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Unconventional hydrogen bonding to organic ions in the gas phase:
Stepwise association of hydrogen cyanide with the pyridine and pyrimidine radical cations and protonated pyridine

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Equilibrium thermochemical measurements using the ion mobility drift cell technique have been utilized to investigate the binding energies and entropy changes for the stepwise association of HCN molecules with the pyridine and pyrimidine radical cations forming the C5H5N+(HCN)n and C4H4N2+(HCN)n clusters, respectively, with n = 1–4. For comparison, the binding of 1–4 HCN molecules to the protonated pyridine C5H4NH+(HCN)n has also been investigated. The binding energies of HCN to the pyridine and pyrimidine radical cations are nearly equal (11.4 and 12.0 kcal/mol, respectively) but weaker than the HCN binding to the protonated pyridine (14.0 kcal/mol). The pyridine and pyrimidine radical cations form unconventional carbon-based ionic hydrogen bonds with HCN (CHδ+···NCH). Protonated pyridine forms a stronger ionic hydrogen bond with HCN (NHδ+···NCH) which can be extended to a linear chain with the clustering of additional HCN molecules (NHδ+···NCH···NCH···NCH) leading to a rapid decrease in the bond strength as the length of the chain increases. The lowest energy structures of the pyridine and pyrimidine radical cation clusters containing 3–4 HCN molecules show a strong tendency for the internal solvation of the radical cation by the HCN molecules where bifurcated structures involving multiple hydrogen bonding sites with the ring hydrogen atoms are formed. The unconventional H-bonds (CHδ+···NCH) formed between the pyridine or the pyrimidine radical cations and HCN molecules (11–12 kcal/mol) are stronger than the similar (CHδ+···NCH) bonds formed between the benzene radical cation and HCN molecules (9 kcal/mol) indicating that the CHδ+ centers in the pyridine and pyrimidine radical cations have more effective charges than in the benzene radical cation. © 2014 AIP Publishing LLC.

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I. INTRODUCTION

Hydrogen bonding is one of the most important intermolecular interactions in chemistry and biology both in gas phase and condensed phase systems.1–3 A special class of this interaction, usually referred to as ionic hydrogen bonds (IHBs), involves hydrogen bonding between radical ions or protonated molecules and neutral molecules.4 IHBs have bond strengths higher than the typical conventional hydrogen bond in neutral systems and they could reach up to 35 kcal/mol, nearly a third of the strength of covalent bonds. These strong interactions are critical in many fields such as ion induced nucleation, ionic clusters, ion solvation, radiation chemistry, electrochemistry, acid-base chemistry, and self-assembly in supramolecular chemistry.1–6 IHBs are also important in biological systems including protein folding, proton transport, membranes, enzyme active centers, and molecular recognition.1–6

Unconventional carbon-based IHBs are formed when the hydrogen donors are ionized hydrocarbons containing CH groups and the hydrogen acceptors are electron lone pairs on hetero atoms such as O or N, olefin double bonds, or aromatic π systems.4 For example, carbon-based CHδ+···O IHBs appear in the hydration of ionized aromatics such as benzene (C6H6δ+), cyclic C3H3δ+ and phenyl acetylene (C8H6δ+).7–10 In addition to water, other polar molecules containing lone pair of electrons such as hydrogen cyanide can participate in hydrogen bonding interactions with the ring hydrogen atoms (CHδ+) of ionized aromatics. Hydrogen cyanide is a useful probe of non-covalent interactions because it is a highly polar molecule (μ = 2.98 D), and it can serve both as a hydrogen donor and as a lone-pair hydrogen acceptor in hydrogen bonds. Furthermore, HCN is an important atmospheric compound known to be produced by biomass burning, and it can be produced in interstellar/nebula environments by the reactions of ammonia and methane.11 In fact, HCN polymers have been shown to exist in meteorites, comets, planets, moons, and in circumstellar envelopes.12–14 Ion-molecule interactions involving HCN, particularly those that lead to the formation of larger species either through chemical addition or association reactions, are of particular interest for the formation of complex organics, clustering, and polymerization in astrochemical environments.13,16

We recently studied the stepwise association of HCN with benzene, substituted benzene, and phenylacetylene radical cations.17–19 In benzene δ+(HCN)n clusters the lig-
II. EXPERIMENTAL SECTION

