

Effects of Sodium Cationization vs. Protonation on the Conformations and N-Glycosidic

Bond Stabilities of Sodium Cationized Uridine and 2'-Deoxyuridine:

Solution Conformation of [Urd+Na]⁺ is Preserved Upon ESI

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Table S1. Mass List of Protonated and Sodium Cationized Urd and dUrd

m/z	Symbol	[Urd+H] ⁺	[dUrd+H] ⁺	[Urd+Na] ⁺	[dUrd+Na] ⁺
267.0	•	–	–	[Urd+Na] ⁺	–
251.0	•	–	–	–	[dUrd+Na] ⁺
245.0	•	[Urd+H] ⁺	–	–	–
229.0	•	–	[dUrd+H] ⁺	–	–
227.0	•	[Urd–W+H] ⁺	–	–	–
211.0	•	–	[dUrd–W+H] ⁺	–	–
203.0	•	[Urd–K+H] ⁺	–	–	–
193.0	•	–	[dUrd–2W+H] ⁺	–	–
185.0	•	[Urd–K–W+H] ⁺	–	–	–
155.0	•	–	–	[Urd–Ura+Na] ⁺	–
139.0	•	–	–	–	[dUrd–Ura+Na] ⁺
135.0	•	–	–	[Ura+Na] ⁺	[Ura+Na] ⁺
131.0	•	–	[Ura+W+H] ⁺	–	–
117.0	•	–	[dUrd–Ura+H] ⁺	–	–
113.0	•	[Ura+H] ⁺	[Ura+H] ⁺	–	–
99.0	•	–	[dUrd–Ura–W+H] ⁺	–	–
81.0	•	–	[dUrd–Ura–2W+H] ⁺	–	–

Table S2. Relative Gibbs Free Energies of the B3LYP/6-311+G(d,p) Optimized Structures of the T1(O2O4'O5') and B1(O2O2') Conformers of [Urd+Na]⁺ and Their Water Adducts ([Urd+Na]⁺•nW, n=1, 2) Calculated Using Different Computational Approaches.^a

[Urd+Na] ⁺	B3LYP		MP2(full)	
	6-311+G(2d,2p)	def2-TZVPPD	6-311+G(2d,2p)	def2-TZVPPD
T1(O2O4'O5')	0.0	0.0	0.0	0.0
B1(O2O2')	6.1	4.5	9.5	9.6
T1(O2O4'O5')•W	1.6	3.1	0.0	0.0
B1(O2O2')•W	0.0	0.0	4.1	4.1
T1(O2O4'O5')•2W	3.9	13.4	1.2	0.0
B1(O2O2')•2W	0.0	0.0	0.0	0.0

^aAll values are given in kJ/mol.

Table S3. Relative Gibbs Free Energies of the B3LYP/6-311+G(d,p) Optimized Structures of the T1, T2, T3 and T4(O2O4'O5') Conformers of [dUrd+Na]⁺ Calculated Using Different Computational Approaches.^a

[dUrd+Na] ⁺	B3LYP		MP2(full)	
	6-311+G(2d,2p)	def2-TZVPPD	6-311+G(2d,2p)	def2-TZVPPD
T1(O2O4'O5')	0.0	0.0	0.0	0.0
T2(O2O4'O5')	3.5	3.3	2.9	3.7
T3(O2O4'O5')	5.5	5.0	1.5	1.9
T4(O2O4'O5')	7.5	7.2	6.1	4.9

^aAll values are given in kJ/mol.

Figure Captions

Figure S1. Designations for nucleobase orientation and pseudorotation phase angle (P) of nucleosides. The pseudorotation phase angle, P , is calculated using eq (S1). E and T forms alternate every 18° .

$$\tan P = \frac{(v_4+v_1)-(v_3+v_0)}{2 \times v_2 \times (\sin 36^\circ + \sin 72^\circ)} \quad (\text{S1})$$

The angles, v_0 , v_1 , v_2 , v_3 and v_4 , represent the $\angle \text{C4'O4'C1'C2'}$, $\angle \text{O4'C1'C2'C3'}$, $\angle \text{C1'C2'C3'C4'}$, $\angle \text{C2'C3'C4'O4'}$ and $\angle \text{C3'C4'O4'C1'}$, respectively. In the upper section of the diagram, $v_2 > 0$, whereas in the lower section of the diagram, $v_2 < 0$. The combination of P and v_2 are used to identify the sugar puckering. If $v_2 > 0$, $P = P$; if $v_2 < 0$ $P \rightarrow P+180^\circ$.

Figure S2. The mass spectra of MS³ experiments of $[\text{dUrd-W+H}]^+$, $[\text{dUrd-2W+H}]^+$, $[\text{dUrd-Ura+H}]^+$ and $[\text{Urd-K+H}]^+$.

Figure S3. Stable low-energy conformers of $[\text{Urd+Na}]^+$. The Na^+ binding modes, orientations of uracil, sugar puckering, and the relative 298 K Gibbs free energies at the B3LYP/6-311+G(2d,2p)//B3LYP/6-311+G(d,p) level of theory are also listed for each structure.

Figure S4. Stable low-energy conformers of $[\text{dUrd+Na}]^+$. The Na^+ binding modes, orientations of uracil, sugar puckering, and the relative 298 K Gibbs free energies at the B3LYP/6-311+G(2d,2p)//B3LYP/6-311+G(d,p) level of theory are also listed for each structure.

Figure S5. Comparison of the experimental IRMPD action spectrum of $[\text{Urd+Na}]^+$ with the B3LYP/6-311+G(d,p) optimized structures and calculated linear IR spectra for *syn* oriented O2 binding conformers with spectral misalignments shaded in red. The B3LYP/6-311+G(2d,2p) relative Gibbs free energies at 298 K is also shown.

Figure S6. Comparison of the experimental IRMPD action spectrum of $[\text{Urd}+\text{Na}]^+$ with the B3LYP/6-311+G(d,p) optimized structures and calculated linear IR spectra for *anti* oriented O2 binding conformers with spectral misalignments shaded in red. The B3LYP/6-311+G(2d,2p) relative Gibbs free energies at 298 K is also shown.

Figure S7. Comparison of the experimental IRMPD action spectrum of $[\text{Urd}+\text{Na}]^+$ with the B3LYP/6-311+G(d,p) optimized structures and calculated linear IR spectra for O4 binding conformers with spectral misalignments shaded in red. The B3LYP/6-311+G(2d,2p) relative Gibbs free energies at 298 K is also shown.

Figure S8. Comparison of the experimental IRMPD action spectrum of $[\text{Urd}+\text{Na}]^+$ with the B3LYP/6-311+G(d,p) optimized structures and calculated linear IR spectra for sugar binding conformers with spectral misalignments shaded in red. The B3LYP/6-311+G(2d,2p) relative Gibbs free energies at 298 K is also shown.

Figure S9. Comparison of the experimental IRMPD action spectrum of $[\text{Urd}+\text{Na}]^+$ with the B3LYP/6-311+G(d,p) optimized structures and calculated linear IR spectra for tautomeric conformers with spectral misalignments shaded in red. The B3LYP/6-311+G(2d,2p) relative Gibbs free energies at 298 K is also shown.

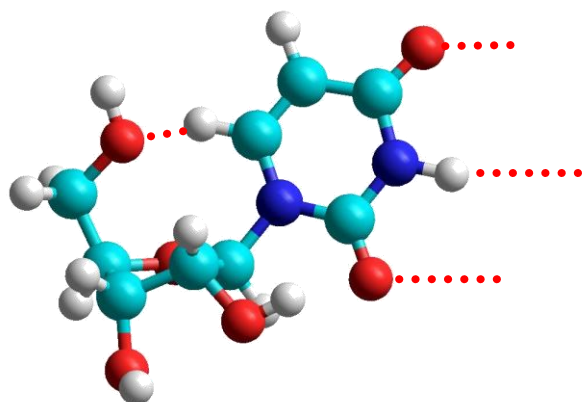
Figure S10. Comparison of the experimental IRMPD action spectrum of $[\text{dUrd}+\text{Na}]^+$ with the B3LYP/6-311+G(d,p) optimized structures and calculated linear IR spectra for T(O2O4'O5') conformers with spectral misalignments shaded in red. The B3LYP/6-311+G(2d,2p) relative Gibbs free energies at 298 K is also shown.

Figure S11. Comparison of the experimental IRMPD action spectrum of [dUrd+Na]⁺ with the B3LYP/6-311+G(d,p) optimized structures and calculated linear IR spectra for bidentate and monodentate O2 binding conformers with spectral misalignments shaded in red. The B3LYP/6-311+G(2d,2p) relative Gibbs free energies at 298 K is also shown.

Figure S12. Comparison of the experimental IRMPD action spectrum of [dUrd+Na]⁺ with the B3LYP/6-311+G(d,p) optimized structures and calculated linear IR spectra for O4 binding conformers with spectral misalignments shaded in red. The B3LYP/6-311+G(2d,2p) relative Gibbs free energies at 298 K is also shown.

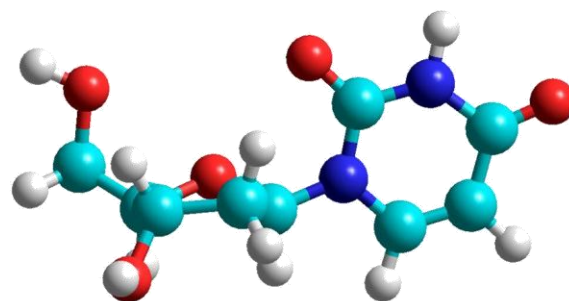
Figure S13. Comparison of the experimental IRMPD action spectrum of [dUrd+Na]⁺ with the B3LYP/6-311+G(d,p) optimized structures and calculated linear IR spectra for sugar binding conformers with spectral misalignments shaded in red. The B3LYP/6-311+G(2d,2p) relative Gibbs free energies at 298 K is also shown.

Figure S14. Comparison of the experimental IRMPD action spectrum of [dUrd+Na]⁺ with the B3LYP/6-311+G(d,p) optimized structures and calculated linear IR spectra for tautomeric conformers with spectral misalignments shaded in red. The B3LYP/6-311+G(2d,2p) relative Gibbs free energies at 298 K is also shown.



