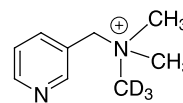


$1\text{--}2 \times 10^{-6}$  torr. Dimethyl ether was added to nominally  $1\text{--}2 \times 10^{-5}$  torr. Helium buffer gas was added to increase the pressure to one millitorr. Solids were introduced to the chamber via a heated direct insertion probe. Trimethyloxonium and trimethyloxonium- $d_6$ , generated from dimethyl ether and dimethyl ether- $d_6$  ions, respectively, was isolated and permitted to react with the neutral analyte for ca. 100 ms. The  $(M+\text{CH}_3)^+$  and  $(M+\text{CD}_3)^+$  product ions were isolated using the apex isolation method<sup>4</sup> and subjected to resonance collisionally activated dissociation (CAD), typically at a  $q_z$  of 0.3.

When **2** is methylated with trimethyloxonium, the only fragment observed from  $(M+\text{CH}_3)^+$  is  $m/z$  108, due to loss of 43 u. The loss of 43 u is best rationalized as the loss of  $\text{CH}_3\text{--N=CH}_2$ , which can readily occur from the dimethylamino portion of the molecule (see Scheme 2). The  $(M+\text{CD}_3)^+$  ion also dissociated exclusively by loss of 43 u to form  $m/z$  111, which means the same loss of  $\text{CH}_3\text{--N=CH}_2$  occurs.

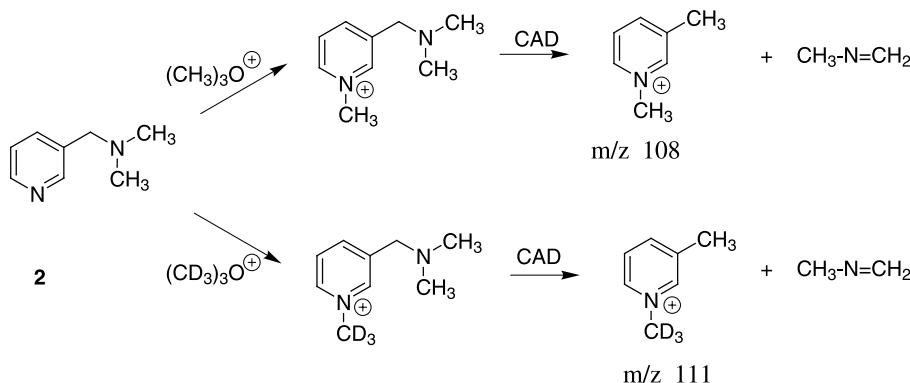
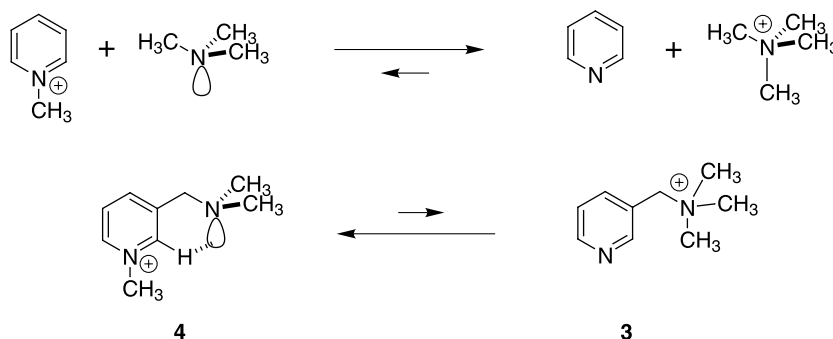
The loss of 43 u from either the  $(M+\text{CH}_3)^+$  or  $(M+\text{CD}_3)^+$  ions of **2** cannot be readily rationalized if deuteriomethylation had occurred on the  $sp^3$ -nitrogen to form **6**. In that case, a mixture of  $\text{CH}_3\text{--N=CH}_2$ ,  $\text{CD}_3\text{--N=CH}_2$  and  $\text{CH}_3\text{N=CD}_2$  fragments would have resulted from the  $(M+\text{CD}_3)^+$  ion, corresponding to loss of 43, 46, and 45 u, because the two methyl and one deuteriomethyl groups have equivalent chances of being eliminated when all are bound to the same nitrogen. The alkylating methyl group is not directly involved in the fragmentation process.

**6**

To evaluate the dichotomy between the gas phase and solution methylation reactions of **2**, we consider in turn gas phase and solution methylation reactions.

The isodesmic reactions shown in Scheme 3 were examined at the RHF/6-31G\*\*//6-31G\*\* level using Cadpac.<sup>5</sup> The reaction between methylpyridinium and trimethylamine is found to favor tetramethylammonium and pyridine by 12.2 kJ mol<sup>-1</sup>, corresponding to a ratio of more than 100:1 at room temperature in the gas phase. In contrast, ion **4** is calculated to be more stable than **3** by 21.4 kJ mol<sup>-1</sup>, corresponding to a room temperature/gas phase ratio of over 5000:1 in favor of **4**.