The gas phase ion association experiments were performed using the Virginia Commonwealth University (VCU) mass-selected ion mobility spectrometer. The details of the instrument can be found in several publications and only a brief description of the experimental procedure is given here.\textsuperscript{8,17,21} In the experiments, the molecular ions of pyridine (C\textsubscript{5}H\textsubscript{4}N\textsuperscript{+}), pyrimidine (C\textsubscript{4}H\textsubscript{4}N\textsubscript{2}\textsuperscript{+}), and C\textsubscript{5}H\textsubscript{5}N\textsuperscript{+} (HCN)\textsubscript{n} clusters with n = 1–4, and investigate the role of hydrogen bonding interactions –CH\textsuperscript{+}···NCH and –NH\textsuperscript{+}···NCH as compared to the ion-dipole N\textsuperscript{+}···NCH interaction in the formation mechanism of these nitrogen-containing cluster ions. The results provide new insights into the factors that determine the structures and energetics of the clusters of polar molecules with N-containing heterocyclic organic ions.

The equilibrium reactions (taken pyridine radical cation C\textsubscript{5}H\textsubscript{5}N\textsuperscript{+} as an example) are represented by Eq. (1),

\[ [C_{5}H_{5}N^{+}(HCN)_{n-1}] + HCN \rightleftharpoons [C_{5}H_{5}N^{+}(HCN)_{n}] \]  

The establishment of equilibrium is verified when: (1) a constant ratio of the integrated intensity of the product to the reactant ions is maintained over the residence time of the ions at constant pressure and temperature, and (2) the ATDs of the reactant and product ions are identical indicating equal residence times. When the equilibrium conditions are well-established, the equilibrium constant, \( K_{eq} \), can be measured using Eq. (2),

\[ K_{eq} = \frac{[C_{5}H_{5}N^{+} \cdot (HCN)_{n}]}{[C_{5}H_{5}N^{+} \cdot (HCN)_{n-1}] \cdot [HCN]} \]

where \([C_{5}H_{5}N^{+} \cdot (HCN)_{n-1}]\) and \([C_{5}H_{5}N^{+} \cdot (HCN)_{n}]\) are the integrated intensities of the ATDs of the reactant and product cluster ions, respectively of reaction (1) and \( P_{HCN} \) is the pressure of HCN (in atmosphere) inside the drift cell. The equilibrium constant, \( K_{eq} \), is measured at different temperatures and from a van’t Hoff plot, \( \Delta H^{o} \) and \( \Delta S^{o} \) values are obtained from the slope and intercept, respectively. The measured values are duplicated at least three times and the average values are reported in Table I the corresponding uncertainties.
TABLE I. Measured thermochemistry (−ΔH°n-1,n and −ΔS°n-1,n) of the formation of C5H5N⁺·(HCN)n and C5H5NH⁺·(HCN)n clusters; with n = 1-4 and the corresponding calculated binding energies for the two lowest energy structures within 2 kcal/mol.

<table>
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<th>Pyridine⁺⁺·(HCN)n</th>
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<tr>
<td>n</td>
<td>−ΔH°a</td>
<td>−ΔS°b</td>
<td>ΔE (M06-2X)c</td>
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<td>21.8</td>
<td>11.9–11.4</td>
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<tr>
<td>n</td>
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<tr>
<td>4</td>
<td>6.1</td>
<td>14.5</td>
<td>8.4–7.0</td>
</tr>
</tbody>
</table>

aΔH°n-1,n units are kcal/mol.

bΔS°n-1,n units are cal/mol K. Error estimate for experimental values from standard deviations of van’t Hoff and from usual uncertainties in clustering equilibrium temperature studies: ΔH° ±1 kcal/mol, ΔS° ±2 cal/mol K.

cBinding energy calculated at the M06-2X/6-311++G(d,p) level with ZPE and BSSE corrections included.

