Electronic Supplementary Information for...

## The Intrinsic Basicity of the Phosphate Backbone Exceeds that of Uracil and Thymine Residues: Protonation of the Phosphate Moiety is Preferred over the Nucleobase for pdThd and pUrd

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## Discussion

**Comparison of Experimental IRMPD and Theoretical IR Spectra of [pdThd+H]<sup>+</sup>.** The measured IRMPD and calculated IR spectra of the **PD**, **PE**, **TB**, **PF**, **TC**, and **O4B** of [pdThd+H]<sup>+</sup> are compared in Figure S3 over the IR fingerprint and hydrogen-stretching regions. Mismatches between the measured and calculated spectra are highlighted. The highly anharmonic IRMPD features measured below ~1650 cm<sup>-1</sup> in the fingerprint region are extensively broadened and thus not diagnostic. The calculated band at 1800 cm<sup>-1</sup> of the **O4B** conformer is shifted to a higher frequency relative to the measured band at 1768 cm<sup>-1</sup>. The calculated bands at 3583, 3595, 3581 and 3615 cm<sup>-1</sup> of **PD**, **TB**, **TC** and **O4B**, respectively, are shifted to higher frequencies relative to the measured band at 3652 cm<sup>-1</sup>. The bands predicted at 3636 cm<sup>-1</sup> for **PE** and **PF** are lower in frequency than the measured band at 3652 cm<sup>-1</sup>. These obvious discrepancies observed between the measured and calculated bands indicate the absence of the **TB**, **PE**, **PF**, and **O4B** conformers in the experiments as well.

Comparison of Experimental IRMPD and Theoretical IR Spectra of  $[pUrd+H]^+$ . The measured IRMPD and calculated IR spectra of the Piii, Piv, PD, PE, TB, Ti and TC conformers of  $[pUrd+H]^+$  are compared in Figure S4 over the IR fingerprint and hydrogen-stretching regions. Mismatches between the measured and calculated spectra are highlighted. Again, the IRMPD features measured below ~1650 cm<sup>-1</sup> in the fingerprint region for all these seven conformers and the hydrogen-bond acceptor hydrogen-stretches predicted at 3685 cm<sup>-1</sup> for Piii, Piv, PD, PE and TC are not diagnostic due to their anharmonicities. The bands predicted at 3636 cm<sup>-1</sup> for Piii,

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**Piv** and **PE** are lower in frequency than the measured band at 3652 cm<sup>-1</sup>. The small IRMPD features predicted at 3615 and 3619 cm<sup>-1</sup> for **Piv** and **Ti** respectively are higher in frequency than the weak band observed at 3588 cm<sup>-1</sup>. The broad band predicted at 3666 cm<sup>-1</sup> for **TB** is higher in frequency than the measured band at 3652 cm<sup>-1</sup>. The calculated bands at 3579 cm<sup>-1</sup> of **PD**, **Ti** and **TC** are lower in frequency than the measured band at 3588 cm<sup>-1</sup>. The modest and broad band predicted at 3588 for TB matches the band position of the weak band observed at 3588 cm<sup>-1</sup>. However, it is too intense relative to that of the measured band. Therefore, the discrepancies between the measured and calculated spectra particularly in the hydrogen-stretching region suggest that the **Piii**, **Piv**, **PD**, **PE**, **TB**, **Ti** and **TC** conformers are not populated in the experiments.

The measured IRMPD and calculated IR spectra of the Pv, PF, Tii, O4B, O4i, Tiii and **O2i** conformers of [pUrd+H]<sup>+</sup> are compared in Figure S5 over the IR fingerprint and hydrogenstretching regions. Mismatches between the measured and calculated spectra are highlighted. The IRMPD features measured below  $\sim 1650 \text{ cm}^{-1}$  in the fingerprint region for all seven conformers and the hydrogen-bond acceptor hydrogen-stretches predicted at 3685 cm<sup>-1</sup> for Pv and PF, and at 3678 cm<sup>-1</sup> for Tiii and O2i are again not diagnostic. The strong IR feature predicted at 1822 cm<sup>-1</sup> for **PF** is not observed in the measured spectrum. The calculated bands at 1814 and 1822 cm<sup>-1</sup> of O4B and O4i respectively are higher in frequency than the measured band at 1797 cm<sup>-1</sup>. The calculated bands at 3632 cm<sup>-1</sup> of **Pv** and **PF** are lower in frequency than the band observed at 3652 cm<sup>-1</sup>. The calculated IR bands in the hydrogen-stretching regions of the Tii, O4B, O4i, Tiii and O2i conformers exhibit very distinct differences as compared to the measured IRMPD bands. In particular, the bands predicted at 3670 cm<sup>-1</sup> for Tii, O4B and O4i are higher in frequency than the measured band at 3652 cm<sup>-1</sup>. The IR features predicted at 3627 cm<sup>-1</sup> for Tii, O4B and O4i are not observed in the measured spectrum. The band predicted at 3691 cm<sup>-1</sup> for **Tii** is too intense relative to the small measured band at that vibrational frequency. The calculated bands at 3519 and 3504 cm<sup>-1</sup> of Tiii and O2i are not observed in the measured spectrum. Therefore, the discrepancies between the measured and calculated spectra particularly in the hydrogen-stretching region suggest that the Pv, PF, Tii, O4B, O4i, Tiii and O2i conformers are not populated in the experiments.