anti orientation

$$\angle C2N1C1'O4' = 90^\circ \text{ to } 270^\circ$$



syn orientation

$$\angle C2N1C1'O4' = -90^\circ \text{ to } 90^\circ$$

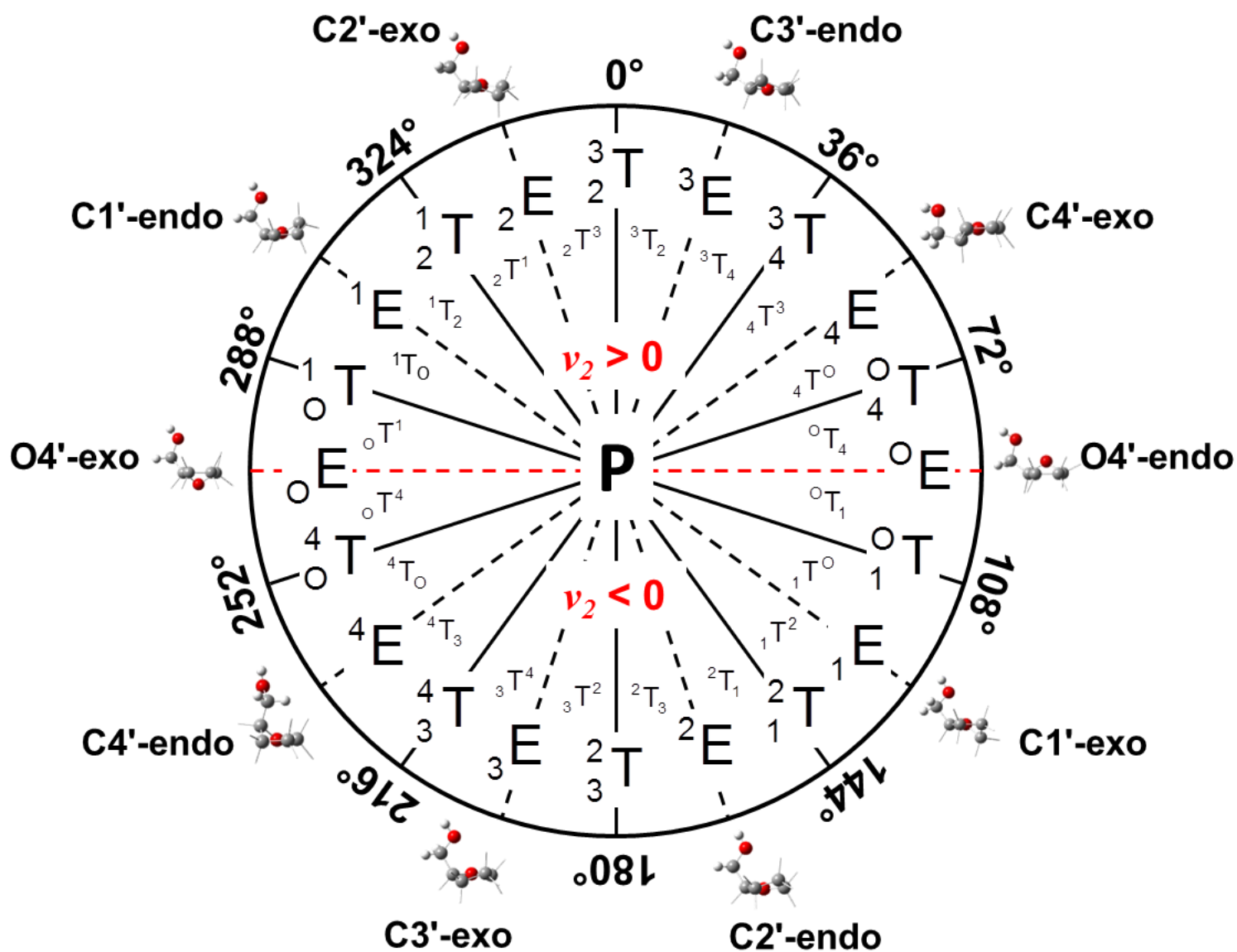
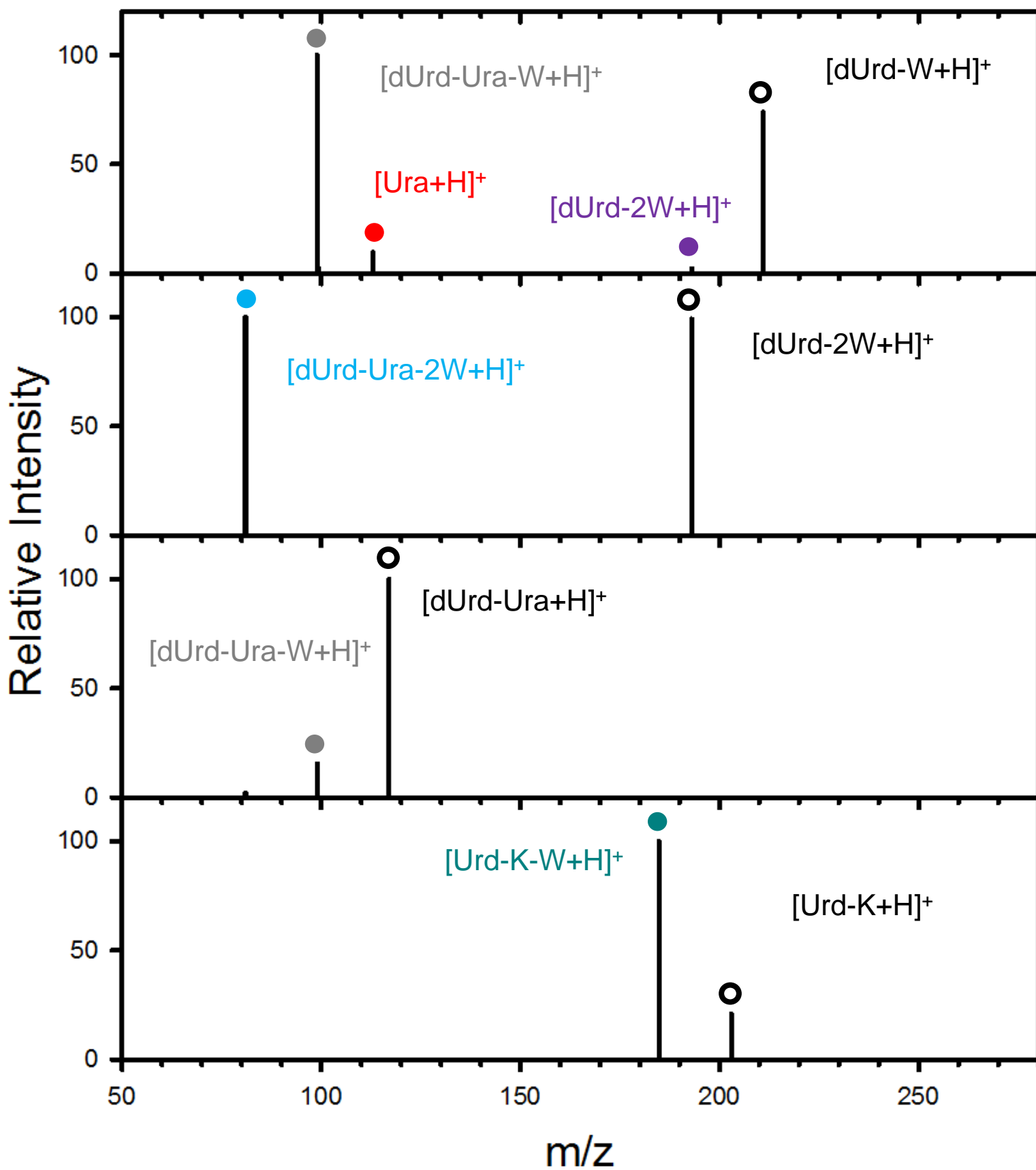
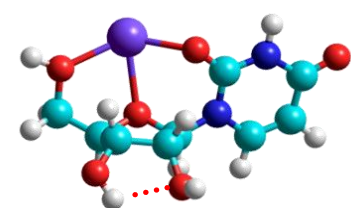
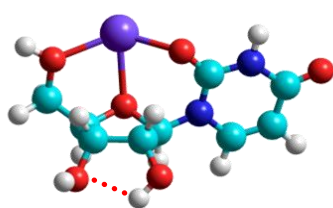


Figure S2.

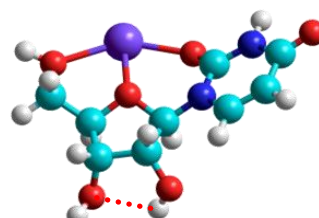




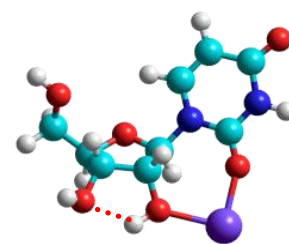
T1(O2O4'O5')
syn, O4'-endo (${}^{\circ}T_4$)
 0.0 kJ/mol



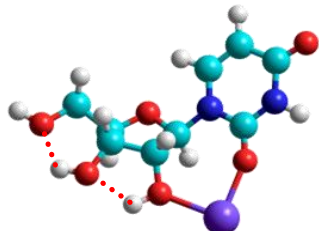
T2(O2O4'O5')
syn, O4'-endo (${}^{\circ}T_1$)
 1.4 kJ/mol



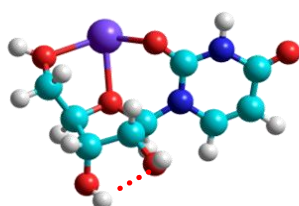
T3(O2O4'O5')
syn, O4'-endo (${}^{\circ}T_1$)
 2.8 kJ/mol



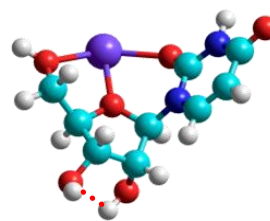
B1(O2O2')
anti, C4'-exo (${}_4T^0$)
 6.1 kJ/mol



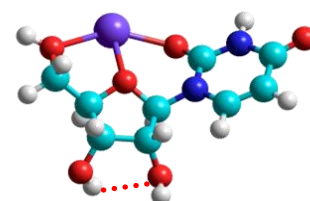
B2(O2O2')
anti, C4'-exo (${}_4T^0$)
 7.1 kJ/mol



T4(O2O4'O5')
syn, C2'-endo (2T_1)
 8.0 kJ/mol



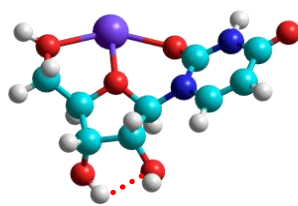
T5(O2O4'O5')
syn, C3'-endo (3T_2)
 8.5 kJ/mol



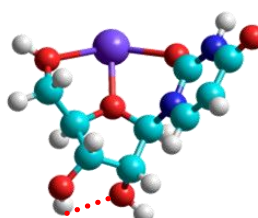
T6(O2O4'O5')
syn, O4'-endo (${}^{\circ}T_4$)
 9.1 kJ/mol



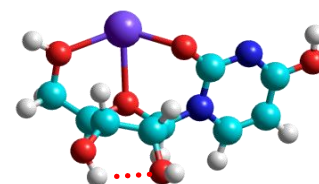
B3(O2O2')
anti, C3'-exo (${}_3T^4$)
 9.2 kJ/mol



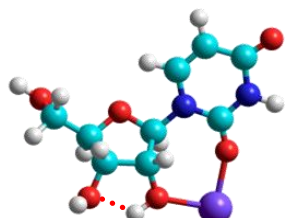
T7(O2O4'O5')
syn, C1'-exo (${}_1T^0$)
 10.2 kJ/mol



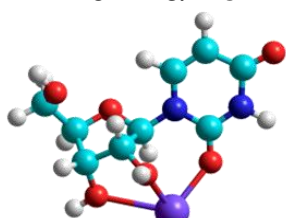
T8(O2O4'O5')
syn, C2'-exo (${}_2T^3$)
 11.7 kJ/mol



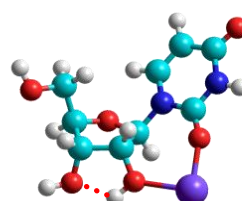
T1(O2O4'O5')t4
syn, O4'-endo (${}^{\circ}T_4$)
 16.6 kJ/mol



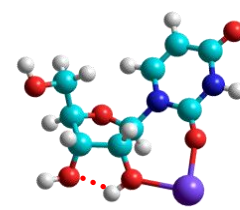
B4(O2O2')
anti, C4'-exo (${}_4T^0$)
 17.7 kJ/mol



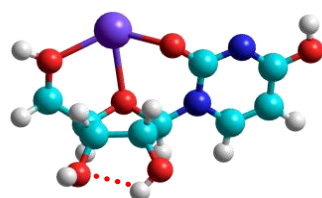
T1(O2O2'O3')
anti, C2'-endo (2T_1)
 18.2 kJ/mol



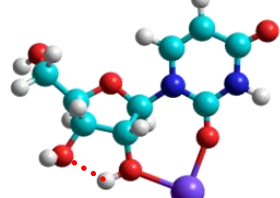
B5(O2O2')
anti, C4'-endo (4T_3)
 18.5 kJ/mol



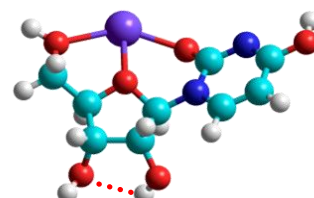
B6(O2O2')
anti, C4'-endo (4T_3)
 18.6 kJ/mol



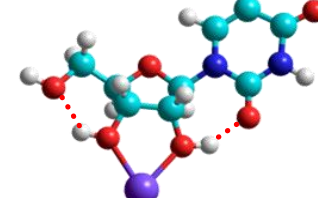
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syn, O4'-endo (${}^{\circ}T_1$)
 19.0 kJ/mol



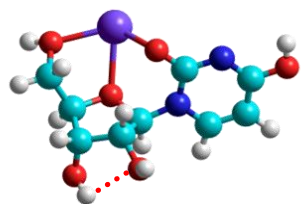
B7(O2O2')
anti, C4'-exo (${}_4T^0$)
 20.5 kJ/mol



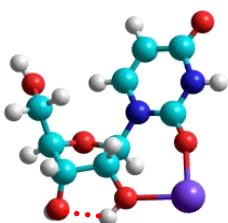
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syn, O4'-endo (${}^{\circ}T_1$)
 21.8 kJ/mol



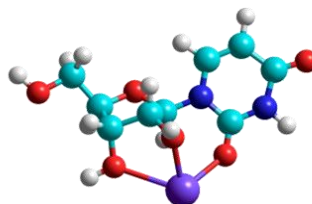
B1(O2'O3')
anti, O4'-endo (${}^{\circ}T_4$)
 22.9 kJ/mol



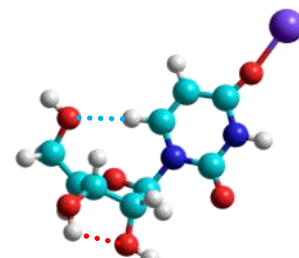
T4(O2O4'O5')t4
syn, C2'-endo (2T_1)
25.9 kJ/mol



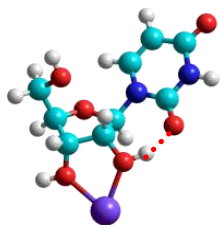
B8(O2O2')
anti, C4'-endo (4T_3)
26.7 kJ/mol



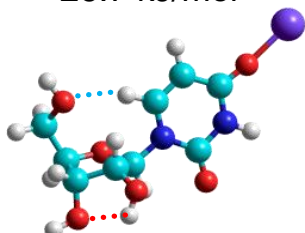
T2(O2O2'O3')
anti, C1'-exo (${}_1T_2$)
26.9 kJ/mol



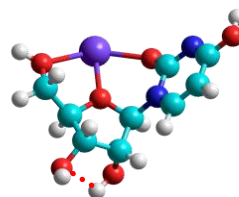
M1(O4)
anti, C3'-endo (3T_2)
27.4 kJ/mol



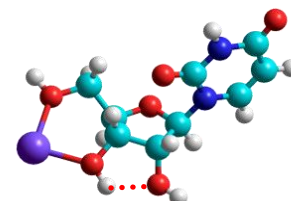
B2(O2'O3')
anti, C3'-exo (${}_3T_2$)
28.5 kJ/mol



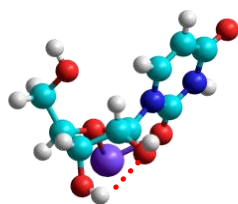
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28.6 kJ/mol



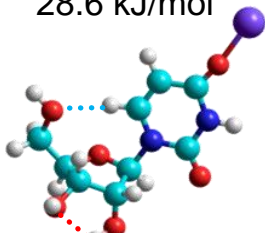
T5(O2O4'O5')t4
syn, C3'-endo (3T_2)
29.0 kJ/mol



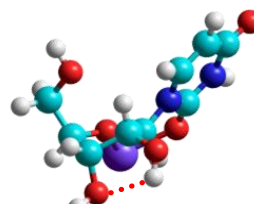
B1(O3'O5')
syn, C3'-endo (3T_4)
29.4 kJ/mol



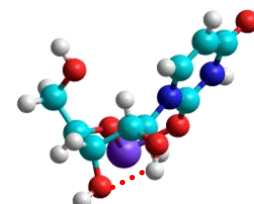
B1(O2O4')
syn, C3'-exo (${}_3T_2$)
30.9 kJ/mol



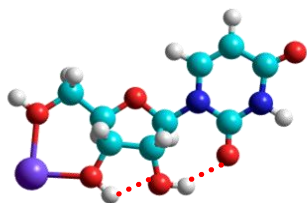
M3(O4)
anti, C3'-endo (3T_2)
31.9 kJ/mol



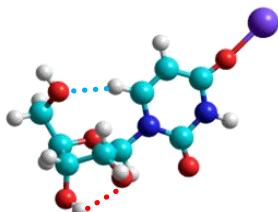
B2(O2O4')
syn, C3'-exo (${}_3T_2$)
32.8 kJ/mol