III. THEORETICAL SECTION

DFT calculations of the lowest energy structures of C5H5N⁺·(HCN)n, C4H4N2⁺·(HCN)n, and C5H5NH⁺·(HCN)n clusters with n = 1–4 were carried out at the M06-2X/6-311++G(d,p) level using the Gaussian 09 suite of programs.22 Frequency calculations have been performed for all the optimized geometries at the same level of theory to obtain the zero point vibrational energy (ZPVE) and to verify the absence of any imaginary frequencies. The calculated binding energies (with respect to C5H5N⁺·(HCN)n+1 + HCN) were corrected for basis set superposition error (BSSE) using the scheme of Boys and Bernardi as described in the Gaussian program.22

IV. RESULTS AND DISCUSSION

A. Association of HCN with the pyridine radical cation and protonated pyridine

Figure 1(a) displays the mass spectra obtained following the injection of the mass-selected pyridine ion into the drift cell containing 1.0 Torr He at 298 K. It is clear that no dissociation products are observed consistent with the low injection energy used (13.8 eV, lab). In the presence of 0.25 Torr HCN vapor in the drift cell at 283 K, the first two association products C5H5N⁺·(HCN)n with n = 1 and 2 are observed as shown in Fig.1(a). As the temperature decreases the ion intensity of C5H5N⁺·(HCN)n significantly decreases and eventually disappears as the equilibrium shifts to higher C5H5N⁺·(HCN)n clusters. At 194 K, the cluster population is dominated by the C5H5N⁺·(HCN)n ions with n = 3 and 4. Similar results are obtained following the injection of the protonated pyridine into the drift cell containing HCN as shown in Fig. 1(b).

![Mass spectra](image_url)

**FIG. 1.** (a) Mass spectra resulting from the injection of the mass-selected pyridine radical cation (Py⁺⁺) into helium gas or HCN vapor at different temperatures using 13.8 eV injection energy (laboratory frame) and 4 V/cm applied field. (b) Mass spectra resulting from the injection of protonated pyridine (H⁺Py) into helium gas or HCN vapor at different temperatures using 14.2 eV injection energy (laboratory frame) and 4 V/cm applied field. The peaks labeled (a) m/z 165 and (b) m/z 192 are due to the association of HCN with a trace impurity (m/z 138) present in the drift cell.
A good test of equilibrium comes from the identical ATDs of the ions coupled by equilibrium. If the \( C_5H_5N^+ \cdot (HCN)_{n-1} \) and \( C_5H_5N^+ \cdot (HCN)_{n} \) ions are in equilibrium, their ATDs must be identical. This is evident from the ATDs shown in Figure S1 of the supplementary material for the \( C_5H_5N^+ \cdot (HCN)_{n} \) ions. The equilibrium constants for the stepwise association of HCN with \( C_5H_5N^+ \cdot (HCN)_{n} \) and \( C_5H_5NH^+ \cdot (HCN)_{n} \) measured at different temperatures yield the van’t Hoff plots shown in Figures 2(a) and 2(b), respectively. The resulting \( \Delta H^o \) and \( \Delta S^o \) values for the formation of the \( C_5H_5N^+ \cdot (HCN)_{n} \) and \( C_5H_5NH^+ \cdot (HCN)_{n} \) clusters are listed in Table I.

The sequential binding energies of HCN to the pyridine radical cation or to the protonated pyridine, shown in Table I, follow the trend of decreasing \( \Delta H^o_{n-1,n} \) with increasing \( n \) which can be expected for association reactions dominated by ion-dipole interactions. Furthermore, the sequential entropy loss values are consistent with ion-dipole interactions where internal rotation and low frequency vibrations are retained. The thermochemical data also show that HCN binds more strongly to the protonated pyridine \( (-\Delta H^o = 14.0 \text{ kcal/mol}) \) than to the pyridine radical cation \( (-\Delta H^o = 11.4 \text{ kcal/mol}) \). However, the difference in binding between the radical cation and the protonated pyridine decreases as the number of HCN molecules increases which is expected since the charge-dipole interaction between protonated pyridine and the associated HCN molecule decreases with increasing the cluster size. Thus the binding energies for the \( n = 4 \) molecule in both the \( C_5H_5N^+ \cdot (HCN)_{n} \) and \( C_5H_5NH^+ \cdot (HCN)_{n} \) clusters are essentially similar (6.4 and 6.1 kcal/mol, respectively).

