

Towards the Strongest Neutral Organic Superbases Based on Intramolecular H-bonds[†]

Danijela Barić and Borislav Kovačević*

 Quantum Organic Chemistry Group, Division of Organic Chemistry and Biochemistry,
 Ruđer Bošković Institute, Bijenička 54, 10000 Zagreb, Croatia

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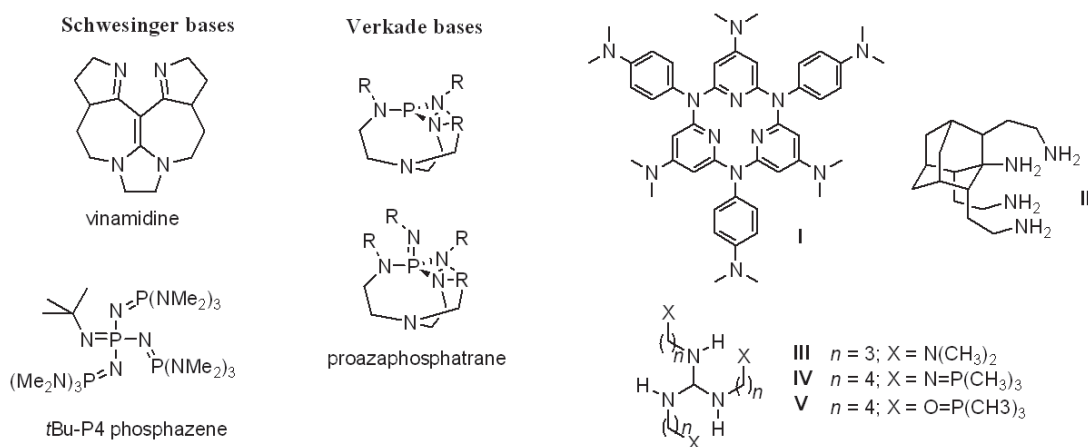
Abstract. Utilizing several different trialkylarsine oxides and substituted pyridine N-oxides as a hydrogen bond acceptors in tri-substituted guanidines we designed several very basic superbases possessing intramolecular hydrogen bonds (IHB-superbases), with proton affinity in the gas phase that comes very close to that of paradigmatic P4-*t*Bu Schwesinger superbase and with pK_a in acetonitrile up to 36 units.

Keywords: organic superbases, intramolecular hydrogen bonds, DFT calculations

INTRODUCTION

Neutral organic bases are important class of compounds in organic synthesis. Large amount of chemical reactions require deprotonation of starting compound as an initial process that activates reactants and enable chemical reaction to proceed. Various types of organic bases of different strength have been developed during the time, trying to achieve higher basicity sometimes needed to abstract the proton from a weak Brønsted acids. The most famous are Schwesinger vinamidines¹ and phosphazenes,² and Verkade proazaphosphatranes³ (Scheme 1). Several other types of molecular skeletons have been developed to achieve high basicity like pro-

ton sponges, or even some supramolecular cages (**I**, Scheme 1).⁴ Advances in the field of design and synthesis of organic bases have been summarized lately in two review articles.^{5,6} One particular molecular motif that leads to a highly basic organic compounds is substitution of guanidines and/or amines with substituents capable of forming intramolecular hydrogen bonds. Using this molecular motif Bachrach studied several very basic amines, with adamantane compound (**II**, Scheme 1) being the most basic one.⁷ More than 10 years ago we designed a substituted guanidine superbase (**III**, Scheme 1) where the basicity of central guanidine core – due to a presence of three intramolecular hydrogen bonds in conjugate acid – was



Scheme 1. Representative examples of some superbases.

[†] Dedicated to Dr. Mirjana Eckert-Maksić on the occasion of her 70th birthday.

