Bis(alkylguanidinium) Receptors for Phosphodiesters: Effect of Counterions, Solvent Mixtures, and Cavity Flexibility on Complexation

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Abstract: Four bis(guanidinium) receptors have been synthesized in which the guanidinium groups are spatially preorganized by an octahydroacridine (meso-3 and d,l-3) or hexahydrodicyclopenta[b,e]pyridine (meso-4 and d,l-4) spacer to complement a phosphodiester. These structures are designed to mimic the active site of staphylococcal nuclease and, thereby, form four hydrogen bonds to a bound phosphodiester with little reorganization of the host structures. The syntheses involve two parts: construction of the spacer and formation of the aminoimidazoline groups via an intramolecular cyclization between an amine and a thiouronium salt. Binding constants between the receptors and dibenzyl phosphate range from 4.0×10^3 to 10 M^{-1} in highly competitive solvent systems such as aqueous DMSO. Each receptor forms both a 1:1 and 2:1 phosphate to host complex. The methods for determining K_1 and K_2 are discussed in detail and involve both ³¹P and ¹H NMR titration experiments followed by a linear treatment of the data. Binding in pure DMSO is worth 3-4 kcal/mol, but the addition of water significantly decreases the degree of complexation. When the guanidinium counterions are tetraphenylboron, the meso forms of the hosts are the best receptors due to preorganization of the guanidinium groups on the same face of the spacer. When the counterions are chloride, the d, I forms can be the best receptors due to a specific ion effect where a chloride is involved in the host-guest complex. Addition of chloride salts increases binding, possibly due to a chaotropic "salting-out" phenomenon. The structures of the host-guest complexes of meso-4 with dibenzyl phosphate and phenyl phosphate have been determined by X-ray analysis. The structures demonstrate the chloride-counter ion assistance and confirm the four hydrogen bonds between the host and the guest. Near-identical structures to the crystal structures are calculated by molecular mechanics for the complex formed between dimethyl phosphate and meso-3 and meso-4. meso-3 has been found to act as an RNA hydrolysis catalyst and is the first step toward the optimization of a functional RNA-cleaving artificial enzyme.

Introduction

The starting point for the design of an artificial enzyme is often the examination of the active site of a natural enzyme. The active site of staphylococcal nuclease (SNase), a DNA- and RNAhydrolyzing enzyme, has several functional groups which contribute to catalysis.¹ Among these are two guanidinium side chains on arginines-35 and -87, a carboxylate of glutamate-43, and a calcium ion. The mechanism of catalysis and the energetic contribution from each of these functional groups have been studied in detail by single² and double-site-directed³ mutants. The guanidiniums are postulated to orient the substrate via twopoint hydrogen bonding and to act as general acids to assist in charge neutralization and leaving-group protonation.⁴ The guanidiniums are responsible for 10^{4,6}-fold out of the total 10¹⁶fold DNA hydrolysis rate enhancement.^{2,3} Guanindinium groups are also known to bind phosphates in other phosphatase enzymes,⁵ and several synthetic monoguanidinium⁶ and diguanidinium⁷ hosts have been developed to study such complexation.

In 1992 and 1993, three independent research groups synthesized artificial enzymes that possess two guanidinium-like moieties in an attempt to electrophilically activate phosphate linkages and thereby achieve a fraction of the rate enhancement imparted by the two arginines of SNase. Göbel reported that compound 1 increased the cleavage of a strained phosphodiester in DMF by a factor of 2000-fold.8 Hamilton reported that compound 2 increased the lutidine-catalyzed transesterification of a p-nitrophenyl-activated RNA analog by nearly 1000-fold in acetonitrile,9 and we reported that compound meso-3 at micromolar concentrations increased imidazole-catalyzed mRNA hydrolysis rates by 20-fold in water.¹⁰ Thus, mimicking the two arginine groups of SNase is a successful strategy for enhancing the rate of phosphodiester P-O bond cleavage under a variety of conditions.

The success of these bis(guanidinium)-SNase mimics resides in an initial binding of the monoanionic tetrahedral phosphodiester substrate followed by an increased electrostatic attraction with

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the dianionic trigonal bipyramidal phosphorane transition state.¹¹ The initial complexation in aqueous media is quite difficult due to competition with the water solvating the guanidinium groups and the phosphate substrate.¹² Since catalysis starts with substrate binding but aqueous media is highly competitive, our initial efforts were focused on maximizing the preorganization and the complementarity of bis(guanidinium) receptors for phospho diester substrates in order to maximize complexation.¹³ Herein, the effects of cavity flexibility, of hydrogen-bonding solvents, and of guanidinium counterions upon complexation strength are discussed.