The DFT lowest energy structures of the \( C_5H_5N^+ \cdot (HCN)_{n} \) and \( C_5H_5NH^+ \cdot (HCN)_{n} \) clusters with \( n = 1–4 \) calculated at the M06-2X/6-311++G(d,p) are shown in Figures 3 and 4, respectively, and the binding energies calculated for the two lowest structures (within 2 kcal/mol) are listed in Table I. The calculated binding energies agree reasonably with the measured \( (-\Delta H^o_{n-1,n}) \) although in most cases, the binding energies of the second lowest energy structures appear to be in better agreement with the experimental values than the lowest energy structures.

Two isomers (1-a and 1-b) with binding energies of 11.9 and 11.4 kcal/mol, respectively, are predicted for the \( C_5H_5N^+ \cdot (HCN)_{n} \) cluster as shown in Fig. 3. These binding energies are in excellent agreement with the experimental value of 11.4 kcal/mol. Both structures show H-bonding between the C–H⋯ of the pyridine cation and the N atom of HCN. However isomer (a) has a bifurcated structure with the N-atom of HCN forming two long H-bonds (2.56 Å and 2.34 Å) with the ortho- and meta-CH groups of the pyridine cation. This structure is similar to the lowest energy isomer of the benzene\(^+ \cdot (HCN)\) cluster.\(^{17} \) The second lowest energy isomer (b) has a 2.07 Å hydrogen bond between the N-atom of HCN and the ortho-aromatic hydrogen atom. The third most stable isomer (1-c) has the HCN molecule attached to the para-aromatic hydrogen atom via 2.2 Å hydrogen bond. This isomer is 2.7 kcal/mol higher in energy than isomer 1-a, and therefore, it gives a binding energy significantly lower (9.3 kcal/mol) than the experimental value of 11.4 kcal/mol. The three lowest energy structures of the \( C_5H_5N^+ \cdot (HCN)_{n} \) cluster confirm that the interaction of HCN with the pyridine cation is dominated by hydrogen bonding between the N of HCN and the ring H atom. In spite of the acidic character of the H atom of HCN, no hydrogen bonding is observed to the N atom of the pyridine cation which carries a net positive charge as shown by the charge density distribution (Figure S2 of the supplementary material).\(^{23} \)

The lowest energy structures of \( C_5H_5N^+ \cdot (HCN)_{n} \) clusters with \( n = 2–4 \) show a tendency for the internal solvation of the pyridine cation by the HCN molecules where bifurcated structures involving multiple hydrogen bonding sites with the ring hydrogen atoms are formed such as shown in structures (2-a), (3-a), (3-b), (4-a), and (4-b). Structures (2-b) and (3-c)

\[ \text{FIG. 2. van’t Hoff plots of the temperature dependence of the equilibrium constants for the stepwise association of HCN with (a) the pyridine radical cation yielding } \frac{C_5H_5N^+ \cdot (HCN)}{n=1-4} \text{, and (b) protonated pyridine yielding } \frac{C_5H_5NH^+ \cdot (HCN)}{n=1-4}. \]
show the formation of hydrogen bonding chains involving two or three HCN molecules, respectively, attached to the pyridine cation through bifurcated structures. However, no chain structures involving four HCN molecules have been found in the \( \text{C}_5\text{H}_5\text{N}^+\cdot(\text{HCN})_4 \) clusters indicating that HCN interactions with the pyridine ring are more favorable than the interactions within the HCN chains. It is interesting to note that the chain structures (2-b, \( \Delta E = 9.4 \text{ kcal/mol} \)) and (3-c, \( \Delta E = 8.2 \text{ kcal/mol} \)) show good agreement with the measured binding energies (8.8 and 7.8 kcal/mol, respectively) unlike the solvated pyridine structures (2-a, \( \Delta E = 10.9 \text{ kcal/mol} \)) and (3-a, \( \Delta E = 10.6 \text{ kcal/mol} \)) which tend to overestimate the binding energies. On the other hand, for the \( \text{C}_5\text{H}_5\text{N}^+\cdot(\text{HCN})_4 \) cluster the solvated pyridine structures (4-b, \( \Delta E = 6.8 \text{ kcal/mol} \)) and (4-c, \( \Delta E = 6.1 \text{ kcal/mol} \)) are in good agreement with the measured binding energy of 6.4 kcal/mol.