* Author to whom correspondence should be addressed. (E-mail: boris@irb.hr)

increased by ≈ 25 kcal mol⁻¹ compared to *N,N,N'*-trimethylguanidine.⁸ Three years later this compound was synthesized and experimental measurements confirmed the proposed structure and basicity.⁹ Subsequently, some modifications of the original idea have been utilized to obtain more or less basic compounds. Glasovac synthesized compounds with methoxy¹⁰ and pyridino¹¹ group at the end of alkyl chain. Recently, utilizing the same motif combined with use of a very strong H-bond acceptors at the end alkyl chain, we designed a series of very strong neutral organic superbases ("IHB-superbases") with a gas phase proton affinity (PA) up to 293.3 kcal mol⁻¹ and p*K*_a in acetonitrile up to 33.2 (**IV**, **V**, Scheme 1).¹² To the best of our knowledge, these are the most basic guanidines designed so far. However, this brings up the question: did we reach the limit of basicity of the substituted guanidines with these compounds? Is it possible to find a stronger H-bond acceptor that would enable design of even stronger guanidines than those designed so far? In this paper we will show that it is feasible to find molecular moieties that can form a H-bonds stronger than alkyl phosphazenes – the strongest H-bond acceptors utilized in our previous study.

COMPUTATIONAL DETAILS

Calculations in the gas phase were carried out at B3LYP/6-311+G(2df,p)//B3LYP/6-31G(d) level of theory. The vibrational analysis was performed to confirm the energy minima on potential energy surface for all structures.

Proton affinities are calculated according to the Equation:

$$PA = H^{298}(\text{B}) - H^{298}(\text{BH}^+) + \frac{5}{2}RT \quad (1)$$

where $H^{298}(\text{B})$ and $H^{298}(\text{BH}^+)$ represent the enthalpies at 298 K of the neutral (B) and protonated (BH⁺) base, calculated at B3LYP/6-311+G(2df,p)//B3LYP/6-31G(d) level of theory. The p*K*_a values in acetonitrile (MeCN) are estimated using a correlation between the experimental data for p*K*_a values in MeCN and calculated basicities:¹³

$$pK_a = 0.545\Delta G'_{\text{a,sol}}(\text{BH}^+) - 133.5 \quad (2)$$

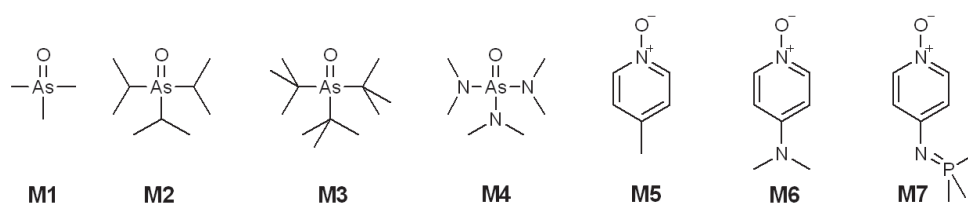


Figure 1. Model molecules used in complexes for estimation of the H-bond strength.

where $\Delta G'_{\text{a,sol}}(\text{BH}^+)$ represents the difference of Gibbs energies between product and reactants for the reaction of deprotonation of BH⁺ in acetonitrile. The Gibbs energy is sum of total electronic energy, the thermal correction to Gibbs energy and the energy of solvation. The solvation energies are calculated using isodensity polarizable continuum (IPCM) method at B3LYP/6-311+G(d,p)//B3LYP/6-31G(d) level of theory.

All computations were performed by the Gaussian03¹⁴ program package.