Results and Discussion

Design Criteria. The design of receptors 3 and 4 has, as its starting point, the crystal structure of SNase with 3',5'-thymidine diphosphate (pdTp), performed by Cotton.¹⁴ Figure 1A is a schematic representation of the position of the catalytic groups found in this crystal structure. Of interest to this work are the two guanidiniums, each of which makes two hydrogen bonds to the dianionic phospho monoester pdTp. Cotton also found that the simple compounds methylguanidinium dihydrogen phosphate¹⁵ and bis(methylguanidinium) monohydrogen phosphate¹⁶ both crystallize to form a similar hydrogen-bonding pattern (Figure 1B). Even in these simple structures, each guanidinium forms two hydrogen bonds to the phosphoric acid moiety. Thus, we reasoned that a design using two guanidiniums in the manner shown in Figure 2 would lead to a successful phospho diester receptor and a possible hydrolysis catalyst.¹⁰

The spacer between the two guanidiniums was designed to converge the hydrogen-bonding groups to complement a bound phospho diester without significant reorganization of the receptor. An octahydroacridine spacer was chosen on the basis of the literature precedent with isophthaloyl spacers.¹⁷ These spacers, when incorporated in a macrocycle, properly orient hydrogenbonding groups for complexation of structures such as barbitu-

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Figure 1. (A) Schematic representation of the staphylococcal nuclease active site. (B) Simple guanidinium phosphate salt structures.



Figure 2. Idealized four-point hydrogen-bonding complexation geometry for bis(guanidinium) receptors with dibenzyl phosphate.

rates.¹⁷ In contrast to an isophthaloyl spacer, however, an octahydroacridine spacer holds the guanidinium groups preorganized and converged without using a macrocycle. Such preorganization has been found to be crucial in "molecular tweezers".¹⁸ In addition, preorganization should lead to a greater tendency for the guanidiniums to bind only one phosphate along a polyphosphate chain as in RNA or DNA. Octahydroacridine was used instead of acridine because of the basicity difference between bis(alkylguanidinium) groups versus that of bis-(arylguanidinium) groups. Alkylguanidiniums are desirable for catalytic studies in water because their pK_{as} (near 13)¹⁹ allow them to remain protonated over a wider pH range than those of arylguanidiniums (pK_a s near 9–10).²⁰ Finally, instead of attaching the parent guanidinium group (with all hydrogens on the nitrogens) to the spacers, aminoimidazoline groups were chosen. The two-point hydrogen-bonding capabilities are the same as those for guanidiniums, and their pK_as are similar, but the methylenes block hydrogen-bonding sites, thereby preventing multiple association like that found in similar systems.²¹

In addition to examining an octahydroacridine spacer in 3, a hexahydrodicyclopenta[b,e]pyridine spacer in 4 was explored.

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Scheme I. Module 1, Synthetic Routes to Spacers 5 and 6 A



The goal of this change was to examine binding characteristics as the angle and the rigidity between the aminoimidazolines changed with other characteristics of the receptor remaining the same. The angle between and the rigidity of the guanidinium groups are related to puckering of the cyclohexeno or cyclopenteno rings in 3 and 4. The cyclopenteno rings allow envelope-type puckering modes, whereas the cyclohexeno rings of 3 allow for each guanidinium group to be either axial or equatorial.

Synthesis. For kinetic studies on phosphodiester hydrolysis, fairly large quantities of the catalysts will be required. In addition, studies of the effects of different spatial positions of the guanidinium groups upon catalysis and the addition of more functional groups require a versatile synthesis. The synthesis involved two parts: the spacer and the aminoimidazolines. The *meso* and d,l forms of receptors 3 and 4 can be typically prepared on 1.5–3-g scales.

The syntheses of 3 and 4 focus initially upon constructing the rigid spacers 5 and 6 that hold the aminoimidazoline groups at specific distances and angles (Scheme I). The routes to diamines 5 and 6 are quite versatile in the sense that many different groups can be placed in the para position of the central pyridine and because the receptor size can be changed by using either cyclohexanone or cyclopentanone. The former allows for future functionalization while the latter allows for a subtle change in cavity size. The synthetic routes to these spacers are based upon extensive work from the Thummel²² and Bell groups.²³

The synthesis of 5 began with an enamine condensation of cyclohexanone with ethyl glyoxylate to form 7 followed by a reflux with ammonium acetate in acetic acid that afforded octahydroacridine 8. Activation of the benzylic positions was effected by refluxing 8 in acetic anhydride with benzaldehyde. Less than refluxing temperature resulted in the isolation of some monoactivated benzylic position with Z-double bond stereo-chemistry, thus blocking a second benzaldehyde addition. The product 9 was isolated by vacuum distillation of the solvent and excess benzaldehyde and crystallization from ethyl acetate to form large yellow plates. The overall yield for the three steps was typically near 25%. In contrast, these three steps can be combined in a one-pot reaction in the synthesis of 6. Refluxing benzaldehyde, cyclopentanone, and ammonium acetate in ethanol

for approximately 1 h formed a yellow crystalline precipitate of crude 10. After recrystallization from ethyl acetate, 10 was also isolated in 25% yield.²⁴

From the point of isolating 9 and 10, the syntheses of 3 and 4 only differ in the method of transforming diketones 11 and 12 to diamines 5 and 6. Both compounds 9 and 10 undergo smooth ozonolysis followed by reductive workup to form diketones 11 and 12. Compound 11 undergoes efficient reductive amination under the typical conditions²⁵ to form diamine 5. In contrast, diketone 12 when subjected to ammonium acetate and NaB-(CN)H₃ immediately turns black, and none of 6 can be isolated. Rather than pursuing other reductive amination conditions for transforming 12 to 6, a two-step method involving the intermediate dioxime 13 was utilized. Compound 13 was easily isolated by collection of the cream-colored precipitate formed upon warming 12, Na₂CO₃, and NH₂OH·HCl in DMF. Treatment of 13 with NaBH₄ and TiCl₄ in dry DME²⁶ effected the reduction to 6 in 50% yield.