The lowest energy structures of the HCN molecules associated with the protonated pyridine in the \( \text{C}_5\text{H}_5\text{NH}^+\cdot(\text{HCN})_n \) clusters are dominated by conventional IHBs between the \( \text{NH}^+ \) group of the protonated pyridine and the nitrogen atom of HCN as shown in all structures displayed in Fig. 4 except structures (1-b) and (1-c). These high energy structures exhibit significantly lower binding energies (1-b and 1-c, \( \Delta E = 9.2 \text{ and 8.1 kcal/mol} \)) as compared to the experimental value of 14.0 kcal/mol for the \( \text{C}_5\text{H}_5\text{NH}^+\cdot(\text{HCN}) \) cluster. Interestingly, HCN extended chain structures are more frequent among the \( \text{C}_5\text{H}_5\text{NH}^+\cdot(\text{HCN})_n \) clusters as shown by the low energy structures (2-b), (3-c), and (4-c).
B. Association of HCN with the pyrimidine radical cation

Figure 5 displays the mass spectra obtained following the injection of the mass-selected pyrimidine ion into the drift cell containing either pure He or HCN vapor at different temperatures. The first association product C4H4N2··(HCN) is observed with 0.2 Torr HCN at 274 K and as the temperature of the drift cell decreases higher association products C4H4N2··(HCN)n with n up to 4 are observed as shown in Fig. 5. The ATDs of the C4H4N2··(HCN)n-1 and C4H4N2··(HCN)n ions indicate the establishment of equilibrium as shown in Figure S3 of the supplementary material. The equilibrium constants for the stepwise association of HCN with C4H4N2·· measured at different temperatures yield the van’t Hoff plots shown in Figure 6, and the resulting ΔH° and ΔS° values for the formation of the C4H4N2··(HCN)n clusters and calculated binding energies at the M06-2X/6-311++G(d,p) are listed in Table II.

The binding energy of HCN to the pyrimidine radical cation (12 kcal/mol) is slightly higher than the corresponding value for the pyridine cation (11.4 kcal/mol) and lower than that of the protonated pyridine (14.0 kcal/mol). This indicates that HCN binds to the pyrimidine cation via a CHδ+···NCH unconventional hydrogen bond, which is relatively weaker than the IHB NHδ+···NCH formed with the protonated pyridine. This also indicates that the pyrimidine cation in our experiments has the conventional radical cation structure (C4H4N2+) and not the distonic structure (C4H3NNH+) which would bind stronger to HCN.
The DFT lowest energy structures of the C₄H₄N₂⁺·(HCN)ₙ clusters with n = 1–4 calculated at the M06-2X/6-311++G(d,p) level are shown in Figure 7. The lowest energy structure of the C₄H₄N₂⁺·(HCN) complex has a T-shaped ion-dipole structure with the N atom of HCN pointing toward the center of the pyrimidine cation ring. The second and third lowest energy structures represent unconventional CH···NCH hydrogen bonds similar to the structures found for the C₃H₃N⁺·(HCN) complex shown in Fig. 3. The strongest H donor site in the pyrimidine cation is the one located between the two nitrogen atoms of the ring and this site appears to form most of the non-bifurcated H bonds with the nitrogen atom of the HCN molecule. The calculated binding energies of C₄H₄N₂⁺·(HCN)ₙ clusters agree reasonably with the measured (−ΔH°ᵣ₋ᵣ₋) except for the C₄H₄N₂⁺·(HCN)₃ cluster where the calculated binding energy (10.7 kcal/mol) is significantly higher than the measured −ΔH°ᵣ₋ᵣ₋ value of 8.6 kcal/mol. This could indicate that the structure of the C₄H₄N₂⁺·(HCN)₃ cluster formed under our experimental conditions may be different from the calculated lowest energy structure which has the bifurcated hydrogen bonding structures of one HCN molecule to two CH⁺ groups of the ring and also two T-shaped ion-dipole structures involving two HCN molecules. This complicated structure may be kinetically unfavorable since the collisions of HCN molecules with the lowest energy structure of the C₄H₄N₂⁺·(HCN)₂ cluster (structure (2-a), Fig. 7) to form structure (3-b) do not require further rearrangements of the HCN molecules as in the case of structure (3-a).

Similar to the trend observed in the structures of the pyridine⁺·(HCN)ₙ clusters, the formation of the solvated cation structures are also observed in the pyrimidine⁺·(HCN)ₙ clusters as shown in structures (4-a),(4-b), and (4-c) displayed in Fig. 7 for the n = 4 cluster.