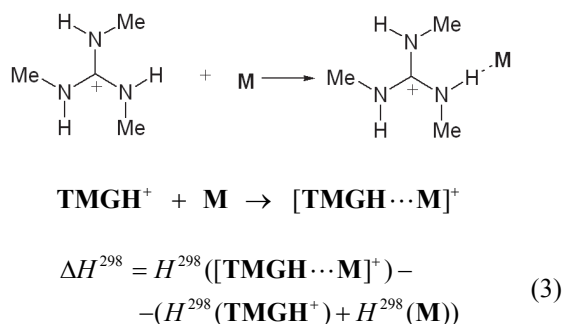
RESULTS AND DISCUSSION

As mentioned in introductory remarks, the strongest IHB-superbases obtained so far were designed utilizing alkylphosphazene, tris(dimethylamino)phosphine oxide, and trimethylphosphine oxide as an H-bond acceptor at the end of alkyl chain.¹² According to the experimentally determined p*K*_{BH} values,¹⁵ it can be expected that alkylsubstituted arsine oxide shows even stronger H-bond accepting ability. The p*K*_{BH} value of Et₃PO is 3.66, whereas Et₃AsO has a p*K*_{BH} value of 4.89. In the same compilation of p*K*_{BH} data it was shown that unsubstituted pyridine N-oxide is a very strong hydrogen bond acceptor (p*K*_{HB} = 2.70), almost as strong as 4-*N,N*-dimethylaminopyridine (p*K*_{BH} = 2.80). Substitution of pyridine N-oxide with strong electron donating group like dimethylamine at *para* position would probably significantly increase its H-bond acceptor strength, just as in the case of pyridine where the difference in p*K*_{HB} between unsubstituted pyridine (p*K*_{HB} = 1.86) and 4-*N,N*-dimethylaminopyridine (p*K*_{HB} = 2.80) is almost one unit.

Previously designed IHB-superbases **IV** and **V** (Scheme 1) consisted of the guanidine serving as a central motif in a superbase, and appropriate *N,N,N'*-substituents capable to make intramolecular hydrogen bond. To select proper substituents for IHB-superbases designed in this work, we calculated the energy of **intermolecular** hydrogen bond(s) in complexes built from protonated *N,N,N'*-trimethylguanidine and model molecules that possess the ability to form a strong H-bond. Our previous study¹² showed that the stabilization energy due to the presence of **intermolecular** H-bond(s) in complexes gave good prediction of basicity that could be obtained in the designed IHB-superbases. Investigated model molecules in this work were alkyl substituted

and dimethylamino substituted arsine oxides (**M1-M4**, Figure 1) since it is expected that they have stronger H-bond acceptor properties when compared to the corresponding phosphine oxides.¹⁵ Second group of model molecules were pyridine-N-oxides substituted with strong electron donating groups (**M5-M7**, Figure 1).

Firstly, the energy for monocomplexes ($\Delta H^{298}(\text{mono-})$) are calculated to compare the H-bond acceptor ability of new molecules with previously obtained ones. H-bond enthalpies for monocomplexes are calculated according to the Equation (3).

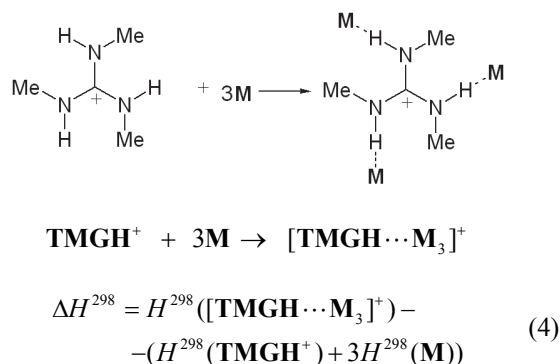


Obtained values of enthalpies for monocomplexes with pyridine N-oxides are in range of -19 to -29 kcal mol⁻¹, whereas enthalpies for monocomplexes with arsine oxides range between -20 and -24 kcal mol⁻¹ (Table 1). This implies that the stabilization upon complexation is better when compared with results from our previous study¹² where the absolute values of enthalpies for mono-complexes did not exceed 23 kcal mol⁻¹. Therefore, it can be expected that some of the IHB-superbases with substituents derived from model molecules **M1-M7** would be even more basic than those obtained previously. The conjugate acid of IHB superbases possesses three intramolecular hydrogen bonds that stabilize protonated central guanidine core.⁸⁻¹² It should be noted that according to the classification, this type of hydrogen bonds belongs to the charge assisted hydrogen bonds.¹⁶

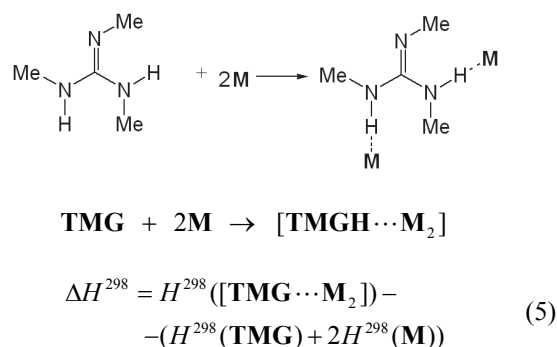
Table 1. The enthalpies of formation of mono-, tri- and di-complexes, and corrected values of enthalpy of tri-complexations