The method of separation of the *meso* and d,l isomers of 5 and 6 is outlined in Scheme II. The *meso* and d,l isomers of both 5 and 6 can be observed to separate on a silica plate with 15% ammonia-saturated methanol in ethyl acetate eluent, but preparative flash chromatography proved difficult due to extensive streaking. On the basis of the assumption that the streaking was due to the highly polar nature of the diamines, a method involving a less polar derivative was pursued. The reaction of 5 and 6 with di-*tert*-butyl dicarbonate in chloroform cleanly gave 14 and 15, from which the *meso* and d,l forms could be easily separated by flash chromatography on silica gel with 50/50 ethyl acetate/hexanes eluent, the d,l isomer eluting first. Clean removal of the tBoc groups was accomplished by standard methods to yield the separated d,l and *meso* isomers of 5 and 6 in nearly quantitative yields.

The isomers were identified by X-ray crystal structures (as discussed below) for compound 4, but the ¹H NMR of the *meso* and d,l isomers of 5 revealed the appropriate stereochemistry of 3. Figure 3 displays the 500-MHz ¹H NMR spectra of the CH₂ moiety of the ethyl ester in both the *meso* and d,l forms of 5. In the *meso* form, the CH₂ group is not diastereotopic and the resonance is a clean quartet. In contrast, the CH₂ group in the d,l form is diastereotopic. The diastereotopic protons form an AB pattern that is further split by the methyl group to give a possible 16-line pattern. Only 12 lines are found due to coincidental overlap.

Formation of the aminoimidazoline groups employed an intramolecular five-membered-ring cyclization between an amine and a thiouronium salt (Scheme III) similar to intermolecular procedures.²⁷ This four-step route was adopted since the reaction of 5 or 6 with 2-(methylthio)-2-imidazoline (which would yield 3 and 4 directly) resulted in exceptionally low yields. The new aminoimidazoline-forming procedure centers around the synthesis of 16, which couples easily with amines to form thioureas. Compound 16 was synthesized by the monoprotection of ethylenediamine by di-tert-butyl dicarbonate followed by treatment with carbon disulfide and DCC.²⁸ The reaction of 5 and 6 with 16 gave high yields of 17 and 18, respectively. S-Alkylation of 17 and 18 followed by deprotection and base-mediated cyclization afforded 3 and 4, respectively, both in near 36% overall yield from their respective precursors 5 or 6. Purification of the free bases 3 and 4 proceeded by silica gel chromatography followed by formation of bis(picrate) salts. The picrates could be converted

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Scheme II. Strategy for Separation of the meso and d, I Isomers of 5 and 6



shift was determined independently. The ³¹P NMR chemical shift of HP could typically be determined to within 0.5 ppm by inspection of the binding isotherm since saturation binding was nearly always achieved. Thus, the variables were reduced to three: the two binding constants and the HP₂ chemical shift.

A method for determining 1:1 and 2:1 complexation constants from NMR data when following the chemical shift of the species whose stoichiometry is two in the 2:1 complex has been discussed.²⁹ We used a similar but slightly modified procedure. Equation 2

$$\delta_{obs} = ([P]/[P]_0)\delta_P + ([HP]/[P]_0)\delta_{HP} + 2([HP_2]/[P]_0)\delta_{HP_2}$$
(2)

shows the relationship of the observed chemical shift to the concentration of the complexes in solution where $[P]_0$ is the total phosphate concentration. If one solves for [P] from the K_1 and K_2 equilibria expressions as a function of total host $[H]_0$ and total phosphate $[P]_0$, eq 3 is obtained.²⁹ Equations 4 and 5 give [HP]

$$K_1 K_2 [P]^3 + K_1 (2K_2 [H]_0 - K_2 [P]_0 + 1) [P]^2 + (K_1 [H]_0 - K_1 [P]_0 + 1) [P] - [P]_0 = 0$$
(3)

$$[HP] = K_1[H]_0[P] / (K_1 K_2[P]^2 + K_1[P] + 1)$$
(4)

 $[HP_2] = K_1 K_2 [H]_0 [P]^2 / (K_1 K_2 [P]^2 + K_1 [P] + 1)$ (5)

and [HP₂], respectively, as a function of [P]. The cubic eq 3 was solved by the Newton-Raphson technique for multiple [H]₀s and with several initial estimates of K_1 and K_2 . Each [P] solved for from eq 3 was used to solve eqs 4 and 5. The resultant [P], [HP], and [HP₂] were substituted into eq 2. The variables K_1 , K_2 , and δ_{HP_2} were visually changed to optimize the theoretical fit to the data. It was found that the use of a nonlinear least-squares fitting routine based upon the simplex algorithm³⁰ did not give significantly different results than those from a visual fit.