C. Comparison with the MP2/6-311++G(d,p) level calculations

In order to compare the results of the DFT/M06-2X/6-311++G(d,p) calculations with a different method, we carried out calculations on the C₄H₄N⁺·(HCN), C₄H₄NH⁺(HCN), and C₄H₄N₂⁺·(HCN) complexes at the MP2/6-311++G(d,p) level. The calculated lowest energy structures, their relative energies, and binding energies of the complexes are shown in Figure 8. The three lowest energy structures of the pyridine⁺· and protonated pyridine complexes with HCN are similar at both the DFT/M06-2X/6-311++G(d,p) and MP2/M06-2X/6-311++G(d,p) levels as shown by comparing the results shown in Figures 3, 4, and 8. The calculated binding energies using the two methods are also similar. Interestingly, the overestimated binding energy of the pyridineH⁺(HCN) complex (16 kcal/mol with the
FIG. 7. Structures of the pyrimidine$^+\cdot$(HCN)$_4$ clusters calculated by DFT at the M06-2x/6-311++G (d,p) level. Relative energies (RE) are the total electronic energies with respect to the lowest energy isomer (0.0). Binding energies (ΔE) are corrected for zero-point energies (ZPE) and basis-set super position errors (BSSE). Energies are in kcal/mol. Distances are in Angstrom.

The only major difference between the DFT and MP2 results is the order of the lowest energy structures of the pyrimidine$^+\cdot$(HCN) complex as shown from the comparison between the results shown in Figures 7 and 8. While DFT predicts a T-shaped ion-dipole structure for the C$_4$H$_8$N$_2$+\cdot(HCN) complex (structure (1-a), Figure 7), the MP2 method shows the unconventional hydrogen bonding structure (CH$^+\cdot\cdot\cdot$NCH) to be the lowest energy isomer (structure 1-a-pyrimidine$^+\cdot$(HCN), Figure 8). With the DFT method, the hydrogen bonding structure was the second lowest energy structure (structure (1-b)) with only 0.4 kcal/mol lower binding energy than the T-shaped structure (structure (1-a)) as shown in Figure 7. However, with the MP2 method the T-shaped structure has substantially lower binding energy (8.3 kcal/mol, structure 1-c-pyrimidine$^+\cdot$(HCN)) as compared to the hydrogen bonding structure (12.2 kcal/mol, structure 1-a) as shown in Figure 8 for the pyrimidine$^+\cdot$(HCN) complex. This indicates that the hydrogen bonding structure ((1-b) in Fig. 7 and 1-a, third row in Fig. 8) is most likely the lowest energy structure for the pyrimidine$^+\cdot$(HCN) complex. This could also
suggest that the ion-dipole interaction is either underestimated or overestimated with the MP2 and DFT methods, respectively.

D. Comparison of HCN interactions with the benzene, pyridine, pyrimidine radical cations, and protonated pyridine

Table III shows a comparison of the sequential binding energies of 1-4 HCN molecules to the benzene, pyridine and pyrimidine radical cations, and the protonated pyridine. Figure 9 compares the trends in binding energies for the four ions. The binding of HCN molecules to the benzene cation is mostly due to unconventional CH$_2^+\cdots$NCH hydrogen bonds directly connected to the CH$_2^+$ sites of the benzene cation and also hydrogen bonding chains (HCN$\cdots$HCN) among the HCN molecules. The small difference in the bond strength of the two types of interactions results in relatively small changes of ($-\Delta H^\circ_{n-1,n}$) for $n = 1-4$ as shown in Fig. 9. HCN binds more strongly to the pyridine and pyrimidine radical cations due to the presence of stronger ion-dipole interactions in addition to the CH$_2^+$—NCH hydrogen bonding interactions. The strongest binding is observed between the protonated pyridine and HCN as a result of IHB that forms between the NH$_2^+$ group of the protonated pyridine and the N atom of HCN. A significant drop in the binding energy (31%) is observed upon the addition of the second HCN molecule to the protonated pyridine in contrast to the smaller changes observed upon the addition of the second HCN molecule to the pyridine or pyrimidine radical cations (23%, and 15%, respectively) as shown in Fig. 9. Despite the strong bonding of HCN to protonated pyridine, the interaction decreases sharply by further addition of HCN molecules and terminal binding for the fourth HCN molecule is only 6.1 kcal/mol. In fact, in all of the studied HCN clusters around ionized or protonated aromatics,\textsuperscript{17-19} the binding energies converge with the addition of 4-5 HCN molecules to the enthalpy of vaporization of HCN liquid ($\Delta H^\circ_{vap}$), which is 6.0 kcal/mol at 298 K.\textsuperscript{24}

