DA A 3 4.465 0.747 -1.470 1.00 0.00 O
 ATOM 45 C3' DA A 3 6.596 -0.236 -1.106 1.00 0.00 C
 ATOM 46 O3' DA A 3 7.469 0.629 -1.835 1.00 0.00 O
 ATOM 47 C2' DA A 3 6.120 0.407 0.197 1.00 0.00 C
 ATOM 48 C1' DA A 3 4.998 1.335 -0.287 1.00 0.00 C
 ATOM 49 N9 DA A 3 3.934 1.516 0.683 1.00 0.00 N
 ATOM 50 C8 DA A 3 3.212 0.532 1.330 1.00 0.00 C
 ATOM 51 N7 DA A 3 2.235 0.999 2.075 1.00 0.00 N
 ATOM 52 C5 DA A 3 2.304 2.371 1.899 1.00 0.00 C
 ATOM 53 C6 DA A 3 1.515 3.440 2.368 1.00 0.00 C
 ATOM 54 N6 DA A 3 0.434 3.256 3.180 1.00 0.00 N
 ATOM 55 N1 DA A 3 1.849 4.697 2.010 1.00 0.00 N
 ATOM 56 C2 DA A 3 2.893 4.870 1.184 1.00 0.00 C
 ATOM 57 N3 DA A 3 3.690 3.946 0.636 1.00 0.00 N
 ATOM 58 C4 DA A 3 3.348 2.712 1.031 1.00 0.00 C
 END

COMPND CAA / D-B3LYP / PO- MODEL / SINGLY IONIZED
 ATOM 1 O5' DC A 1 -7.105 -0.754 -0.217 1.00 0.00 O
 ATOM 2 C5' DC A 1 -6.688 -1.783 -1.115 1.00 0.00 C
 ATOM 3 C4' DC A 1 -5.173 -1.909 -1.089 1.00 0.00 C
 ATOM 4 O4' DC A 1 -4.565 -0.750 -1.694 1.00 0.00 O
 ATOM 5 C3' DC A 1 -4.567 -2.015 0.319 1.00 0.00 C
 ATOM 6 O3' DC A 1 -3.373 -2.801 0.218 1.00 0.00 O
 ATOM 7 C2' DC A 1 -4.226 -0.561 0.640 1.00 0.00 C

ATOM 8 C1' DC A 1 -3.788 -0.037 -0.733 1.00 0.00 C
 ATOM 9 N1 DC A 1 -4.016 1.400 -0.928 1.00 0.00 N
 ATOM 10 C2 DC A 1 -2.982 2.235 -1.424 1.00 0.00 C
 ATOM 11 O2 DC A 1 -1.894 1.727 -1.751 1.00 0.00 O
 ATOM 12 N3 DC A 1 -3.209 3.569 -1.515 1.00 0.00 N
 ATOM 13 C4 DC A 1 -4.420 4.063 -1.242 1.00 0.00 C
 ATOM 14 N4 DC A 1 -4.578 5.395 -1.319 1.00 0.00 N
 ATOM 15 C5 DC A 1 -5.530 3.226 -0.894 1.00 0.00 C
 ATOM 16 C6 DC A 1 -5.275 1.899 -0.749 1.00 0.00 C
 ATOM 17 P DA A 2 -2.753 -3.433 1.619 1.00 0.00 P
 ATOM 18 OP1 DA A 2 -3.774 -4.339 2.253 1.00 0.00 O
 ATOM 19 OP2 DA A 2 -2.129 -2.335 2.458 1.00 0.00 O
 ATOM 20 O5' DA A 2 -1.531 -4.330 0.976 1.00 0.00 O
 ATOM 21 C5' DA A 2 -1.229 -4.503 -0.409 1.00 0.00 C
 ATOM 22 C4' DA A 2 -0.043 -3.656 -0.843 1.00 0.00 C
 ATOM 23 O4' DA A 2 -0.435 -2.264 -0.933 1.00 0.00 O
 ATOM 24 C3' DA A 2 1.168 -3.708 0.127 1.00 0.00 C
 ATOM 25 O3' DA A 2 2.357 -3.768 -0.685 1.00 0.00 O
 ATOM 26 C2' DA A 2 0.989 -2.406 0.903 1.00 0.00 C
 ATOM 27 C1' DA A 2 0.543 -1.500 -0.233 1.00 0.00 C
 ATOM 28 N9 DA A 2 -0.021 -0.232 0.177 1.00 0.00 N
 ATOM 29 C8 DA A 2 -0.906 0.012 1.185 1.00 0.00 C
 ATOM 30 N7 DA A 2 -1.183 1.316 1.350 1.00 0.00 N
 ATOM 31 C5 DA A 2 -0.419 1.928 0.422 1.00 0.00 C
 ATOM 32 C6 DA A 2 -0.267 3.292 0.030 1.00 0.00 C
 ATOM 33 N6 DA A 2 -1.020 4.245 0.562 1.00 0.00 N
 ATOM 34 N1 DA A 2 0.663 3.607 -0.904 1.00 0.00 N
 ATOM 35 C2 DA A 2 1.348 2.616 -1.461 1.00 0.00 C
 ATOM 36 N3 DA A 2 1.229 1.274 -1.259 1.00 0.00 N
 ATOM 37 C4 DA A 2 0.351 0.992 -0.327 1.00 0.00 C
 ATOM 38 P DA A 3 3.817 -3.061 -0.333 1.00 0.00 P
 ATOM 39 OP1 DA A 3 4.810 -3.691 -1.269 1.00 0.00 O
 ATOM 40 OP2 DA A 3 4.036 -2.971 1.156 1.00 0.00 O
 ATOM 41 O5' DA A 3 3.530 -1.495 -0.799 1.00 0.00 O
 ATOM 42 C5' DA A 3 3.790 -1.033 -2.124 1.00 0.00 C
 ATOM 43 C4' DA A 3 4.903 0.005 -2.099 1.00 0.00 C
 ATOM 44 O4' DA A 3 4.484 1.209 -1.428 1.00 0.00 O
 ATOM 45 C3' DA A 3 6.159 -0.458 -1.352 1.00 0.00 C
 ATOM 46 O3' DA A 3 7.268 0.226 -1.937 1.00 0.00 O
 ATOM 47 C2' DA A 3 5.893 0.011 0.087 1.00 0.00 C
 ATOM 48 C1' DA A 3 5.005 1.252 -0.106 1.00 0.00 C
 ATOM 49 N9 DA A 3 3.913 1.362 0.863 1.00 0.00 N
 ATOM 50 C8 DA A 3 3.185 0.377 1.504 1.00 0.00 C
 ATOM 51 N7 DA A 3 2.182 0.841 2.220 1.00 0.00 N
 ATOM 52 C5 DA A 3 2.240 2.211 2.034 1.00 0.00 C
 ATOM 53 C6 DA A 3 1.485 3.289 2.545 1.00 0.00 C
 ATOM 54 N6 DA A 3 0.437 3.112 3.383 1.00 0.00 N
 ATOM 55 N1 DA A 3 1.826 4.547 2.190 1.00 0.00 N
 ATOM 56 C2 DA A 3 2.869 4.716 1.363 1.00 0.00 C
 ATOM 57 N3 DA A 3 3.663 3.788 0.814 1.00 0.00 N
 ATOM 58 C4 DA A 3 3.309 2.554 1.192 1.00 0.00 C
 END