The HP₂ complex under all our experimental conditions possessed a lower field chemical shift than free dibenzyl phosphate, whereas the 1:1 complex possessed a higher field chemical shift. Figure 5 and 6 show theoretically generated ³¹P NMR binding isotherms generated by the method discussed above. In Figure 5, the chemical shifts of P, HP, and HP₂ are held constant while

Figure 3. ¹H NMR spectra of the methylene ethyl hydrogens of *meso*-5 and d_i -5.

to chlorides by anion-exchange chromatography, and the chlorides could be converted to tetraphenylborates by a metathesis with sodium tetraphenylborate in acetonitrile or water.

Complexation Studies. Methods. The binding constants between receptors 3 and 4 were primarily determined from ³¹P NMR titrations. The dibenzyl phosphate concentration was held constant, and the concentration of the receptor was incrementally changed to span a host to guest ratio between 0 and 6. In low percentages of H₂O in DMSO, curves were generated that indicated 1:1 and 2:1 phosphate (P) to host (H) complex formation. As water percentages approached and exceeded 20, only 1:1 phosphate to host complexation was observed. Two representative binding isotherms are shown in Figure 4. The inflections in the isotherms are indicative of the two equilibria shown in eq 1. Since

$$H + P \stackrel{K_1}{\longleftarrow} HP \quad HP + P \stackrel{K_2}{\longleftarrow} HP_2 \qquad (1)$$

the isotherms begin with dibenzyl phosphate in excess, the first complex is HP_2 which exists in highest concentration near 0.5 equiv of host. As the host concentration increases, the equilibria

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Scheme III. Module 2, Synthetic Routes for Construction of the Aminoimidazoline Groups



Figure 4. Typical ³¹P NMR binding isotherms: $\triangle = 1.15 \times 10^{-2}$ M dibenzyl phosphate in 15% D₂O/DMSO-d₆ with increasing meso-4 and $\bigcirc = 8.6 \times 10^{-3}$ M dibenzyl phosphate in 15% D₂O/DMSO-d₆ with increasing d,l-4. The lines drawn are the calculated fits. The y-axis errors represent $\pm 15\%$ estimated error in determining the host/guest ratio. The error in reading the chemical shift is within the size of the marker.



Figure 5. Theoretical ³¹P NMR binding isotherms. All curves have $\delta_P = 0$ ppm, $\delta_{HP} = -1$ ppm, and $\delta_{HP_2} = 1$ ppm. (A) $K_1 = 5000 \text{ M}^{-1}$, $K_2 = 1000 \text{ M}^{-1}$; (B) $K_1 = 1000 \text{ M}^{-1}$, $K_2 = 1000 \text{ M}^{-1}$; (C) $K_1 = 1000 \text{ M}^{-1}$, $K_2 = 100 \text{ M}^{-1}$; (D) $K_1 = 5000 \text{ M}^{-1}$, $K_2 = 100 \text{ M}^{-1}$; and (E) $K_1 = 200 \text{ M}^{-1}$, $K_2 = 20 \text{ M}^{-1}$.

the magnitudes and the ratios of K_1 and K_2 are varied. With line A, both K_1 and K_2 are relatively large and only 5-fold different from each other. This creates a binding isotherm that is slow to

Figure 6. Theoretical ³¹P NMR binding isotherms. All curves have K_1 = 1000 M⁻¹, K_2 = 400 M⁻¹, δ_{HP} = -1 ppm, and δ_P = 0 ppm. (A) δ_{HP_2} = 2 ppm, (B) δ_{HP_2} = 1 ppm, (C) δ_{HP_2} = 0.5 ppm, (D) δ_{HP_2} = 0 ppm, (E) δ_{HP_2} = -0.5 ppm, (F) δ_{HP_2} = -1 ppm, and (G) δ_{HP_2} = -2 ppm.

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plateau as a result of a significant amount of the HP₂ complex being present over a wide range of concentrations. Lines D and E represent ratios of K_1 to K_2 of 10 and 50, respectively, with a relatively low K_2 binding constant. In these cases, there is very little inflection in the isotherms found at 0.5 equiv of host, giving as a result a quick plateau. Thus, visual examination of the experimentally determined isotherms yields insight into the relative ratios of the binding constants. In Figure 6, the magnitudes of K_1 , K_2 and δ_{HP} are held constant and δ_{HP_2} is varied. Experimental examples similar to these theoretical curves yield insight into the chemical shifts of HP and HP₂.

Figure 4 shows two representative experimental isotherms (not our best nor our worst matches between experiment and theory) for the *meso* and *d*,*l* isomers of 4 in 15% aqueous DMSO. The differences in binding characteristics between the two isomers are evident by inspection. The isotherm for the *d*,*l* isomer shows a downfield shift initially followed by an upfield plateau. In contrast, the isotherm for the *meso* isomer shows a continuous upfield shift. One explanation for the difference between the *meso* and *d*,*l* isotherms at 0.5 host to guest ratio is that both K_{2S} are similar but δ_{HP_2S} are different. A confirmation of these chemical shift differences will be discussed with respect to Figure 6. In addition, the curve for the *d*,*l* isomer is still moving upfield while the curve for the *meso* isomer has nearly reached a plateau at a host to guest ratio of 6. Therefore, K_1 is larger for the *meso*



 $\chi_{HG} / \Delta \delta_{obs}$

Figure 7. Linear treatment (eq 6) of the experimental data given in Figure 4: \blacksquare , data from *meso*-4, and \blacktriangle , data from *d*,*l*-4.

compound on the basis of the higher extent of saturation at 6 equiv of host. These observations are confirmed by theory which yields $K_1 = 550$ and 400 M⁻¹ for *meso*-4 and *d*,*l*-4, respectively, and K_2 for both isomers being 40 M⁻¹ (Table II).