Species	Conformer		MP2(full)			B3LYP	
_		$\Delta H_0$	$\Delta H_{298}$	$\Delta G_{298}$	$\Delta H_0$	$\Delta H_{298}$	$\Delta G_{298}$
[pdThd+H] <sup>+</sup>	$\mathbf{PA}^{b}$	0.0	0.0	0.2	0.0	0.0	0.8
	$\mathbf{PB}^{b}$	2.1	2.5	0.0	1.5	1.9	0.0
	PC	15.1	14.4	17.1	4.5	3.8	7.1
	ТА	28.5	28.5	28.1	24.2	24.1	24.3
	$\mathbf{O2A}^b$	16.0	11.6	24.8	5.9	1.5	15.3
	O4A	25.2	24.0	26.7	14.6	13.4	16.7
$\left[\text{pdUrd+H}\right]^+$	$\mathbf{PA}^{b}$	0.0	0.3	2.1	0.3	0.8	10.4
	$\mathbf{PB}^{b}$	1.1	0.0	4.0	0.9	0.0	11.8
	PC	8.0	12.2	0.0	0.0	4.4	0.0
	ТА	25.1	26.8	23.5	21.4	23.3	27.9
	O4A	21.1	20.2	22.4	10.2	9.5	19.4

**Table S1.** Relative Enthalpies and Free Energies at 0 and 298 K in kJ/mol of Select Low-Energy Conformers of  $[pdThd+H]^+$  and  $[pUrd+H]^+$  in a Polarizable Continuum<sup>*a*</sup>

<sup>*a*</sup>Single point energy calculations using the B3LYP/6-311+G(d,p) optimized structures are performed at the MP2(full)/6-311+G(2d,2p) and B3LYP/6-311+G(2d,2p) levels of theory with the IEFPCM polarizable continuum model and include ZPE and thermal corrections. The relative stabilities of the ground conformers are indicated in boldface. <sup>*b*</sup>Low-energy conformers that are populated in the gas phase and contribute to the measured IRMPD action spectra.

## **Figure Captions**

**Figure S1.** B3LYP/6-311+G(d,p) low-energy conformers of  $[pdThd+H]^+$  and their relative Gibbs free energies at 298 K calculated at the MP2(full)/6-311+G(2d,2p) (in red) and B3LYP/6-311+G(2d,2p) (in black) levels of theory. The site of protonation/tautomeric form, nucleobase orientation, and sugar puckering are also indicated for each conformer.

**Figure S2.** B3LYP/6-311+G(d,p) low-energy conformers of  $[pUrd+H]^+$  and their relative Gibbs free energies at 298 K calculated at the MP2(full)/6-311+G(2d,2p) (in red) and B3LYP/6-311+G(2d,2p) (in black) levels of theory. The site of protonation/tautomeric form, nucleobase orientation, and sugar puckering are also indicated for each conformer.

**Figure S3.** Comparison of the measured IRMPD action spectrum of  $[pdThd+H]^+$  with the theoretical linear IR spectra of the **PD**, **PE**, **TB**, **PF**, **TC**, and **O4B** conformers of  $[pdThd+H]^+$  and the corresponding B3LYP/6-311+G(d,p) optimized structures. Also shown are the MP2(full)/6-311+G(2d,2p) (in red) and B3LYP/6-311+G(2d,2p) (in black) relative Gibbs free energies at 298 K. The site of protonation/tautomeric form, nucleobase orientation, and sugar puckering are also indicated for each conformer. To facilitate comparison of the measured and computed spectra, the IRMPD spectrum is overlaid (in grey) with each computed spectrum and scaled to match the intensity of the most intense feature in each region.