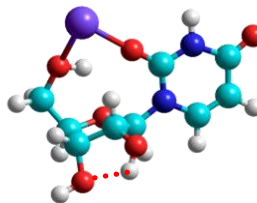
B3(O2O4')
syn, C3'-exo (${}_3T_2$)
32.9 kJ/mol



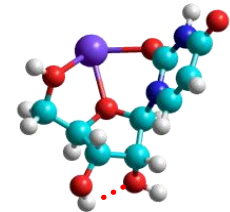
B2(O3'O5')
anti, O4'-endo (0T_4)
33.5 kJ/mol



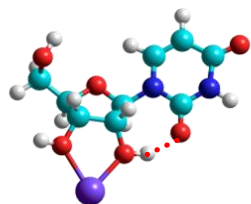
M4(O4)
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34.3 kJ/mol



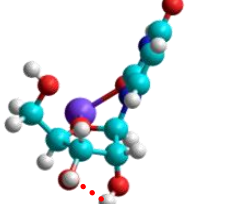
B1(O2O5')
syn, C2'-endo (2T_1)
34.5 kJ/mol



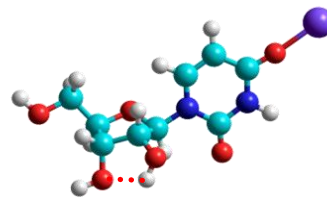
T9(O2O4'O5')
syn, C3'-endo (3T_2)
34.6 kJ/mol



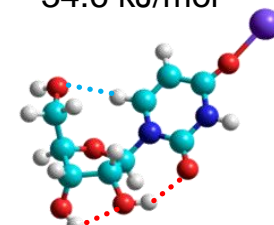
B3(O2'O3')
anti, O4'-endo (0T_4)
34.7 kJ/mol



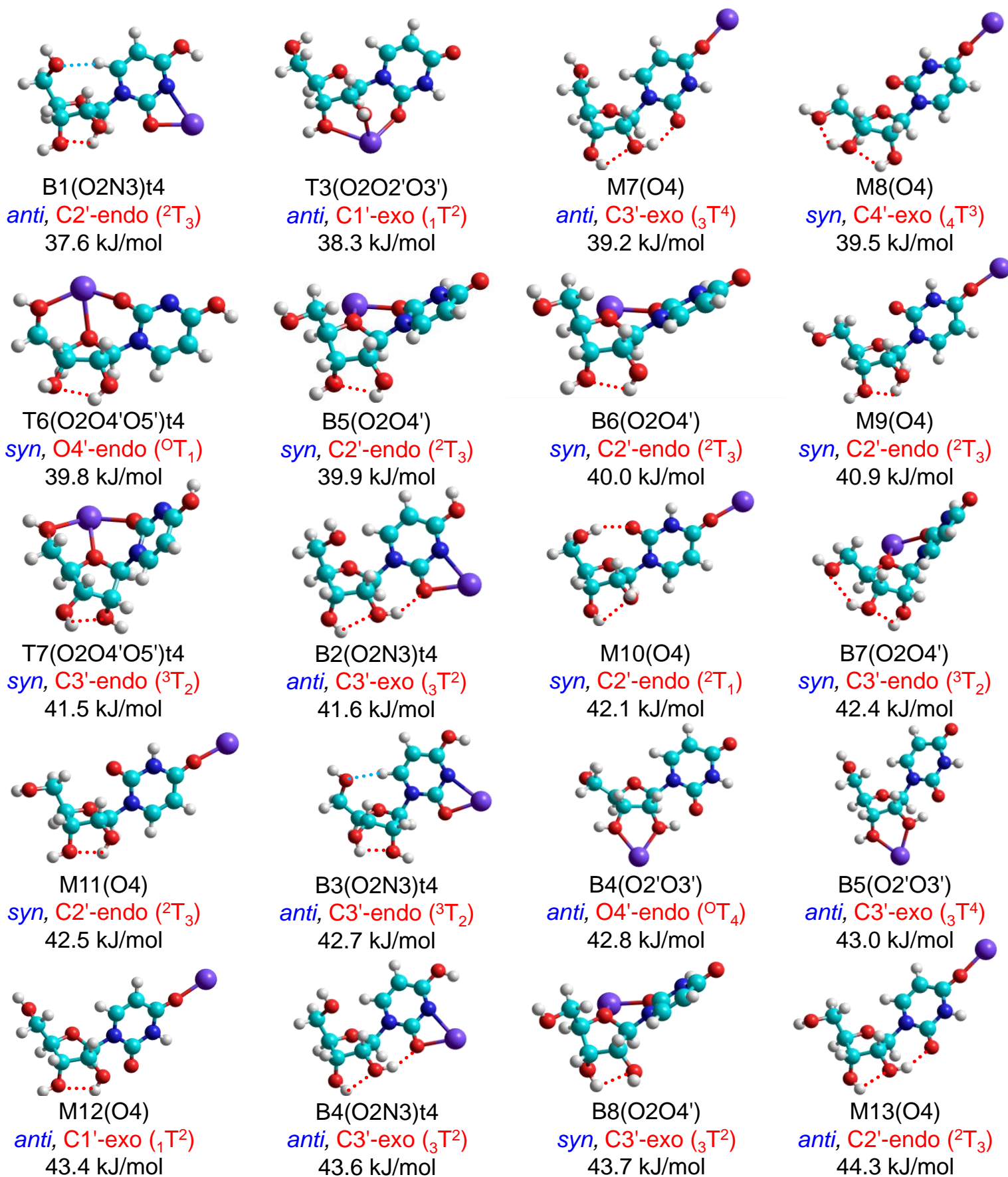
B4(O2O4')
syn, C2'-exo (${}_2T_3$)
35.1 kJ/mol

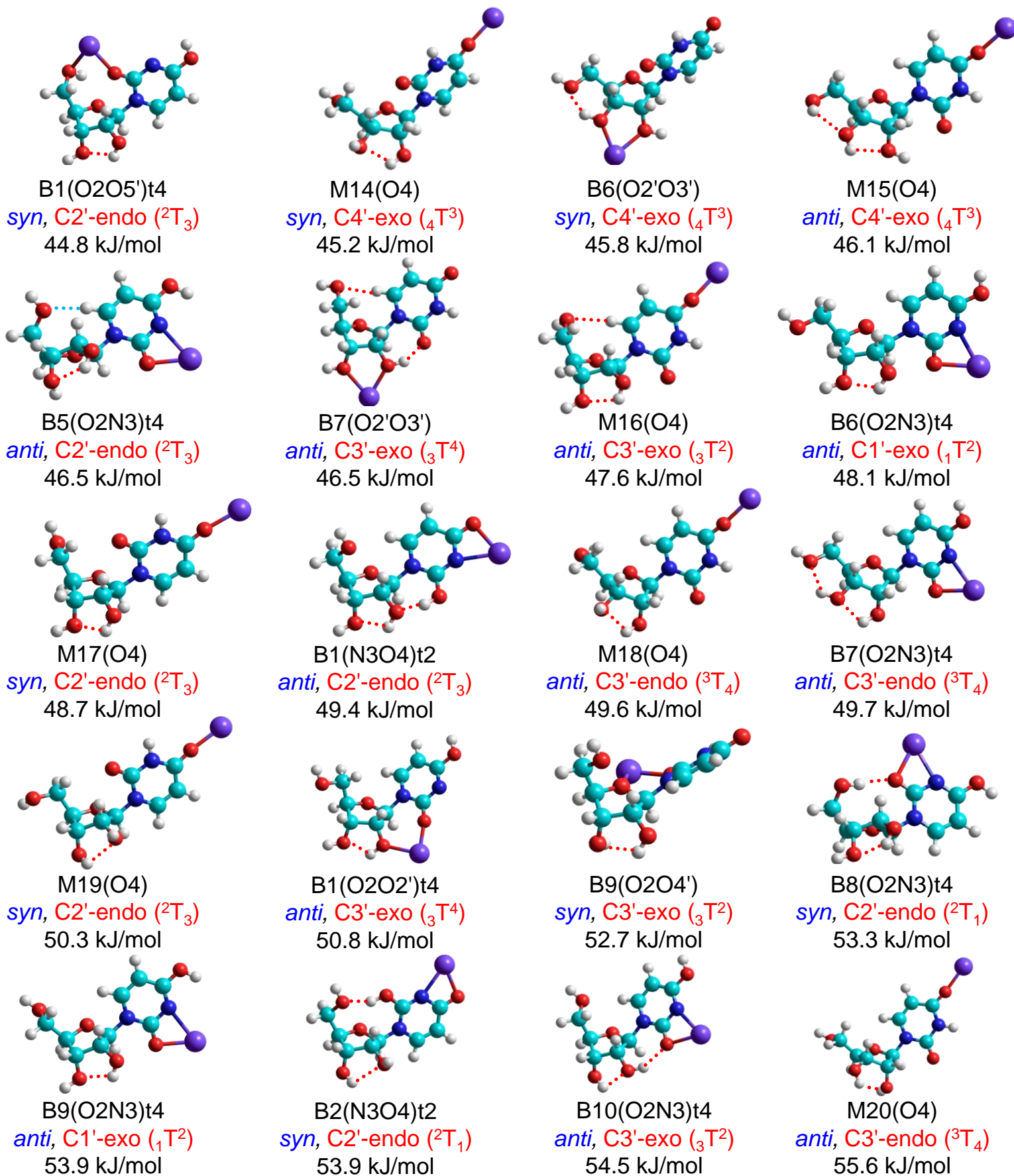


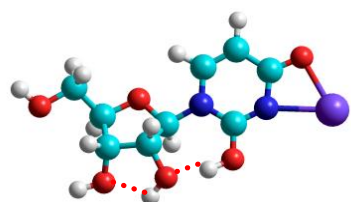
M5(O4)
anti, C1'-exo (${}_1T_2$)
36.8 kJ/mol



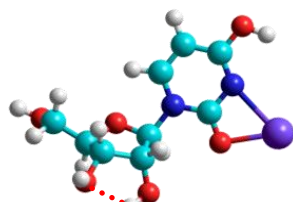
M6(O4)
anti, C3'-exo (${}_3T_4$)
37.5 kJ/mol



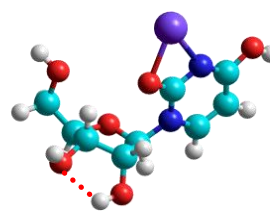




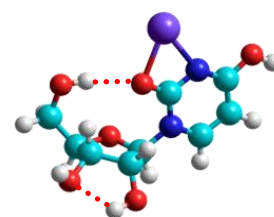
B3(N3O4)t2
anti, C1'-exo ($_1T^2$)
56.3 kJ/mol



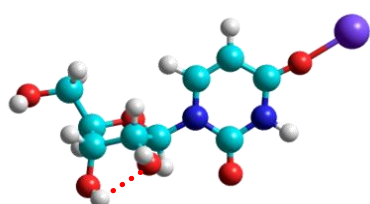
B11(O2N3)t4
anti, C4'-exo ($_4T^3$)
56.3 kJ/mol



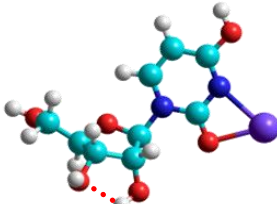
B12(O2N3)t4
syn, C4'-exo ($_4T^3$)
56.5 kJ/mol



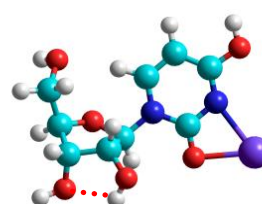
B13(O2N3)t4
syn, C4'-exo ($_4T^0$)
56.6 kJ/mol



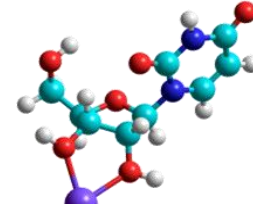
M21(O4)
anti, C2'-endo (2T_1)
56.8 kJ/mol



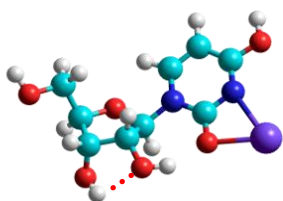
B14(O2N3)t4
anti, C4'-exo ($_4T^3$)
56.9 kJ/mol



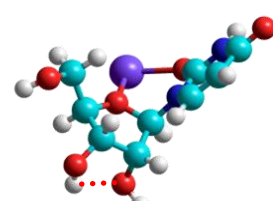
B15(O2N3)t4
anti, C3'-exo ($_3T^2$)
58.0 kJ/mol



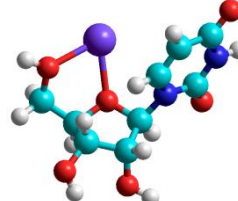
B8(O2'O3')
syn, C4'-exo ($_4T^3$)
59.0 kJ/mol



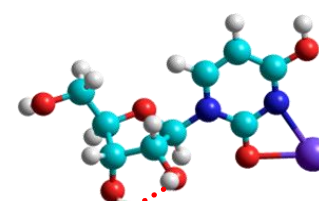
B16(O2N3)t4
anti, C2'-endo (2T_3)
59.2 kJ/mol



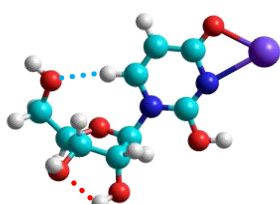
B10(O2O4')
syn, C2'-exo ($_2T^1$)
61.2 kJ/mol



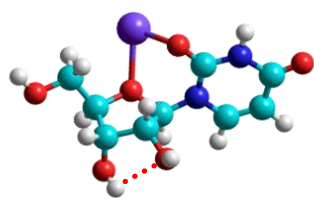
B1(O4'O5')
anti, C4'-exo ($_4T^3$)
62.1 kJ/mol



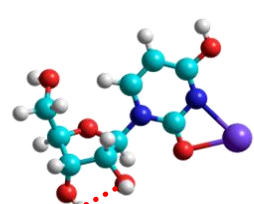
B17(O2N3)t4
anti, C2'-endo (2T_1)
62.3 kJ/mol