In all the cases studied, the K_1 binding constant is a reliable value but complications can arise with respect to K_2 when there is no prior knowledge of the magnitude of K_2 and δ_{HP_2} . In these cases, the same experimental binding isotherm can be fit if one K_2 is arbitrarily chosen and the δ_{HP_2} is optimized or vice versa. The isotherms, however, could not be fit with significantly different K_1 s. The magnitude of K_2 , however, can be determined by a ¹H NMR titration binding experiment.

To confirm the binding constants measured by the ³¹P NMR technique, several were also measured by ¹H NMR titrations. In Tables I and II (discussed below), the binding constants that were confirmed by ¹H NMR are marked with an asterisk. In these experiments the host concentrations were held constant and either the guanidinium hydrogen resonances or the aminoimidazoline methylene resonances were followed as a function of increasing concentrations of dibenzyl phosphate. The algorithm for fitting a 1:1 and 2:1 NMR isotherm when following the chemical shift of the species whose stoichiometry is one has been discussed.³¹ The agreements with the binding constants obtained by the ¹H NMR method were typically within 15% or less of those obtained by the ³¹P NMR method. The agreement with the ³¹P NMR data was best for the meso forms of 3 and 4, for which little scatter in the experimental proton data was observed. However, for the d,l forms, the chemical shift of the protons changed less and therefore possessed more scatter and the absolute values were not as reliable. However, it is clear that the general trend as measured by ³¹P NMR is confirmed by the ¹H NMR.

As a further confirmation of the validity of the binding constants reported in Tables I and II, a simple algebraic manipulation of eq 2 can be performed that creates a linear form giving a check on K_1 and K_2 .²⁹ If we define $\Delta\delta s$ as the change in chemical shift from that of δ_P and χ as mole fractions, then eq 2 can be transformed to eq 6. Substituting the mole fractions of HP and

$$\frac{(1/\Delta\delta_{\rm HP_2}) - (\Delta\delta_{\rm HP}/\Delta\delta_{\rm HP_2}) (\chi_{\rm HP}/\Delta\delta_{\rm obs}) = 2(\chi_{\rm HP_2}/\Delta\delta_{\rm obs}) \quad (6)}{b - m \quad x = y}$$

HP₂ (calculated for each [H]₀ and [P]₀ by using the K_1 and K_2 determined by the Newton-Raphson technique) into eq 6 and plotting ($\chi_{HP}/\Delta\delta_{obs}$) versus $2(\chi_{HP_2}/\Delta\delta_{obs})$ should yield a straight line with $(1/\Delta\delta_{HP_2})$ and $(\Delta\delta_{HP}/\Delta\delta_{HP_2})$ as the intercept and the slope, respectively. Figure 7 shows the results of checking the data from Figure 4. As discussed above, the curves of Figure 4 suggest different δ_{HP_2} values but similar δ_{HP} values. This is

confirmed by the fact that the theoretical fit to the experimental curves of Figure 4 yields lines by the eq 6 treatment and that the y intercepts representing $(1/\Delta\delta_{HP_2})$ are quite different. Thus, the overall procedure for measuring the binding constants involved independent ¹H and ³¹P NMR titrations followed by linearizing the ³¹P NMR data to eq 6.

The binding studies that involved simple one to one complexes were performed by following the ³¹P NMR dibenzyl phosphate chemical shift as a function of the concentration of various receptors. The data were then fit to the conventional 1:1 binding algorithm that has been previously discussed.³² On the basis of repetitive measurements of the binding constants by either the ¹H or ³¹P NMR technique, we estimated at worst a 25% error on all the binding constants reported.

Effect of Host Cavity Size and Counterions. Table I lists the binding constants between the meso and d,l forms of 3 and 4 with dibenzyl phosphate. In pure DMSO, both hosts show stronger binding in the meso form than in the d, l form when the counterion is tetraphenylborate. This suggests that the aminoimidazolines are acting cooperatively in the meso forms to complex the phosphate and that the cooperativity is at a maximum when the aminoimidazolines are positioned on the same face of the spacer. meso-3 shows stronger binding than meso-4. This supports a more complementary cavity with 3 than with 4. We postulate that the octahydroacridine spacer in 3 is more flexible than that in 4 and that this flexibility allows the cavity to adjust to the phosphodiester substrate. The difference, however, between the meso forms of 3 and 4 is less than 0.3 kcal/mol, and thus both structures are quite comparable. In contrast, both K_1 and K_2 for the d,l form of 4 are similar, indicating that the guanidiniums act independently. Thus, the spacer in 4 does not have enough flexibility to allow the guanidiniums of both the meso and d,lforms to adopt similar spatial positions when binding a phosphodiester. The d,l form of 4 must therefore restrict the aminoimidazoline groups to be on different sides of the spacer and to not be appropriately positioned to act cooperatively to complex the phosphodiester.