Molecule	$\Delta H^{298} / \text{kcal mol}^{-1}$			
	mono-	tri-	di-	tri-corr
M1	-23.8	-56.7	-7.3	-49.4
M2	-24.1	-56.0	-4.9	-51.1
M3	-23.7	-52.1	-2.6	-49.5
M4	-20.5	-48.4	-3.4	-45.0
M5	-19.7	-47.2	-6.5	-40.7
M6	-25.1	-57.8	-8.5	-49.2
M7	-29.1	-64.9	-9.4	-55.5

To estimate the stabilization energy of three H-bonds on protonated **TMG** cation, we calculated the stabilization enthalpies of tri-complexes ($\Delta H^{298}(\text{tri-})$) according to the Equation (4).



Additionally, it should be noted that neutral form of IHB-superbases possess two intramolecular H-bonds between substituents and two hydrogen atoms covalently bonded to nitrogen on central guanidine. Usually, these H-bonds are relatively weak, however, they contribute to the stabilization of neutral form of a molecule. We calculated the stabilization enthalpy, $\Delta H^{298}(\text{di-})$, between neutral **TMG** and two molecules **M** according to the Equation (5).



Since the proton affinity is the difference between the energy of neutral and protonated form of the base in question, the increase in PA due to a presence of H-bonds is a result of the difference in H-bond energy in neutral base and H-bond energy in the protonated one. Accordingly, we calculated the 'corrected' H-bond enthalpies $\Delta H^{298}(\text{tri-corr})$ as a difference between $\Delta H^{298}(\text{tri-})$ and $\Delta H^{298}(\text{di-})$ values. The enthalpies of formation of mono-, tri- and di-complexes, and corrected values of enthalpy of tri-complexations are presented in Table 1. We expect that the PA of newly designed superbases that are built with substituents based on model molecules **M** could roughly be the PA of **TMG** molecule plus the absolute value of $\Delta H^{298}(\text{tri-corr})$. Since the PA of **TMG**¹² is 249.9 kcal mol⁻¹, the PA values of new bases could be in range of 290 to 300 kcal mol⁻¹.

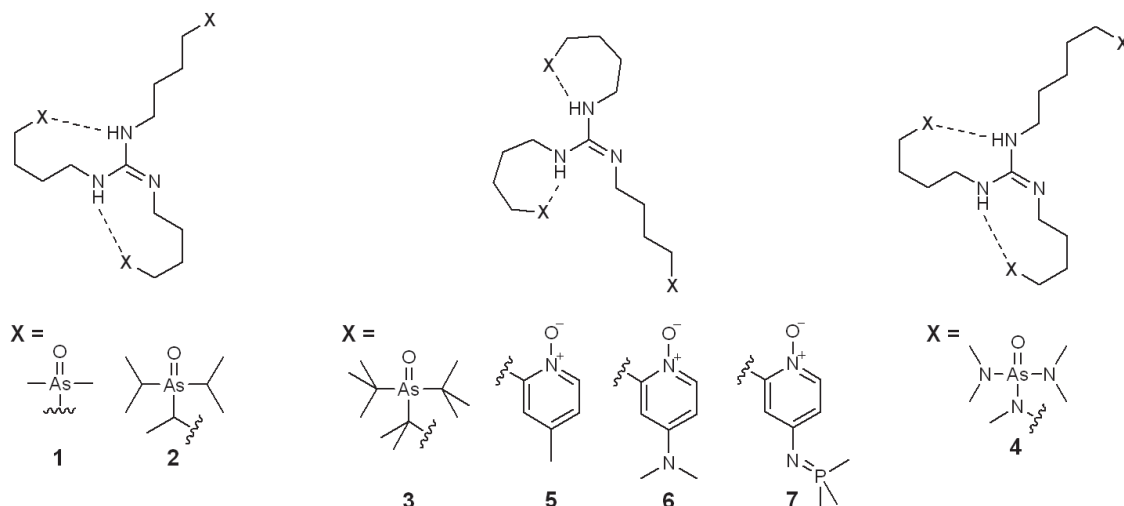


Figure 2. Schematic representation of newly designed IHB-superbases.