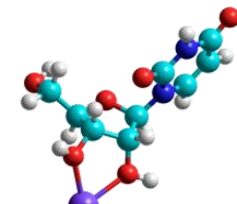
B4(N3O4)t2
anti, C3'-endo (3T_4)
63.1 kJ/mol



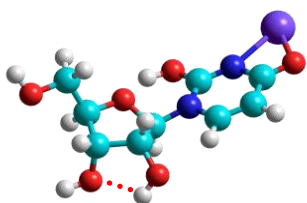
B11(O2O4')
syn, C1'-exo ($_1T^2$)
64.3 kJ/mol



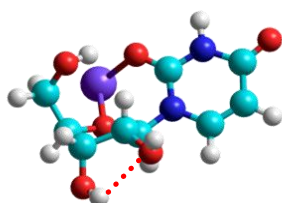
B18(O2N3)t4
anti, C3'-exo ($_3T^2$)
64.6 kJ/mol



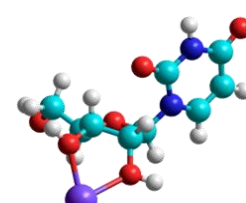
B9(O2'O3')
syn, C4'-exo ($_4T^3$)
64.7 kJ/mol



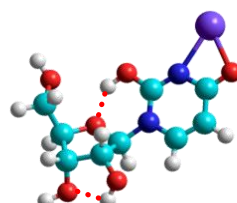
B5(N3O4)t2
syn, C1'-exo ($_1T^2$)
64.9 kJ/mol



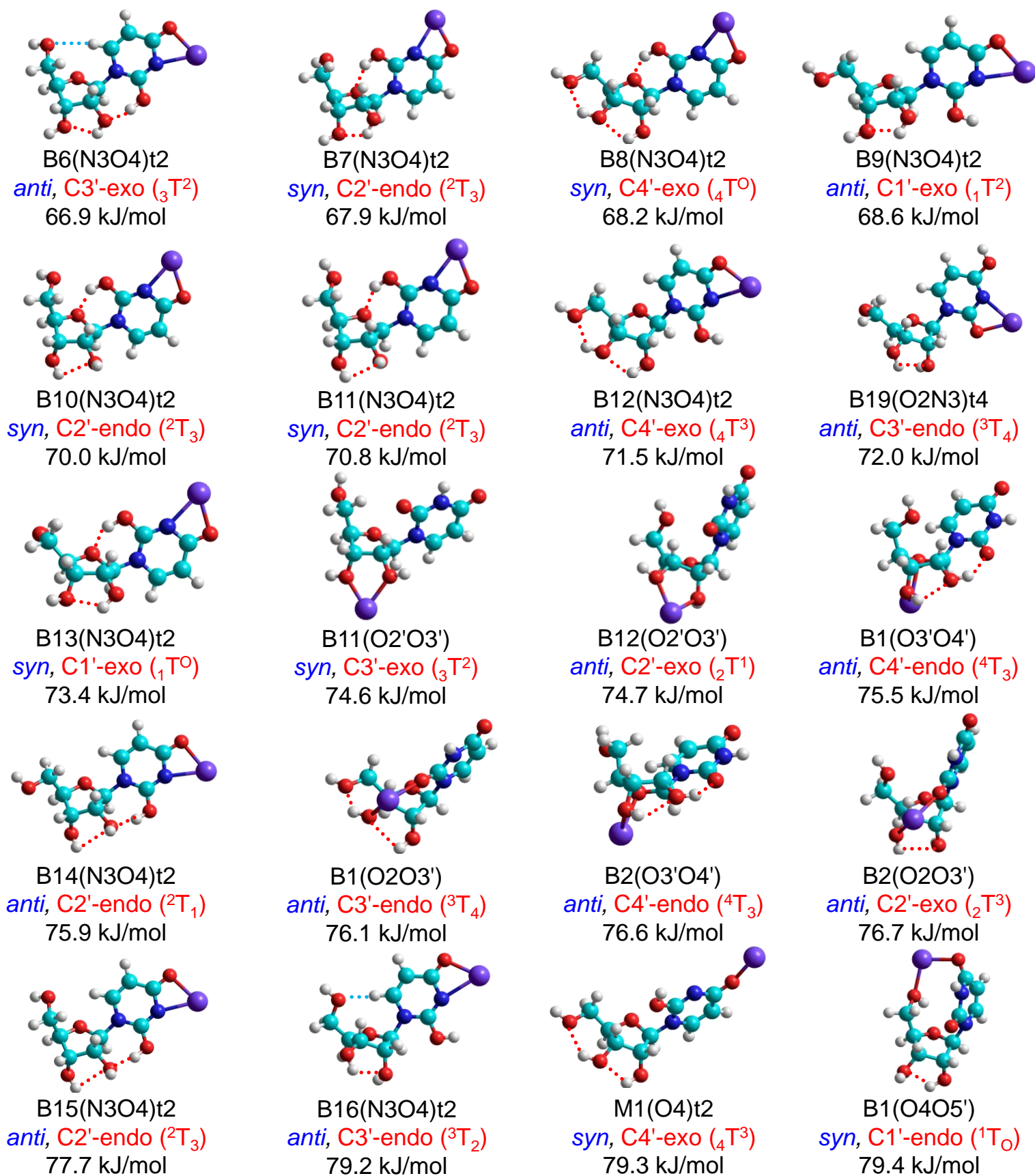
B12(O2O4')
syn, C2'-endo (2T_3)
65.6 kJ/mol

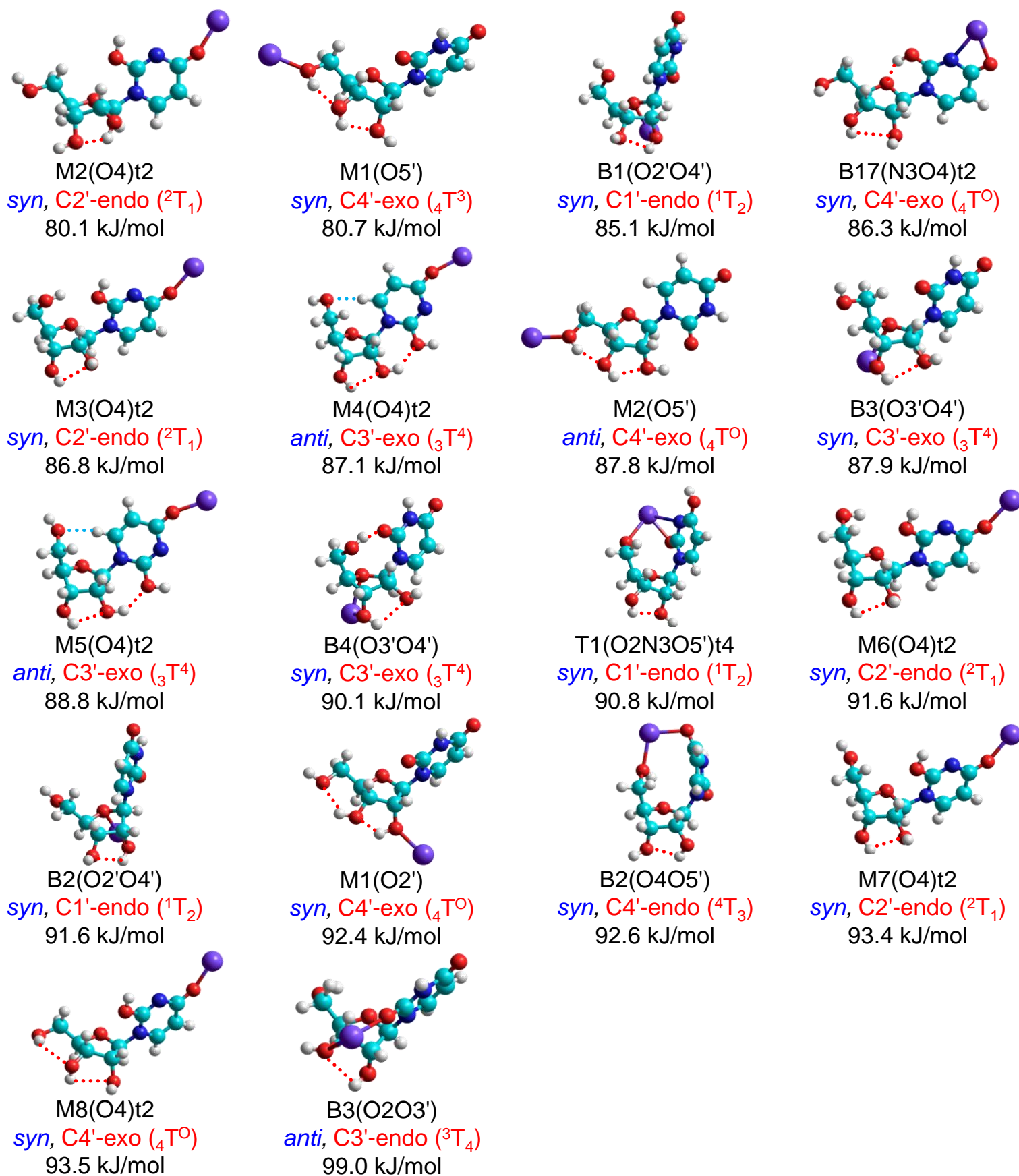


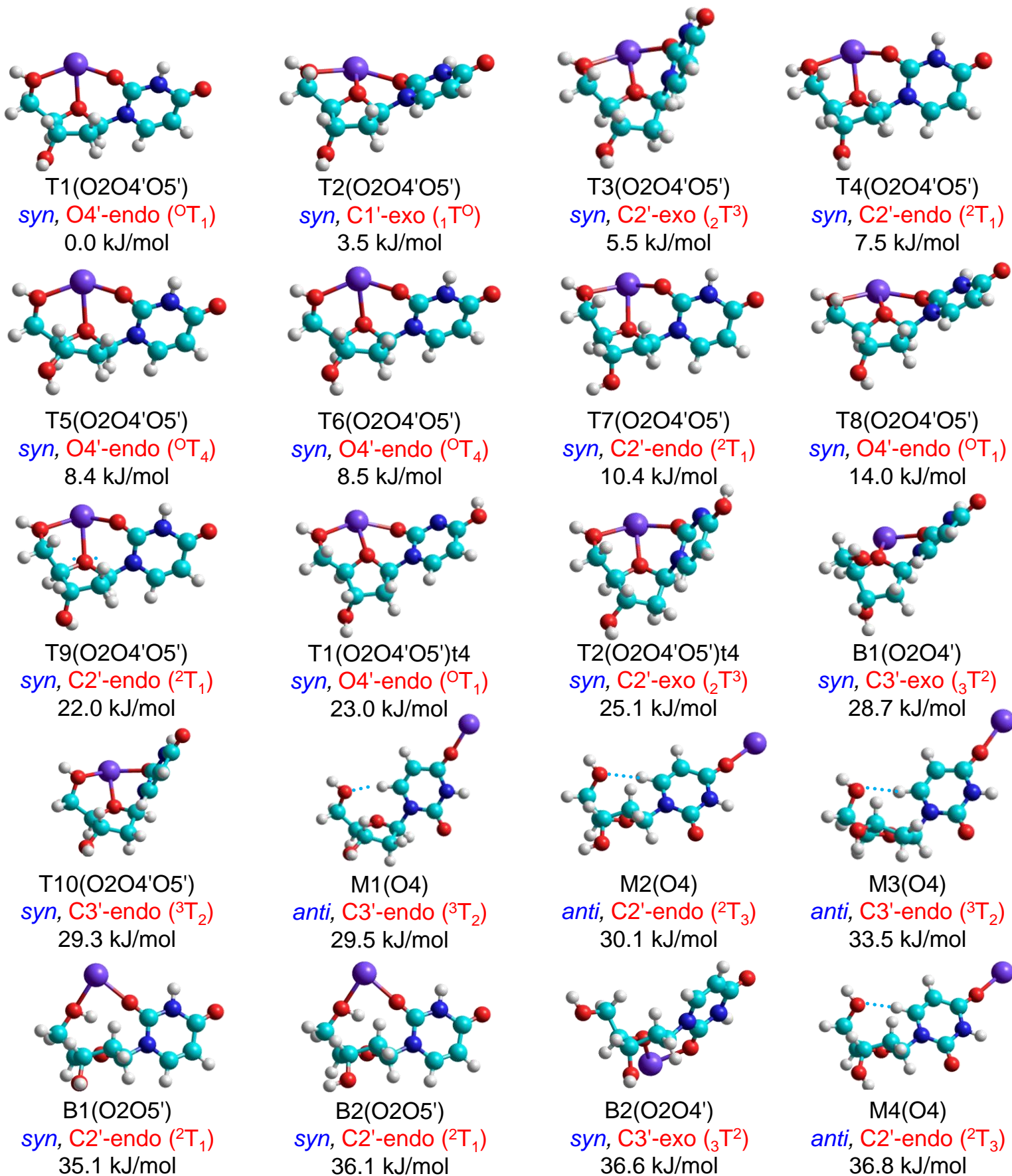
B10(O2'O3')
syn, C4'-exo ($_4T^3$)
65.8 kJ/mol