The effect of these differences of flexibility and the relative sizes of the binding constants on the ability of these receptors to act as phosphodiester hydrolysis catalysts has not yet been determined. It will be interesting to discover if the more flexible receptors are better catalysts due to better accommodation of the geometry change between tetrahedral substrates and trigonal bipyramidal-like transition states or if the more rigid *meso-4* will be better due to a wider cavity that is more complementary to the expanded coordination sphere of the transition state.

Cooperativity for phosphodiester binding between the aminoimidazolines in *meso-3* and *meso-4* is further supported by comparison with the binding constant of dibenzyl phosphate to **19**. K_1 , using either chloride or tetraphenylborate counterions, is 350 M⁻¹. In comparison, compounds **3** and **4** have binding constants greater than that of **19**. This supports the postulate



that the guanidinium groups of 3 and 4 are acting cooperatively to bind the dibenzyl phosphate guest and that a 0.8-1.1 kcal/mol advantage is gained by preorganizing two guanidiniums in a binding array similar to that of SNase (calculated from the tetraphenylborate salts of 3 and 4). This energy advantage can result from either a chelation effect due to two additional hydrogen bonds formed from the second guanidinium or binding to one

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Table I. K_1 and K_2 Binding Constants for Dibenzyl Phosphate with the Different Isomers of 3 and 4 in DMSO, Solved for by the ³¹P NMR Titration Method^a

conditions	compound							
	meso-3.2Cl-	d,l-3-2Cl-	meso-4.2Cl-	d,1-4-2C1-	meso-3*·2B(Ph)4-	<i>d</i> , <i>l</i> -3·2B(Ph) ₄ -	meso-4*.2B(Ph)4-	d,l-4*·2B(Ph)4-
0% H ₂ O/100% DMSO	2.2×10^{2} 1.8×10^{2}	2.5×10^{3} 1.0×10^{2}	4.0×10^{3} 1.0×10^{2}	1.0×10^{3} 4.0×10^{2}	2.2 × 10 ³ 10	8.0×10^2 10	1.4×10^{3} 30	9.6×10^2 8.8×10^2

^a Those entries marked with an asterisk indicate that the binding constants were verified by ¹H NMR titrations. All M⁻¹. Error estimated at 20%.

guanidinium with a neighboring second positive charge. We prefer the former explanation based upon crystal structures and molecular mechanics (discussed below).

The term cooperativity as used above is not meant to imply binding above twice the ΔG value for 19 but only an increase in binding with 3 and 4 compared to that in 19. The addition of the ΔG for 19 (3.4 kcal/mol) would predict a binding constant of 9.7×10^4 M⁻¹ (6.8 kcal/mol) for 3 and 4 in DMSO. Such a large increase, however, is not predicted since the second guanidinium does not form an ion pair with the phosphodiesters as does the first guanidinium. Chelation by the second guanidinium provides only two more hydrogen bonds to the complex. Such hydrogen bonds are not expected to contribute greatly to binding in highly competitive solvents such as DMSO and water, and apparently, the hydrogen bonds increase binding by only about 1 kcal/mol as discussed above. The bis(alkylguanidinium) receptors are, however, designed to achieve a significant chelate effect in the binding of dianionic phosphorane intermediates along the hydrolysis reaction pathway. In contrast to the binding of the monoanionic substrates, catalysis of phosphodiester hydrolysis using receptors 3 and 4 would be expected to benefit from a large increase in binding in the transition state since the phosphorane intermediate is formally minus two and thus a full second ion pair is formed in the transition state that is lacking in the ground state. We are currently working to verify this prediction.

Since both the chloride and the noncoordinating tetraphenylboron salts of 19 bind dibenzyl phosphate similarly in DMSO, the chloride counterion must not compete for complexation with the guanidinium. In contrast, complexation displays unusual results with the chloride salts of the bis(guanidinium) receptors. For the chloride salts, meso-4 binds more strongly than d, l-4 but d.l-3 binds more strongly than meso-3. This is a different trend than that observed for the tetraphenylborate salts. In addition, the chloride salts of 3 and 4 typically have larger association constants with dibenzyl phosphate than the tetraphenylborate salts. This at first seems counterintuitive since the noncoordinating tetraphenylborates should present no competition in coordinating to the guanidiniums and thus these counterions should yield the largest binding constants for the guanidiniums to the dibenzyl phosphate. The results, however, clearly indicate that the chloride counterion is assisting the phosphodiester binding. As will be discussed below with respect to the effect of added salts and structural studies, the chloride counterion appears to be intimately associated in the 1:1 host to guest complex, yielding a specific ion effect that assists phosphodiester complexation.