Once we selected an appropriate molecules **M** that should serve as H-bond accepting substituents, the next phase in designing of IHB-superbases was to find an appropriate alkyl chain that binds central guanidine core and selected substituent based on **M**. To find a proper chain, we applied the principle of minimal perturbation of arrangement obtained in tri-complexes. The alkyl chain was constructed in such a way to maximally preserve an 'ideal' orientation of molecules **M** around the central guanidine as it was in tricomplexes. The optimal alkyl chain for **1–3** and **5–7** was found to be buthyl, while for base **4** it was pentyl. The newly designed IHB-superbases are presented in Figure 2.

In molecules **3**, **5**, **6** and **7** the oxygen atom on substituents on guanidine forms an intramolecular H-bonds with H atom attached to the same nitrogen as substituent, thus forming pseudo seven-membered ring. In molecules **1** and **2** the oxygen atom on substituents forms a hydrogen bond with H atom attached to a neighboring N atom, forming an pseudo nine-membered ring. The same pattern is observed in base **4**, however, due to a longer alkyl chain, the pseudo ten-membered ring is assembled upon formation of intramolecular hydrogen bonds.

Table 2. The calculated gas phase proton affinities (PA) and pK_a values in acetonitrile for IHB superbases **1–7**

Superbase	PA / kcal mol ⁻¹	pK_a
1	293.1	32.1
2	296.9	36.7
3	296.0	32.7
4	290.9	-(a)
5	289.0	29.4
6	295.9	32.6
7	299.4	-(a)

(a) No convergence in the IPCM calculation.

The proton affinities and pK_a values for bases **1–7** are presented in Table 2. Inspection of data in Table 2 reveals that newly designed IHB-superbases possess gas phase PA between 289 and 299 kcal mol⁻¹, which is by 3 to 6 kcal mol⁻¹ higher than PAs obtained for IHB-superbases in our previous study.¹² However, the pK_a values are on average not significantly higher. Nonetheless, the superbase **2** with pK_a of 36.7 is by 3.5 units more basic than superbase **V** (Scheme 1) thus representing the most basic IHB-superbase designed so far.

As already mentioned, the effect of **intra**-molecular hydrogen bonds on PA of IHB-superbases can be roughly estimated by calculating the **inter**molecular H-bonds in complexes between model molecules **M** and **TMG**. However, a more specific analysis performed earlier¹² showed that intramolecular H-bonds are not the only factor that influences the basicity of IHB-superbases; the inductive effect of the bridging alkyl chain and the inductive effect of the H-bond accepting substituent at the end of the chain (**X**, Figure 2) also make significant impact.

We proposed that the total increase in PA (ΔPA_{total}) for each superbase compared to the corresponding reference molecule **TMG** may be represented as a sum of three influences: (i) the strength of intramolecular H-bonds (ΔPA_{IHB}), (ii) the inductive effect of alkyl chain (ΔPA_{alkyl}), and (iii) the inductive effect of the substituent **X** derived from the model molecules **M** (ΔPA_X):

$$\begin{aligned} \Delta PA_{total} &= PA(\text{superbase}) - PA(\text{TMG}) \\ &= \Delta PA_{IHB} + \Delta PA_{alkyl} + \Delta PA_X \end{aligned} \quad (6)$$

Here, ΔPA_{IHB} is obtained as a difference between the PA of superbases in folded conformation when H-bonds are present, and PA of unfolded (zig-zag) con-

Table 3. Analysis of various contributions to the proton affinities (PA) of IHB-superbases 1–7