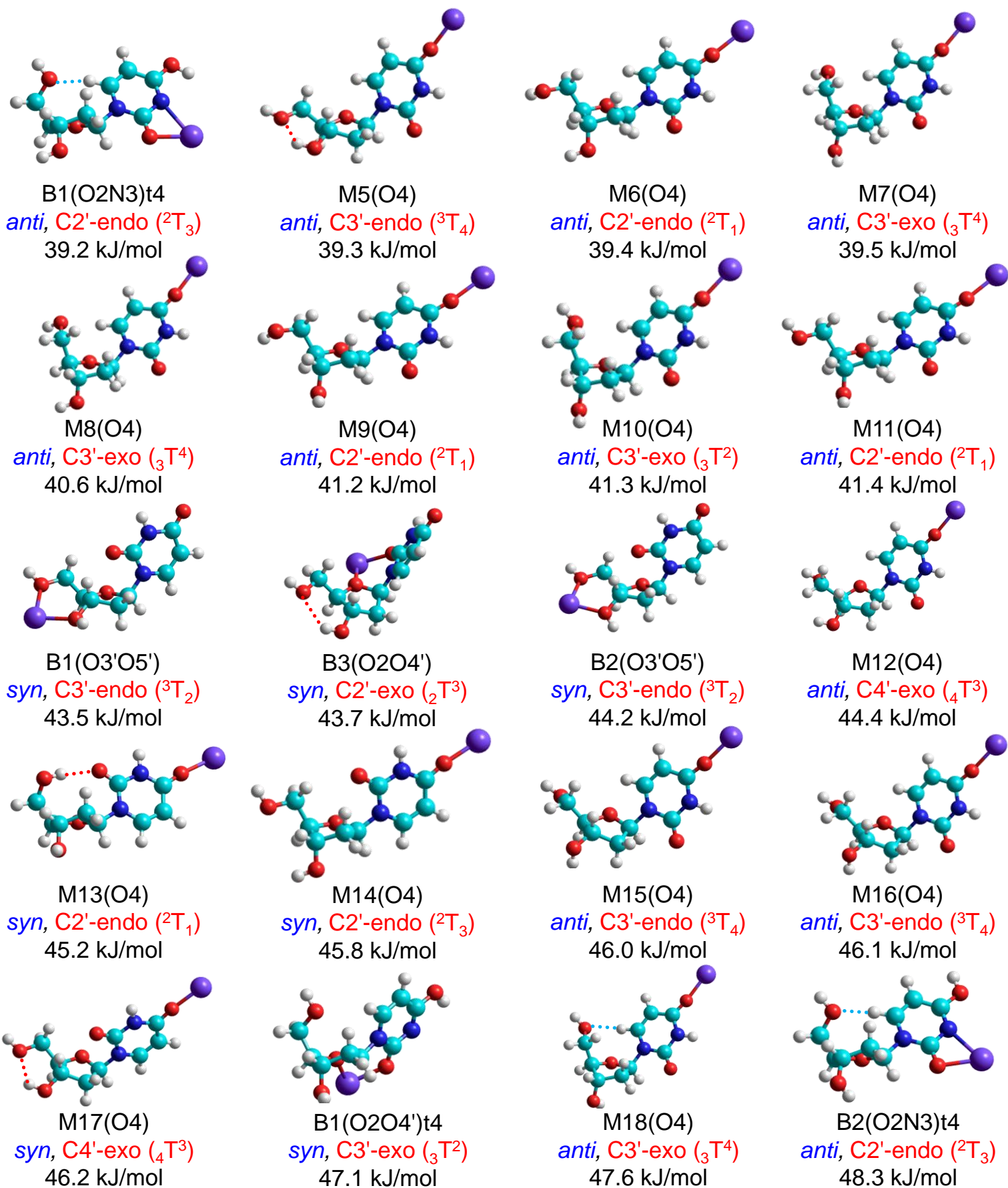


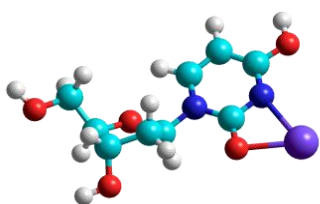
B14(O2N3)
syn, C2'-endo (2T_3)
66.1 kJ/mol

[Urd+Na]⁺

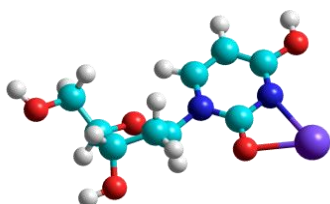




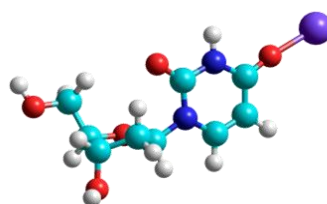




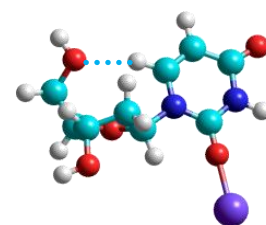
B3(O2N3)t4
anti, C2'-endo (2T_1)
48.5 kJ/mol



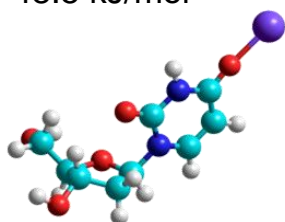
B4(O2N3)t4
anti, C2'-endo (2T_1)
48.6 kJ/mol



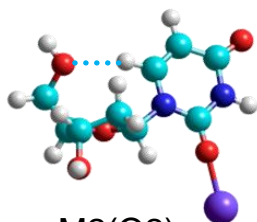
M19(O4)
syn, C2'-endo (2T_3)
48.9 kJ/mol



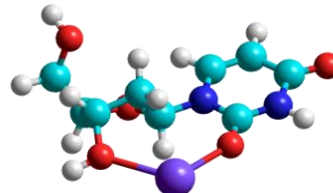
M1(O2)
anti, O4'-endo (0T_4)
49.0 kJ/mol



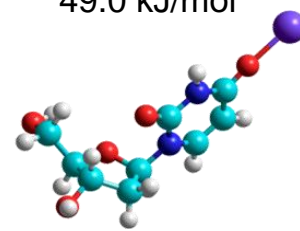
M20(O4)
syn, C4'-exo (${}_4T^3$)
49.5 kJ/mol



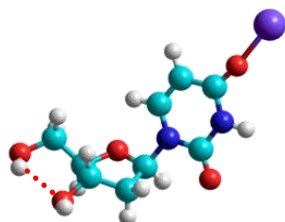
M2(O2)
anti, C2'-endo (2T_3)
49.7 kJ/mol



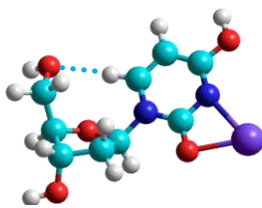
B1(O2O3')
anti, C2'-endo (2T_1)
49.9 kJ/mol



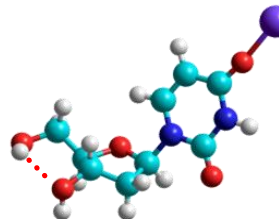
M21(O4)
syn, C4'-exo (${}_4T^3$)
50.3 kJ/mol



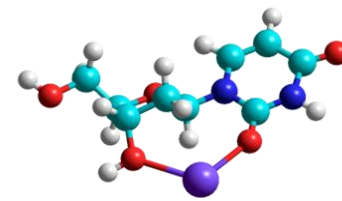
M22(O4)
anti, C3'-endo (3T_4)
51.1 kJ/mol



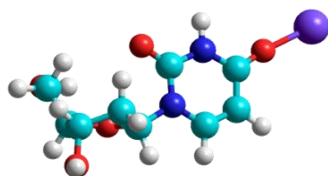
B5(O2N3)t4
anti, C3'-exo (${}_3T^4$)
51.4 kJ/mol



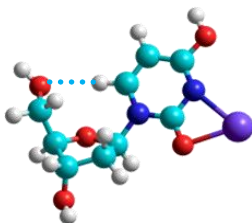
M23(O4)
anti, C3'-endo (3T_4)
51.8 kJ/mol



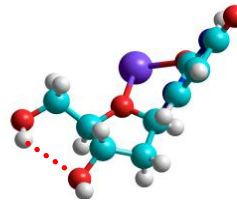
B2(O2O3')
anti, C1'-exo (${}_1T^2$)
51.9 kJ/mol



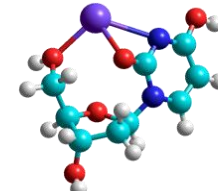
M24(O4)
syn, C2'-endo (2T_3)
52.3 kJ/mol



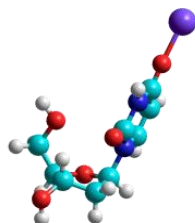
B6(O2N3)t4
anti, C3'-exo (${}_3T^2$)
52.7 kJ/mol



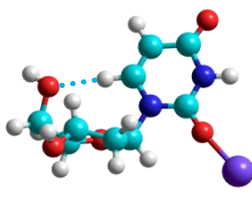
B4(O2O4')
syn, C3'-endo (3T_2)
55.6 kJ/mol



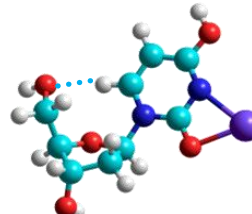
T1(O2N3O5')t4
syn, C3'-exo (${}_3T^2$)
55.8 kJ/mol



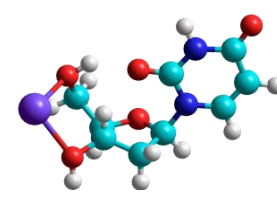
M25(O4)
anti, C2'-exo (${}_2T^1$)
56.0 kJ/mol



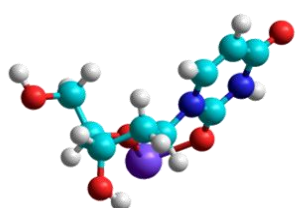
M3(O2)
anti, C3'-endo (3T_4)
56.6 kJ/mol



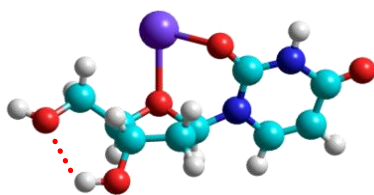
B7(O2N3)t4
anti, C3'-exo (${}_3T^4$)
57.6 kJ/mol



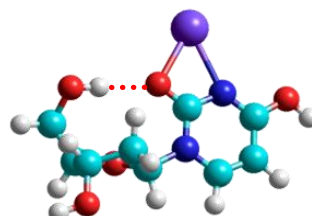
B2(O3'O5')
syn, C3'-endo (3T_2)
57.8 kJ/mol



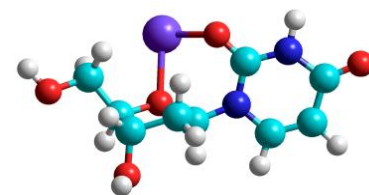
B5(O2O4')
syn, C3'-exo (${}_3T_2$)
58.4 kJ/mol



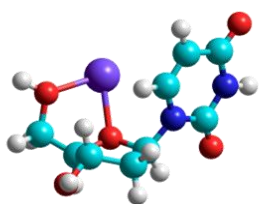
B6(O2O4')
syn, O4'-endo (0T_1)
58.9 kJ/mol



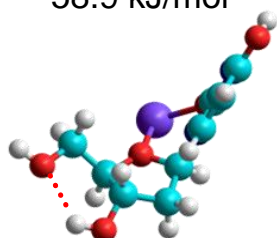
B8(O2N3)t4
syn, C2'-endo (2T_1)
59.6 kJ/mol



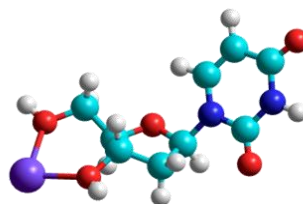
B7(O2O4')
syn, C1'-exo (${}_1T_2$)
59.7 kJ/mol



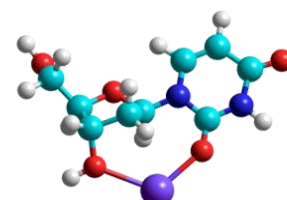
B1(O4'O5')
anti, C4'-exo (${}_4T_3$)
60.6 kJ/mol



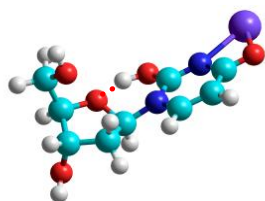
B2(O2O4')t4
syn, C2'-exo (${}_2T_3$)
60.9 kJ/mol



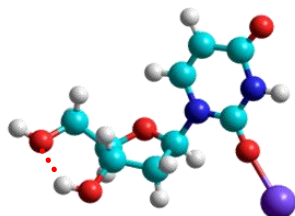
B3(O3'O5')
anti, C4'-exo (${}_4T_3$)
62.4 kJ/mol



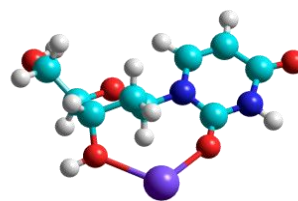
B3(O2O3')
anti, C1'-exo (${}_1T_2$)
63.0 kJ/mol



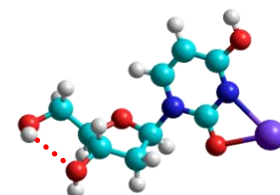
B1(N3O4)t2
syn, C2'-endo (2T_3)
63.8 kJ/mol



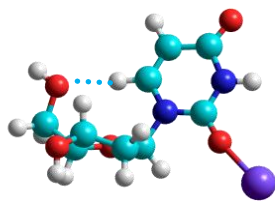
M4(O2)
anti, C3'-endo (3T_4)
63.9 kJ/mol



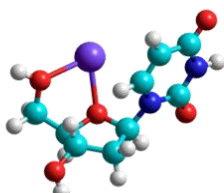
B4(O2O3')
anti, C1'-exo (${}_1T_2$)
64.2 kJ/mol



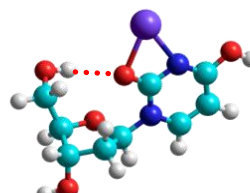
B9(O2N3)t4
anti, C3'-endo (3T_4)
65.0 kJ/mol



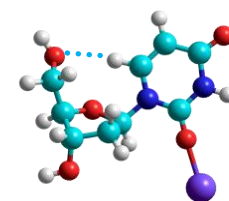
M5(O2)
anti, C3'-endo (3T_2)
65.3 kJ/mol



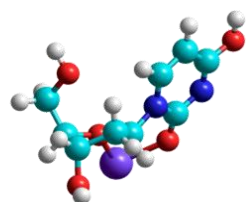
B2(O4'O5')
anti, C4'-exo (${}_4T_3$)
66.1 kJ/mol



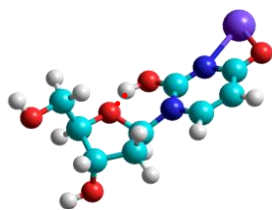
B10(O2N3)t4
syn, C2'-endo (2T_3)
66.6 kJ/mol