Effect of Solvent System. Binding studies between 3 or 4 with dibenzyl phosphate in different ratios of DMSO/water were performed. In percentages of water between 0 and 15, 2:1 binding was observed (Table II). At 20% water, the 2:1 binding vanished and only 1:1 binding was observed which decreased with increasing percentages of water. At 40–50% water, the binding constants became negligible (<10 M⁻¹). Weak binding is, however, still observed for meso-4 (K_1 near 30 M⁻¹) even in 50% H₂O/DMSO. DMSO is a strong hydrogen-bond acceptor and thus competes with the phosphodiester for hydrogen bonding to the hosts,³³ but H₂O is able to solvate both the host and the guest by hydrogen-bond accepting and donating, respectively. In addition, the higher

Table II. K_1 Binding Constants for Dibenzyl Phosphate with the Different Isomers of 3 and 4 (either Cl⁻ or BPh₄⁻ counterions) in Various Percentages of D₂O in DMSO- d_6^a

conditions	compound						
	meso- 4 . 2Cl ⁻	<i>d,1-</i> 4 . 2C1 ⁻	meso-3. 2B(Ph)4	<i>d,l-3</i> . 2B(Ph) ₄ -	mix-3*. 2Cl ⁻		
5% H ₂ O	2.0×10^{3} 6×10^{1}						
10% H ₂ O					3.0×10^2 6×10^1		
15% H ₂ O	5.5×10^{2} 4×10^{1}	4.0×10^{2} 4×10^{1}					
20% H ₂ O			7×10^{1} 7×10^{1}	1.0×10^{2} 1×10^{1}	6×10^{1} 2 × 10 ¹		
25% H ₂ O	3.3×10^2 6.5×10^1						
33% H ₂ O 50% H ₂ O	6×10^{1} 3×10^{1}				3 × 10 ¹		

^a All M⁻¹. Error estimated at 10-20%.



Figure 8. K_1 complexation constants between 3 and dibenzyl phosphate as a function of the ionic strength in (67:33) DMSO/H₂O: \blacktriangle = LiCl, \square = NaCl, \bigcirc = KCl, \blacksquare = NaClO₄, and \bigcirc = NaSCN.

dielectric effect of water tends to shield the electrostatic attraction of the host and the guest better than that of DMSO.

Effect of Added Salts. Binding studies with a 1:1 mixture of *meso-3* and *d,l-3* and dibenzyl phosphate were performed with varying ionic strengths. The effect of the ionic strength on the phosphodiester chemical shift was examined prior to this experiment because changes in the ³¹P NMR chemical shift with ionic strength have been reported.³⁴ A typical change in ionic strength during a binding study was estimated at ~0.02 M. Chemical shift changes due to this ionic strength change were found to be smaller than 0.012 ppm, which is much smaller than the 1–2 ppm change due to complexation between host and guest.

The results of binding studies with several salts (LiCl, NaCl, KCl, NaClO₄, and NaSCN) in DMSO/H₂O (67/33) are shown in Figure 8. All systems showed only 1:1 binding in this high percentage of water. The effect of added salts seemed to be classified into two categories. One category includes LiCl, NaCl, and KCl, which increased the host-guest binding. The other is NaClO₄ and NaSCN, which hindered the host-guest binding. These results cannot be interpreted as a dielectric effect because

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Figure 9. X-ray crystal structure of *meso-4* with chloride and dibenzyl phosphate counterions.

an increase in ionic strength should cause a decrease in the electrostatic attractive force between host and guest with all five salts. These results, however, could be interpreted by a salting-in and salting-out chaotropic phenomenon. LiCl, KCl, and NaCl are known as salting-out salts,35 in which the structure of water is more icelike and therefore a worse solvent for organic groups. In contrast, NaClO₄ and NaSCN are known as salting-in salts³⁵ which destroy the water structure and increase the interaction between organic compounds and water. Evidence for a chaotropic effect on the binding-constant trend is the experimental observation that LiCl increases binding more than NaCl, which in turn increases binding more than KCl. This is the expected trend if the best salting-out compounds would increase binding the most. Chaotropic effects, however, are not the only interactions that lead to increased binding with added chloride salts. Table I shows that the dichloride salts of 3 and 4 bind dibenzyl phosphate better than the tetraphenylborate salts in DMSO. This observation, taken along with the increasing binding constants with increasing NaCl and KCl concentrations, gives evidence that the chloride is part of the 1:1 host to phosphate structure, either coordinated to a guanidinium or in the solvent sphere for charge neutrality. Structure studies (discussed below) support a structure in which the chloride is coordinated to a guanidinium with a water molecule bridging the chloride and the phosphate (Figure 9). The reason for the intimate association of the chloride is the nature of the solvent. DMSO is a good solvator³³ of cations but not of anions, and as a result, association of the chloride to a guanidinium yields charge neutrality as well as hydrogen-bonddonor solvation of the chloride.

Alternatively, chloride may be selectively and preferentially solvated by water, as could Li, Na, and K, thus making the binary solvent system more DMSO-like and increasing the binding constants. Such hetero- and homoselective solvations have been observed previously.³⁶ Since salting-out phenomenon are typically associated with hydrophobic compounds, however, it is quite interesting to find a trend between the salting-out ability and the aggregation of charges. Clearly this effect deserves further study.

Incorporation of a chloride into the receptor-substrate complex should be detrimental to the catalysis of phosphodiester hydrolysis.