Superbase	Proton affinities / kcal mol ⁻¹					
	PA	ΔPA_{total}	PA_{unfolded}	ΔPA_{IHB}	ΔPA_{alkyl}	ΔPA_{X}
1	293.1	43.2	255.8	37.2	6.7	-0.8
2	296.9	47.0	267.6	29.3	6.7	11.0
3	296.0	46.1	252.3	43.7	6.7	-4.3
4	290.9	41.0	259.2	31.8	7.5	1.8
5	289.0	39.1	253.6	35.5	6.7	-3.0
6	295.9	46.0	256.8	39.2	6.7	0.2
7	299.4	49.5	265.1	34.2	6.7	8.7

formers where intramolecular H-bonds do not exist. The term ΔPA_{alkyl} represents the difference in PAs between trisalkyl-substituted guanidine and the corresponding reference molecule **TMG**. As stated above, we found that for bases 1–3 and 5–7 the optimal chain was butyl whereas for base 4 the optimal chain is pentyl. The proton affinity of *N,N,N'*-trisbutylguanidine is 6.7 kcal mol⁻¹ higher than the PA of **TMG**, whereas the PA of *N,N,N'*-trisbutylguanidine is increased by 7.5 kcal mol⁻¹ compared to **TMG**.

The last term, ΔPA_{X} , which represents the contribution to the increase of PA by inductive effect of the H-bond accepting substituent **X** at the end of the chain, is calculated as a difference in PAs between the unfolded superbase and molecule of trisbutyl-substituted guanidine (PA = 256.6 kcal mol⁻¹) or trispentyl-substituted guanidine (PA = 267.4 kcal mol⁻¹). A detailed analysis for superbases 1–7 is presented in Table 3.

In all IHB-superbases the biggest contribution to the increase of PA are intramolecular H-bonds, represented by term ΔPA_{IHB} . The strongest influence of H-bonds is noticed in 3 and 6, however, they are not the most basic molecules here. The reason for that lies in the fact that the inductive effect of their substituents **X** (*tert*Bu-arsine-oxide and dimethylamino-pyridine N-oxide) are negative (being -4.3 kcal mol⁻¹ for molecule 3) and neutral (0.2 kcal mol⁻¹ for superbase 6). Since the ΔPA_{X} represents the contribution to the increase of PA due to the presence of substituent **X** when alkyl chain is unfolded, the negative value of ΔPA_{X} means that **X** withdraws electrons from the central guanidine and thus diminishes a proton affinity. At the same time, that positively influences the H-bond accepting ability of **X**, resulting in a relatively strong H-bonds as found in 3 and, in less extent, in 6.

The most basic is molecule 7, with phosphazeno-substituted pyridine-N-oxide as a substituent **X**. The positive inductive effect of **X** is strong, being almost 9 kcal mol⁻¹, which combined with also strong intramolecular H-bonds (≈ 34 kcal mol⁻¹) results in the biggest increase of PA when compared with **TMG**.

CONCLUSIONS

Based on the step by step approach in designing of IHB-superbases with the guanidine as a core fragment and heteroalkyl substituents, we designed seven new molecules that possess the gas phase proton affinity higher than those obtained so far. The substituents are chosen according to their H-bond accepting ability, that was firstly tested by building the complexes between model molecules and *N,N,N'*-trimethylguanidine. Alkyl chains bind the substituent with a central core and at the same time they enable optimal arrangement of the atoms which are involved in formation of H-bonds – *i.e.* the geometry of the complexes should be preserved in a great extent. The values of the PAs for designed IHB-superbases range from 289.0 to 299.4 kcal mol⁻¹, while pK_{a} values vary between 29.4 and 36.6 units. IHB-superbases owe their high basicity to the presence of three intramolecular hydrogen bonds in protonated form of the molecule, however, two more additional contributions have an influence to their PA. These contributions are the inductive effect of alkyl chains present in a molecule, and the inductive effect of the substituent **X**. The inductive effect of alkyl is always positive, but the inductive effect of the **X** depends on the character of the **X**. If **X** donates electrons through the alkyl chain toward the central core, then the basicity is additionally increased, as in superbases 2 and 7. In molecules 3 and 5, the **X** has negative inductive effect behaving as an electron acceptor that withdraws electrons from the central guanidine, thus having negative influence to basicity. On the other side, this electron-accepting behaviour of **X** results in a stronger H-bonds, so the final increase of PA is still preserved. For molecules 1, 4 and 6 the inductive effect of **X** is negligible and does not influence basicity.

We believe that superbases 1–7 should be synthetically available and we look forward for their preparation and application.

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