As indicated earlier, HCN is a useful probe of non-covalent interactions because it can serve both as a hydrogen donor and as a lone-pair hydrogen acceptor in...
different types of hydrogen bonds. Therefore, it is instructive to compare the structures of the HCN complexes of ionized heterocyclic aromatics such as pyridine and pyrimidine radical cations with the corresponding neutral complexes. In the neutral systems, e.g., pyridine···HCN, the existence of the aromatic conjugated system and the localized lone pair of electrons in pyridine provides the possibility of competitive interactions for various types of hydrogen bonds. For example, hydrogen bonding of the hydrogen atom in HCN to the \( \pi \)-system of pyridine prefers the T-shaped geometry while the interaction with the nitrogen lone pair of electrons leads to planar structures since the lone pair is located in the plane of the aromatic ring. However, in ionized aromatics only the –CH\(^+\)···NCH type (or the –NH\(^+\)···NCH in protonated pyridine) of hydrogen bonds is observed since the T-shaped structures ((4-c), Fig. 3 and (1-a), (3-d), Fig. 7) do not involve H-bonding as they represent ion-dipole interaction between the positively charged ring and the negative end of the dipole moment of HCN. It should be noted that the T-shaped hydrogen bonding of HCN to the \( \pi \)-system of the neutral aromatic becomes repulsive in ionized aromatics because of the positive charge on the ring and therefore, the HCN molecule must rotate 180° to adopt a new T-shaped structure where the nitrogen atom of HCN is now facing the positively charged ring as in structures (1-a) and (3-d) in Fig. 7.

### E. Applications in astrochemistry

Hydrogen cyanide is a significant component of interstellar clouds and solar nebulae.\(^{11-14}\) Aromatic molecules such as benzene, pyridine, pyrimidine, and polycyclic aromatic hydrocarbons (PAHs) could produce stable molecular ions under conditions of ionizing radiation. In these low-temperature environments, HCN can condense on the ions forming organic-doped ice grains. The protonation of the aromatic molecules and PAHs could result in significant binding of the polar space molecules such as HCN to the even-electron ions. The present systems model these processes, in particular, because the astrochemical condensation also involves stepwise addition of gas phase molecules.

### V. SUMMARY AND CONCLUSIONS

Equilibrium thermochemical measurements using the ion mobility drift cell technique have been utilized to investigate the binding energies and entropy changes for the stepwise association of HCN molecules with the pyridine and pyrimidine radical cations forming the \( \text{C}_5\text{H}_5\text{N}^+\cdot\cdot\cdot\text{(HCN)}_n \) and \( \text{C}_5\text{H}_4\text{N}_2^+\cdot\cdot\cdot\text{(HCN)}_n \) clusters, respectively, with \( n = 1-4 \). For comparison, the binding of 1-4 HCN molecules to the protonated pyridine \( \text{C}_5\text{H}_4\text{N}^+\cdot\cdot\cdot\text{(HCN)}_n \) has also been investigated. The binding energies of HCN to the pyridine and pyrimidine radical cations are nearly equal (11.4 and 12.0 kcal/mol, respectively) but weaker than the HCN binding to the protonated pyridine (14.0 kcal/mol). The pyridine and pyrimidine radical cations form unconventional carbon-based ionic hydrogen bonds with HCN (\( \text{CH}^+\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\cdot\·

![Fig. 9. Binding energy versus number of HCN molecules for the stepwise association with the benzene, pyridine and pyrimidine radical cations, and protonated pyridine.](image-url)
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23 See supplementary material at http://dx.doi.org/10.1063/1.4890372 for the arrival time distributions (ATDs) for the reactant and product ions to verify the attainment of equilibrium in the pyridine+·(HCN)n (Figure S1) and pyrimidine+·(HCN)n clusters (Figure S3), and calculated atomic charges and geometries of pyridine and pyrimidine radical cations, and protonated pyridine (Figure S2).