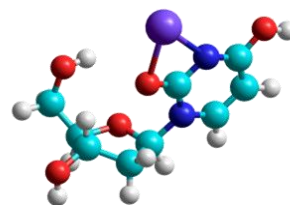
M6(O2)
anti, C3'-exo (3T_2)
67.6 kJ/mol



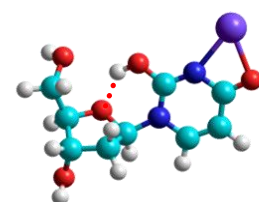
B3(O2O4')t4
syn, C3'-exo (${}_3T_2$)
68.8 kJ/mol



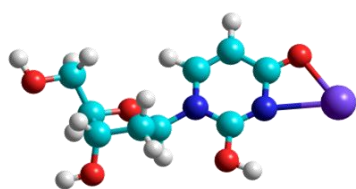
B2(N3O4)t2
syn, C2'-endo (2T_1)
69.9 kJ/mol



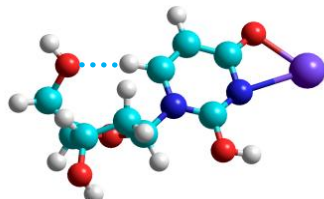
B11(O2N3)t4
syn, C4'-exo (${}_4T_3$)
70.1 kJ/mol



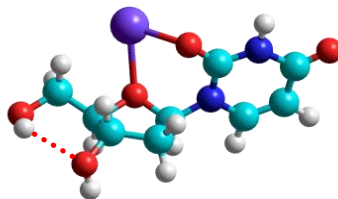
B3(N3O4)t2
syn, C2'-endo (2E)
70.4 kJ/mol



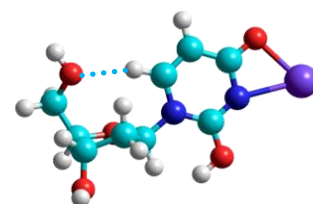
B4(N3O4)t2
anti, C2'-endo (2T_1)
72.3 kJ/mol



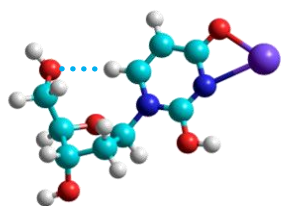
B5(N3O4)t2
anti, C2'-endo (2T_3)
72.5 kJ/mol



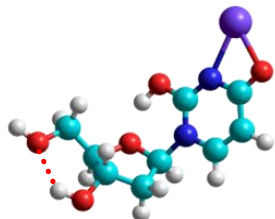
B8(O2O4)'
syn, O4'-endo (0T_4)
75.9 kJ/mol



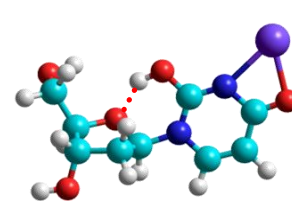
B6(N3O4)t2
anti, C2'-endo (2T_3)
76.6 kJ/mol



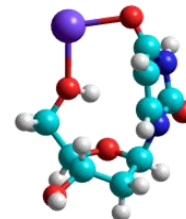
B7(N3O4)t2
anti, C3'-exo (${}_3T^2$)
77.5 kJ/mol



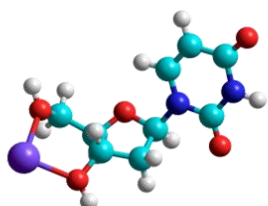
B8(N3O4)t2
syn, C3'-endo (3T_4)
77.6 kJ/mol



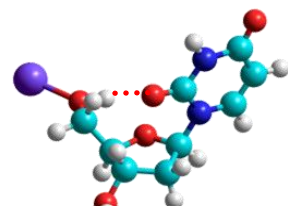
B9(N3O4)t2
syn, C1'-exo (${}_1T^2$)
78.9 kJ/mol



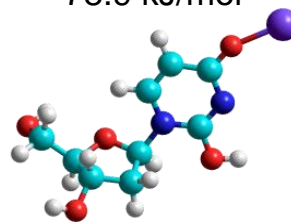
B1(O4O5)'
syn, C2'-exo (${}_2T^3$)
79.0 kJ/mol



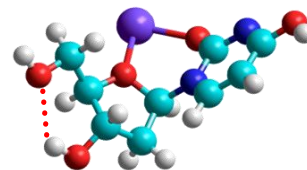
B4(O3'O5)'
anti, C4'-exo (${}_4T^3$)
80.9 kJ/mol



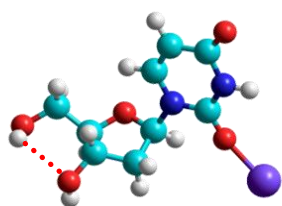
M1(O5)'
syn, C3'-endo (3T_2)
81.8 kJ/mol



M1(O4)t2
anti, C4'-exo (${}_4T^3$)
82.2 kJ/mol



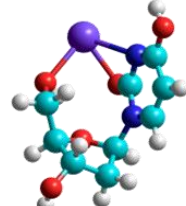
B4(O2O4)'t4
syn, C2'-exo (${}_2T^3$)
82.7 kJ/mol



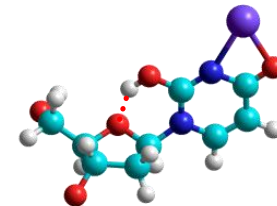
M7(O2)
anti, C3'-endo (3T_4)
83.7 kJ/mol



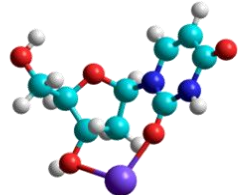
B1(O2O3)'t4
anti, C1'-exo (${}_1T^2$)
85.1 kJ/mol



T2(O2N3O5)'t4
syn, C1'-endo (1T_2)
86.0 kJ/mol



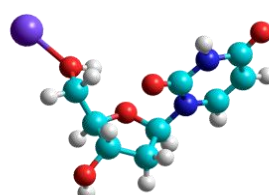
B10(N3O4)t2
syn, O4'-endo (0T_4)
86.4 kJ/mol



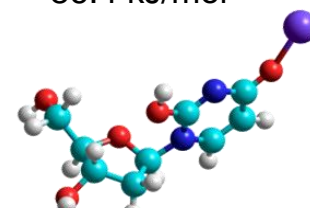
B5(O2O3)'
anti, C3'-endo (3T_2)
88.1 kJ/mol



M1(O3)'
syn, C4'-exo (${}_4T^3$)
88.3 kJ/mol

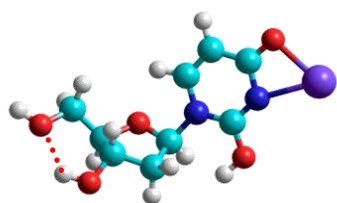


M2(O5)'
syn, C3'-endo (3T_2)
89.7 kJ/mol

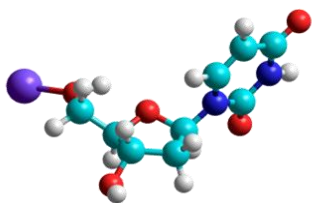


M2(O4)t2
syn, C4'-exo (${}_4T^0$)
91.3 kJ/mol

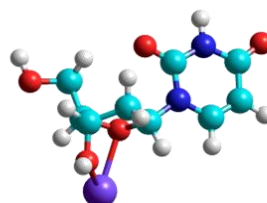
[dUrd+Na]⁺



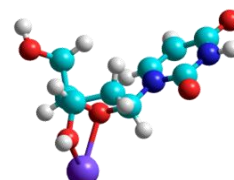
B11(N3O4)t2
anti, C3'-endo (3T_4)
92.3 kJ/mol



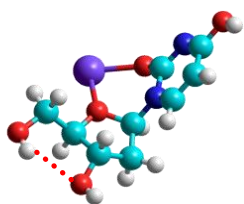
M3(O5')
anti, C4'-exo (${}^4T^0$)
93.0 kJ/mol



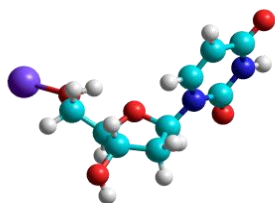
B1(O3'O4')
syn, C3'-exo (${}^3T^4$)
93.5 kJ/mol



B2(O3'O4')
anti, C4'-endo (4T_3)
94.1 kJ/mol



B5(O2O4')t4
syn, C3'-endo (3T_2)
95.2 kJ/mol



M4(O5')
anti, C4'-exo (${}^4T^0$)
99.8 kJ/mol

Figure S5.

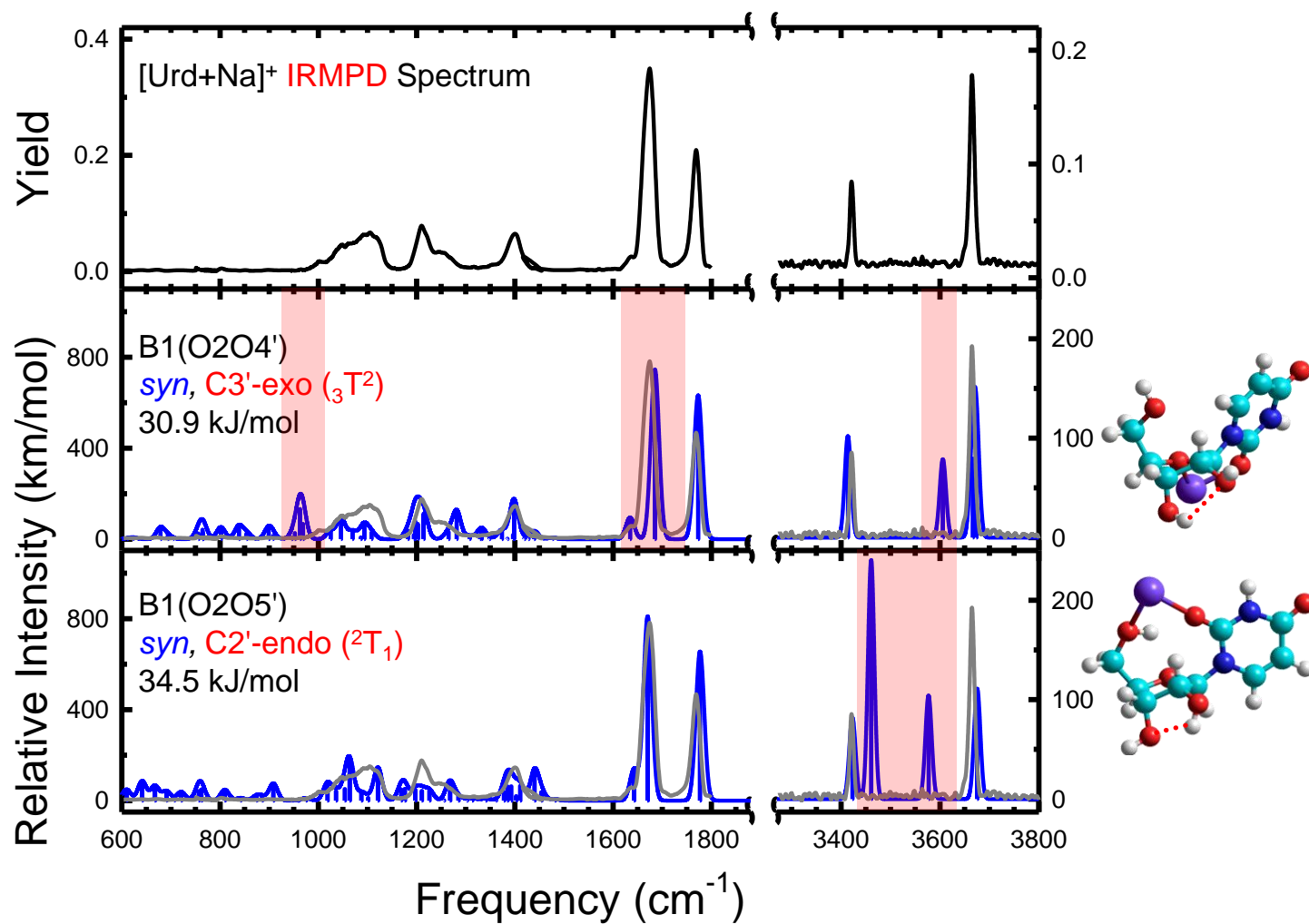


Figure S6.

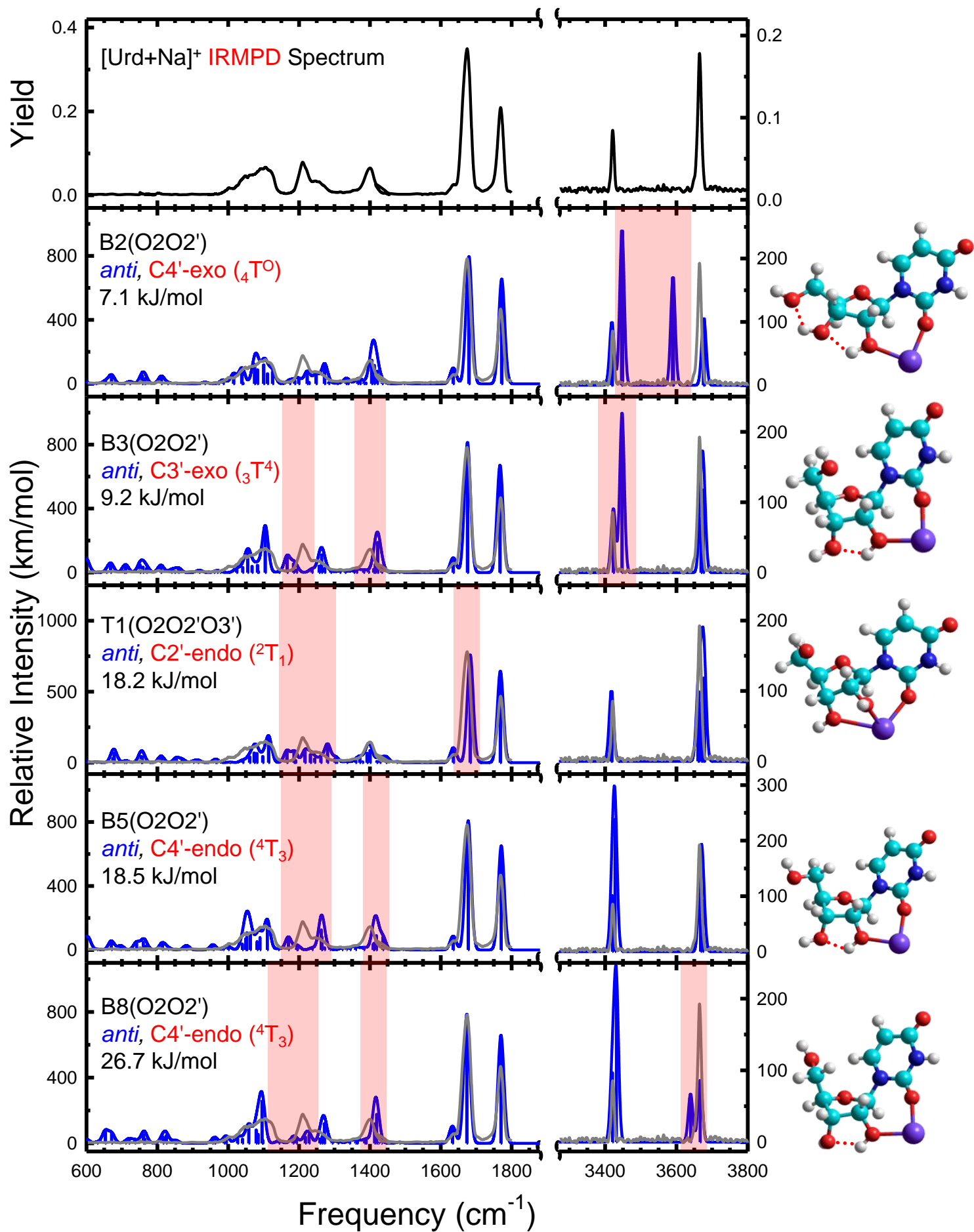


Figure S7.