**Figure S4.** Comparison of the measured IRMPD action spectrum of  $[pUrd+H]^+$  with the theoretical linear IR spectra of the **Piii**, **Piv**, **PD**, **PE**, **TB**, **Ti** and **TC** conformers of  $[pUrd+H]^+$  and the corresponding B3LYP/6-311+G(d,p) optimized structures. Also shown are the MP2(full)/6-311+G(2d,2p) (in red) and B3LYP/6-311+G(2d,2p) (in black) relative Gibbs free energies at 298 K. The site of protonation/tautomeric form, nucleobase orientation, and sugar puckering are also indicated for each conformer. To facilitate comparison of the measured and computed spectra, the IRMPD spectrum is overlaid (in grey) with each computed spectrum and scaled to match the intensity of the most intense feature in each region.

**Figure S5.** Comparison of the measured IRMPD action spectrum of  $[pUrd+H]^+$  with the theoretical linear IR spectra of the **Pv**, **PF**, **Tii**, **O4B**, **O4i**, **Tiii** and **O2i** conformers of  $[pUrd+H]^+$  and the corresponding B3LYP/6-311+G(d,p) optimized structures. Also shown are the MP2(full)/6-311+G(2d,2p) (in red) and B3LYP/6-311+G(2d,2p) (in black) relative Gibbs free energies at 298 K. The site of protonation/tautomeric form, nucleobase orientation, and sugar puckering are also indicated for each conformer. To facilitate comparison of the measured and computed spectra, the IRMPD spectrum is overlaid (in grey) with each computed spectrum and scaled to match the intensity of the most intense feature in each region.

Figure S1.

S5



**PA**, syn, C2'-endo 0.0, 0.7 kJ/mol



PD, syn, C2'-endo 11.1, 15.2 kJ/mol



**TB**, anti, C2'-endo 11.8, 7.9 kJ/mol



TC, syn, C2'-endo 14.4, 9.0 kJ/mol



**PB**, syn, C2'-endo 0.2, 0.3 kJ/mol



**TA**, syn, C3'-endo 11.7, 4.3 kJ/mol



**PF**, syn, C2'-endo 12.4, 2.3 kJ/mol



**O4A**, anti, C2'-endo 18.7, 6.4 kJ/mol

[pdThd+H]<sup>+</sup>



PC, syn, C3'-endo 10.6, 0.0 kJ/mol



PE, syn, C2'-endo 11.7, 5.2 kJ/mol



**O2A**, syn, C2'-endo 13.0, 2.9 kJ/mol



**O4B**, anti, C3'-endo 24.8, 9.8 kJ/mol

**Pi**, syn, C2'-endo 0.0, 3.9 kJ/mol



**PB**, syn, C2'-endo 5.0, 7.3 kJ/mol



Piii, syn, C3'-endo 9.7, 2.2 kJ/mol



PE, syn, C2'-endo 13.0, 8.1 kJ/mol



Pii, syn, C2'-endo 0.2, 3.5 kJ/mol



PC, syn, C3'-endo 6.0, 0.0 kJ/mol



Piv, syn, C2'-endo 10.8, 8.1 kJ/mol



**TB**, anti, C2'-endo 13.1, 8.8 kJ/mol

[pUrd+H]<sup>+</sup>



**PA**, syn, C2'-endo 4.2, 7.0 kJ/mol



**TA**, syn, C3'-endo 7.3, 2.1 kJ/mol



PD, syn, C2'-endo 11.9, 16.5 kJ/mol



Ti, syn, C3'-endo 13.4, 8.4 kJ/mol

Figure S2.

Figure S2.

**S**7



TC, syn, C2'-endo 13.7, 6.9 kJ/mol



**Tii**, anti, C2'-endo 15.9, 14.0 kJ/mol



**O4i**, anti, C2'-endo 23.5, 14.2 kJ/mol



Pv, syn, C2'-endo 14.6, 9.2 kJ/mol



**O4A**, anti, C2'-endo 18.1, 6.5 kJ/mol



**Tiii**, anti, C2'-endo 27.5, 20.8 kJ/mol

[pUrd+H]<sup>+</sup>



**PF**, syn, C2'-endo 15.8, 8.2 kJ/mol



**O4B**, anti, C3'-endo 23.3, 11.6 kJ/mol



**O2i**, anti, C2'-endo **39.1**, 32.2 kJ/mol





**S**8

Figure S4.





S10