This contrast may in part be explained by the decrease in the basicity of the  $sp^3$ -amine in **2** and **4**, because of the presence of the electron withdrawing properties of the pyridine ring. In addition, calculations reveal a weak hydrogen bond between the  $sp^3$ -amine's lone pair of electrons and the C(2)–H and C(4)–H of the pyridine ring in **2** and **4**. This is seen by examining the preferred conformations of these and related model compounds, as determined by theory. The side chain of **3** is perpendicular to the plane of the pyridine ring, i.e.  $\tau_1[\text{C}(2 \text{ or } 4)\text{--C}(\text{ipso})\text{--C}_\alpha\text{--N}] = \text{ca. } 90^\circ$ , similar to that found for ethylbenzene,<sup>6</sup> propylbenzene,<sup>6</sup> and isobutylbenzene<sup>7</sup> **7**

**Scheme 2.****Scheme 3.**

and the monoprotonated **8** (see Fig. 1). However, in the pyridine ring methylated **4** and 3-(dimethylamino)-methylpyridine **2**, a very different conformation is found. Here, the lone pair on the side chain nitrogen points towards either the C(2)- or C(4)-hydrogen (see data in Fig. 1). The torsional angle found for *N,N*-dimethylbenzylamine **9** in this work is consistent with the literature experimental data and the highest level of theory in the earlier calculations.<sup>12</sup> The *sp*<sup>3</sup>-nitrogen lone pair is within hydrogen bonding distance to the C(2)- and C(4)-hydrogens. This is reminiscent of the formyl hydrogen bond recently identified by Corey et al.<sup>8,9</sup> and investigated computationally by one of us (J.G.).<sup>10,11</sup>

If this analysis is correct, the strength of the nitrogen lone pair–aromatic ring hydrogen interaction would increase as the polarization of the C(aromatic)–H bond increases. This would lead to a decrease in the torsional angle, as the acyclic nitrogen lone-pair–ring hydrogen interaction would be more able to counteract the steric effects pushing the side chain out of the plane. The calculated geometries are consistent with this interpretation. As the *ortho* aromatic-hydrogen increases its positive charge, the torsional angle shifts from perpendicular toward the plane of the aromatic ring, indicating an increased nitrogen lone pair–hydrogen bond. This trend is shown in Fig. 1. The nitrogen lone pairs in **4**, **2** and **9** are oriented to point in the direction of an aromatic hydrogen. The calculations show that for **4**, the nitrogen lone pair prefers to interact with the C(2)–H by ca. 5 kJ/mol. For **2**, it prefers the C(4)–H by ca. 2 kJ/mol. This switch in orientation is consistent with a polarization phenomenon, the interaction increasing with C(2)–H as the pyridine nitrogen becomes more positively charged.

The regiochemistry of the gas phase methylation of **2** may now be explained based on both kinetic and thermodynamic features: the *sp*<sup>3</sup>-amine's nucleophilicity is decreased by virtue of its weak hydrogen bonding to

the C(2)–H and presumably C(4)–H. In addition, as **4** is significantly more stable than **3**, thermodynamic considerations also favor pyridine methylation.

Acetonitrile is known to more effectively solvate pyridine than either ammonia or trimethylamine.<sup>7</sup> By analogy, acetonitrile would more effectively solvate the pyridine ring nitrogen than the dimethylaminomethyl nitrogen in **2**. In acetonitrile, pyridine methylation may be energetically less favored in part because pyridine desolvation must occur first.

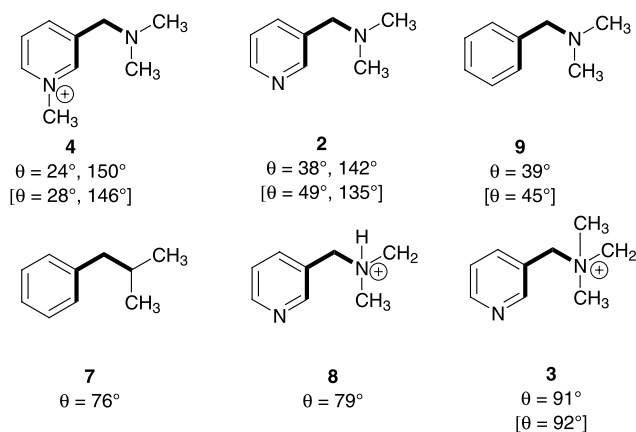
In conclusion, a reversal in regiochemistry is observed for gas-phase<sup>13</sup> and condensed methylations: A counterbalancing of effects explain the results.<sup>14</sup> In the gas phase, *sp*<sup>2</sup>-nitrogen methylation is explained by a weak hydrogen bonding interaction between the *sp*<sup>3</sup>-nitrogen lone pair and the C(2)- and C(4)-hydrogen atoms which acts against *sp*<sup>3</sup>-methylation. In acetonitrile, preferential pyridine solvation acts against *sp*<sup>2</sup>-nitrogen methylation.

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**Figure 1.** Torsion angles  $\tau_1(\text{C}_{\text{aromatic}}-\text{C}_{\text{ipso}}-\text{C}_{\alpha}-\text{N})$  or  $\tau_1(\text{C}_{\text{aromatic}}-\text{C}_{\text{ipso}}-\text{C}_{\alpha}-\text{C}_{\beta})$  from RHF/3-21G calculations and [in brackets] RHF/6-31G\*\* calculations.

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13. For a leading discussion of related gas phase reactions, see: Zhong, M.; Brauman, J. I. *J. Am. Chem. Soc.* **1996**, 118, 636–641.
14. The two reactions being compared herein (Scheme 1) differ not only in terms of the reaction medium, but also in terms of alkylating reagent and reaction type (i.e. formation of a salt from two neutral molecules versus reaction of one neutral molecule with a carbocation, to form another neutral molecule and another carbocation). These additional issues will be discussed in detail in our full paper.