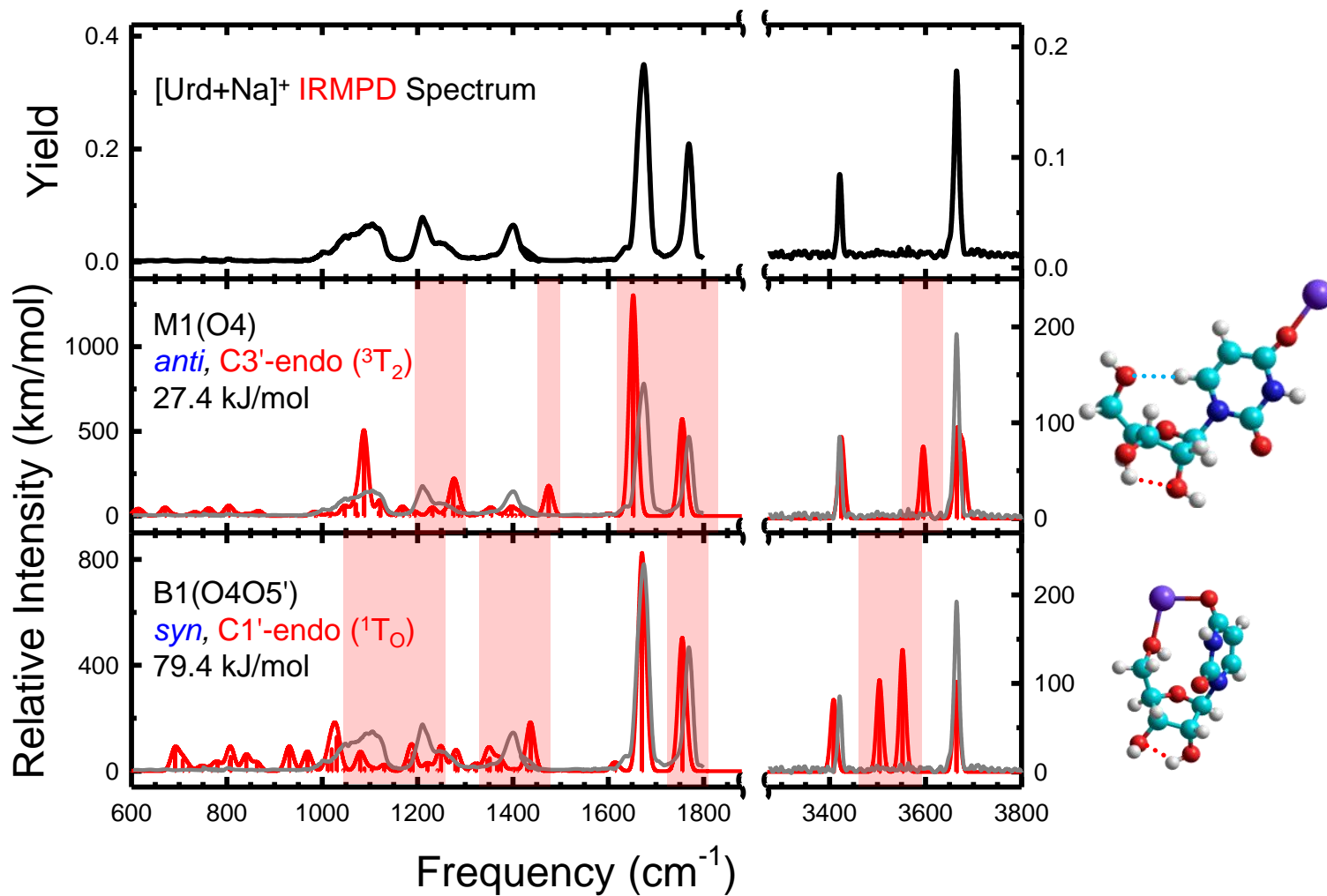


Figure S8.

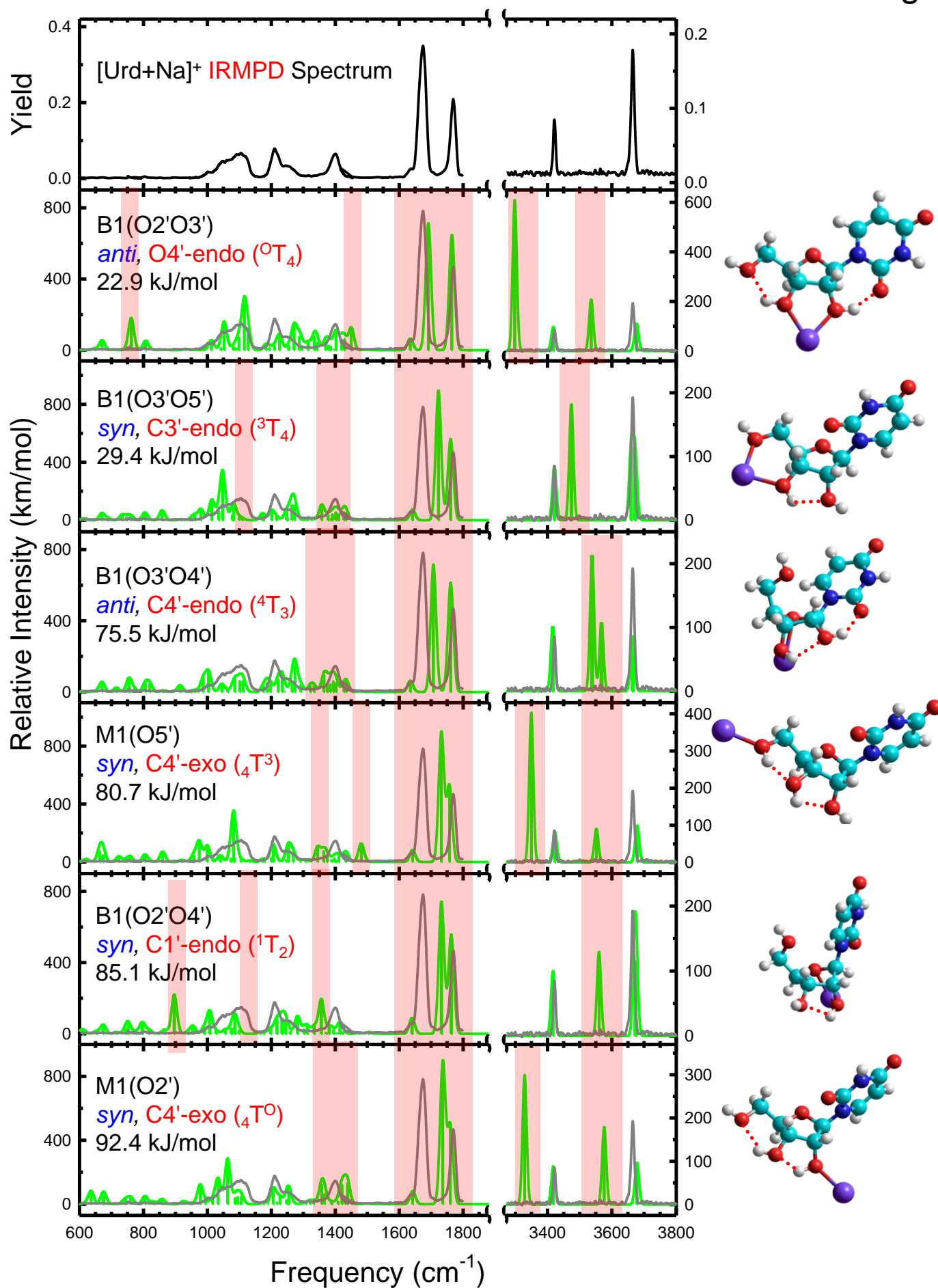
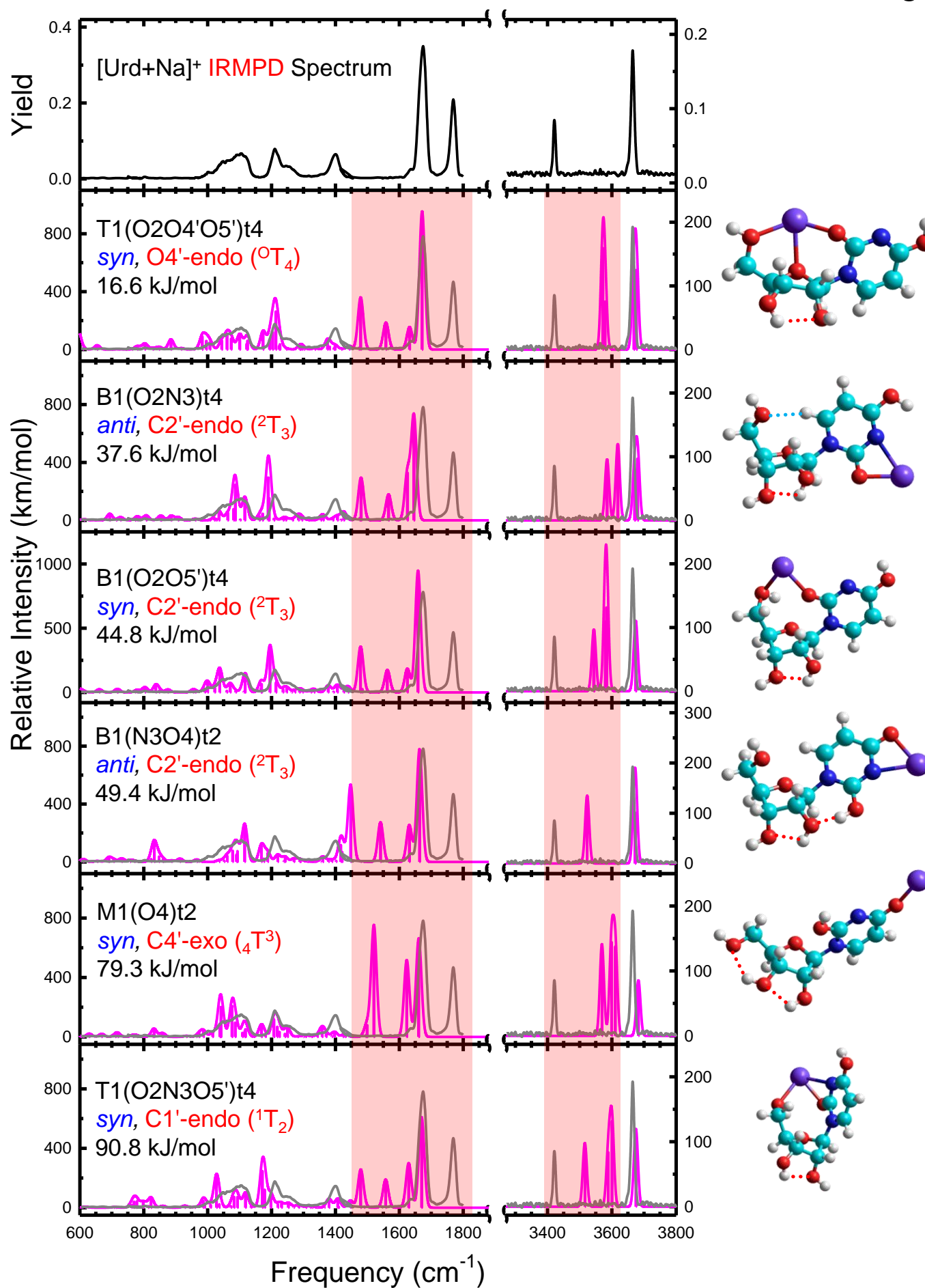
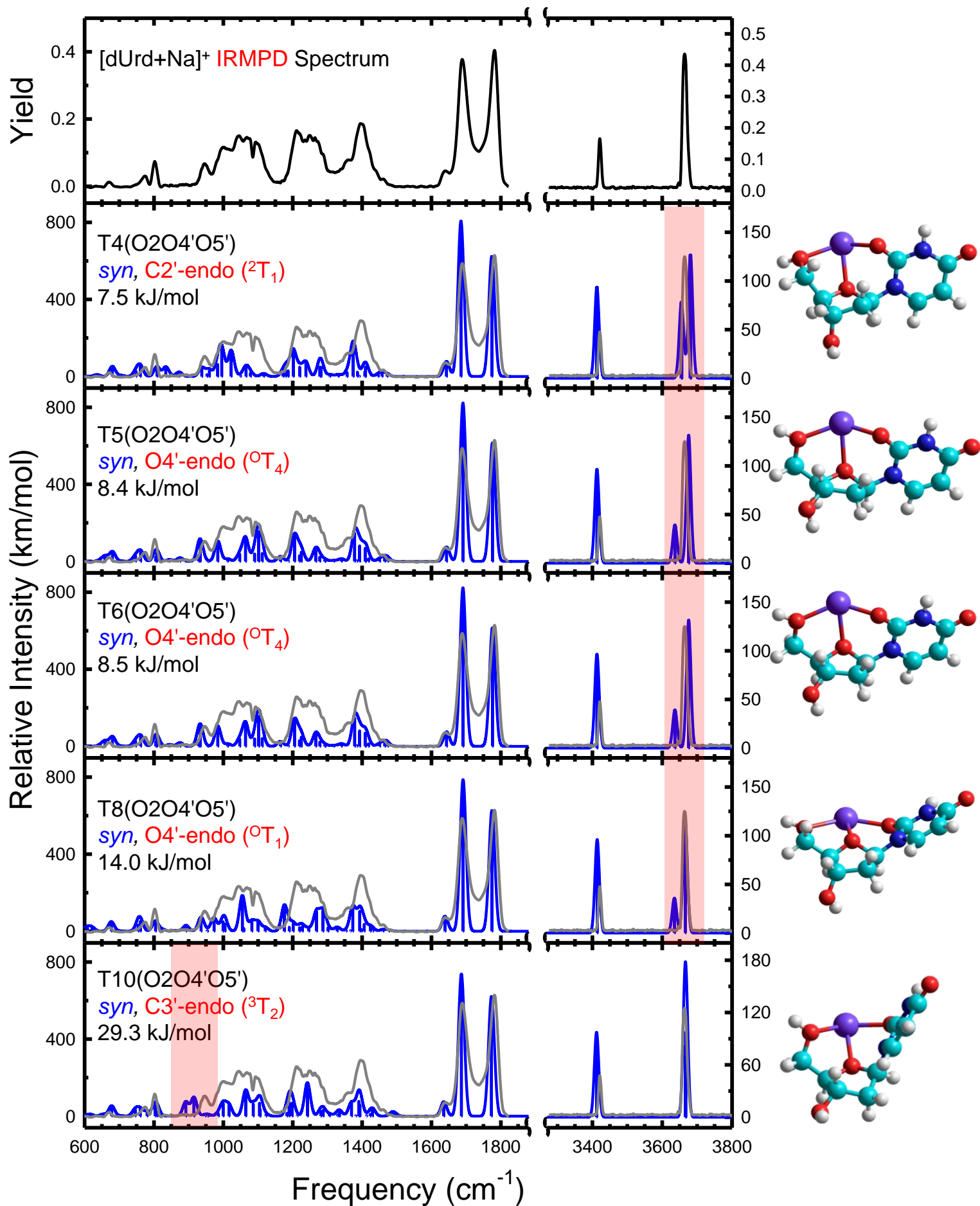
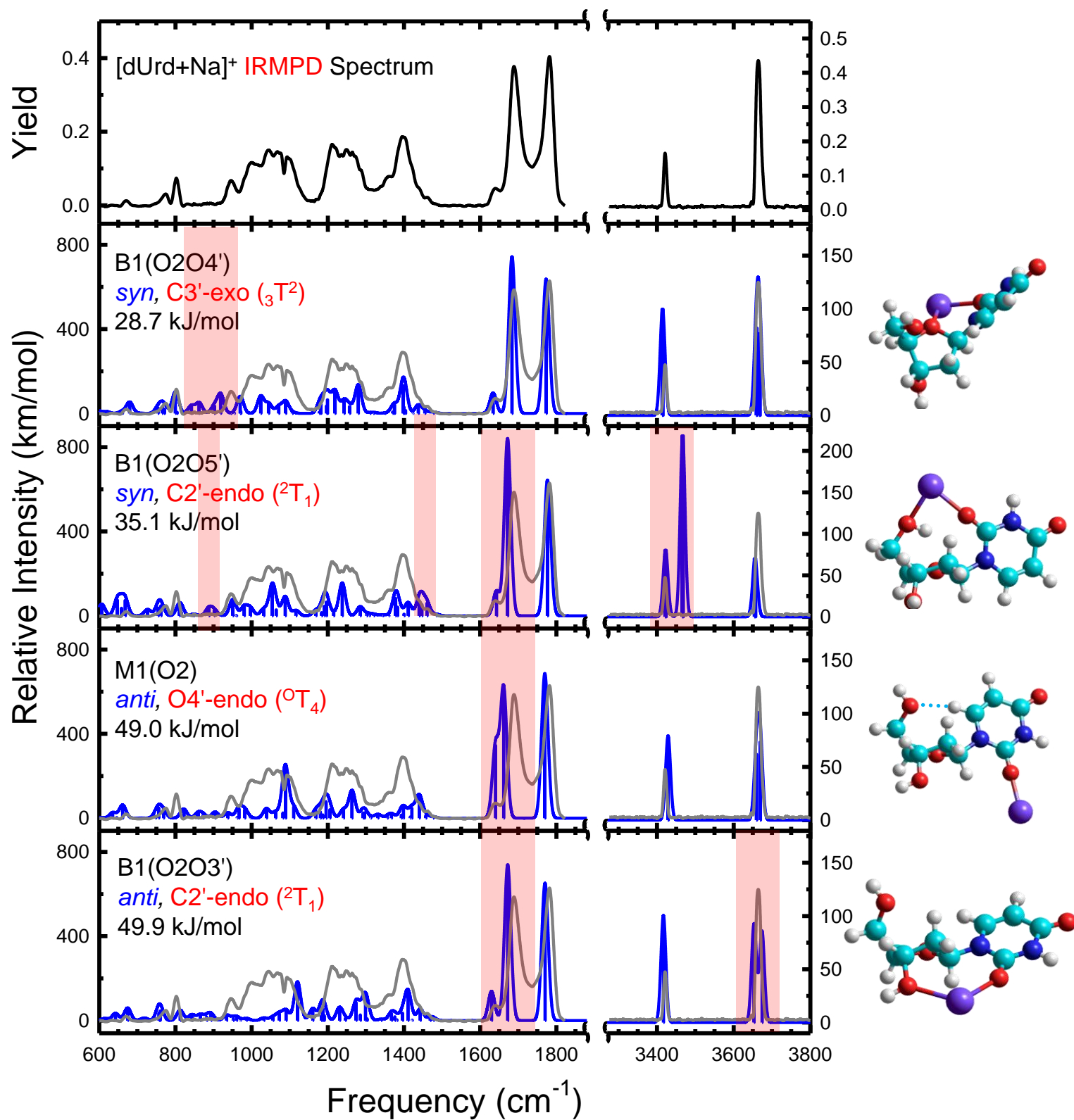
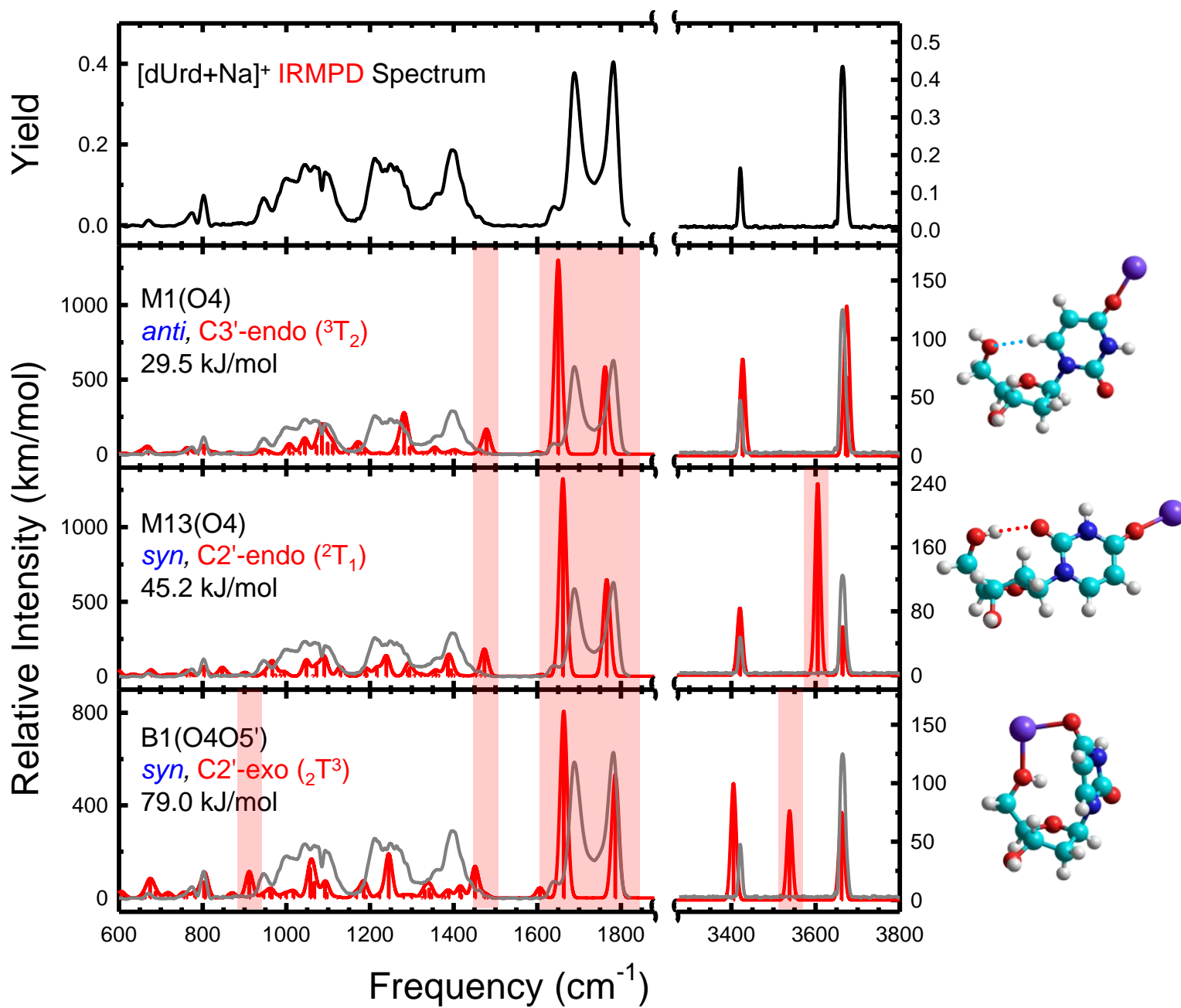


Figure S9.









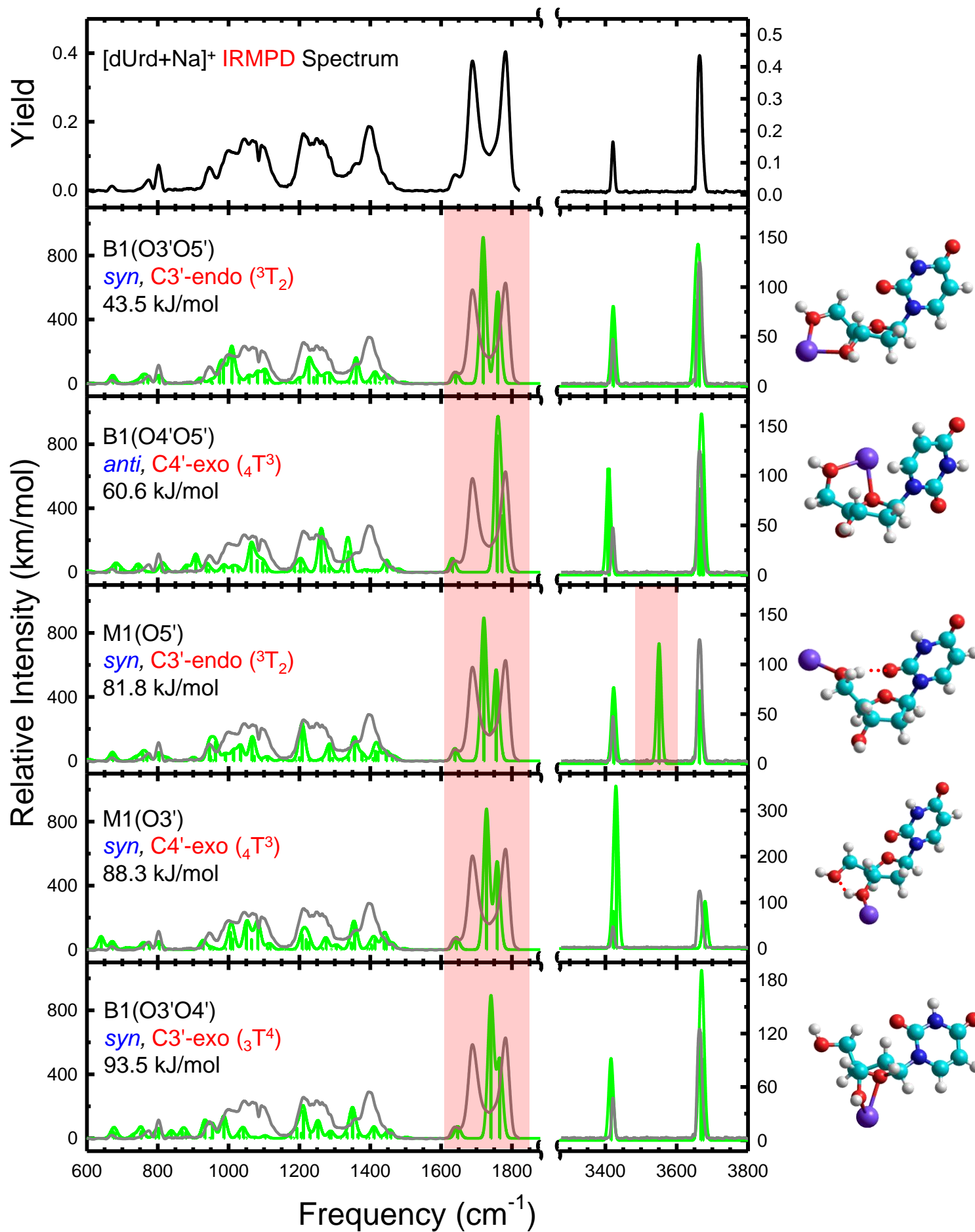


Figure S14.