Table III. Hydrogen Bond Distances and Angles for the Structure Shown in Figure 9^a

D-H···A	DA	Н…А	D-H-A
N9-H9-02	2.843(5)	1.945(5)	175.4(4)
N14-H14-O1	2.769(5)	2.052(5)	135.8(4)
N15-H15-01W	3.012(4)	2.326(4)	132.9(4)
N17–H17…C11	3.168(4)	2.309(4)	159.7(5)
O1W-HWA-C11	3.212(5)	2.208(5)	164.0(4)
O1W-HWB-O2	2.881(5)	1.790(5)	153.3(4)

^a The hydrogen atom positions were calculated, and the esd's for the parameters involving them are underestimated.

The chloride forms the second ion pair in the ground state that is postulated to be the driving force for increased binding of the dianionic transition state and the phosphorane intermediate. Since the catalysis strategy resides in complexing both guanidiniums to one phosphodiester substrate, the incorporation of another anion into the complex is a complication to catalysis that must be continually monitored.

Structural Studies. The host-guest complexes are less soluble in water than in DMSO, such that in 30% water, the meso and d,l forms of 3 or 4 and dibenzyl phosphate precipitate from solution, and are quite insoluble in 50% water. X-ray data revealed a monoclinic crystal of space group $P2_1/c$ with a = 18.010(8), b = 10.708(3), and c = 19.989(7) Å, $\beta = 104.83(3)^{\circ}$, R = 0.0664, and a goodness-of-fit being 1.597. The crystal structure of meso-4 and dibenzyl phosphate shows the phosphate bound by two hydrogens of one guanidinium moiety, while a chloride ion is bound to the other guanidinium (Figure 9). The chloride is required in the lattice for charge neutrality. A water molecule bridges the phosphate to the chloride. This structure is likely to exist in the solution state and explains the increased 1:1 binding with the chloride salts relative to tetraphenylborate (Table I) and the increased K_1 s with increasing concentrations of NaCl and KCl (Figure 8). Table III shows the hydrogen bond angles and distances within the host-guest complex. The incorporation of the chloride and the water in the crystal structure leads to a widening of the cavity spreading apart the aminoimidazoline moieties. This forces a binding arrangement between the phosphodiester and one of the aminoimidazolines which is nonoptimal. The hydrogen bonds between these groups have heteroatom to heteroatom distances between 2.85 and 2.77 Å, but the N14-H14-O1 bond is severely bent with a N-H-O angle of only 136°. The hydrogen bonding between the water and the aminoimidazoline is also nonoptimal with a heteroatom distance of 3.012(4) Å and a bond angle of 133°. The hydrogenbond donations from the water to the chloride and the phosphodiester are both near 160°.

To avoid the requirement of two counterions in the crystal lattice of 3 or 4 with dibenzyl phosphate, crystals were grown of a complex between meso-4 and a dianionic phosphomonoester, phenyl phosphate (Figure 10). The X-ray data revealed a monoclinic $P2_1/c$ space group with a = 10.578(5), b = 8.838(4), and c = 32.07(2) Å, $\beta = 96.11(4)^{\circ}$, R = 0.0579, and a goodnessof-fit being 1.343. The crystal structure of meso-4 with phenyl phosphate shows the binding expected for a phosphoester, with two-point bonding from each guanidinium moiety to the guest, as is found in SNase and simple phosphate-guanidinium salts (Figure 1). All the heteroatom-heteroatom hydrogen bond distances between the host and the guest are less than 2.9 Å (Table IV). The hydrogen bond angles are also near optimal, being at the most only 11° from 180°. Thus, the spacers used in 3 and 4 position the aminoimidazoline groups complementary to a tetrahedral phosphoester.

The structure between *meso-4* and phenyl phosphate is particularly important for the studies of phosphodiester hydrolysis, not only because it demonstrates host-guest complementarity and the predicted four hydrogen bonds but also because the phenyl phosphate may serve as a suitable model for the dianionic

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⁽³⁶⁾ Reichardt, C. Solvent and Solvent Effects in Organic Chemistry, 2nd ed.; VCH: Weinheim, Germany, 1988; pp 35-38.



Figure 10. X-ray crystal structure of meso-4 and phenyl phosphate.

Table IV. Hydrogen Bond Distances and Angles for the Structure Shown in Figure 10^a

D-H-A	DA	HA	D-H-A	
N9-H9-02	2.868(6)	2.11(5)	178(5)	
N11-H11-01	2.751(6)	1.84(5)	172(4)	
N15-H15-O2	2.840(6)	1.95(5)	169(4)	
N17-H17-O3	2.658(7)	1.69(6)	176(5)	

^a The hydrogen atoms were found and refined.



Figure 11. Superposition of *meso*-4 from the crystal structure with bound chloride and dibenzyl phosphate (---) and phenyl phosphate (---), revealing the different puckering modes and bond rotations.

intermediate and the transition state for phosphodiester hydrolysis. Unfortunately, we have not yet measured complexation constants for phosphomonoesters to 3 and 4 due to immediate precipitation upon mixing host and guest in a variety of solvent systems.

The structure of meso-4 with a bound chloride and phosphodiester demonstrates that the receptors possess enough flexibility to bind two separate counterions. The cavities of both 3 and 4 can widen via puckering modes of the cyclohexeno and cyclopenteno rings. Another degree of freedom involves rotation along the aminoimidazoline bond to the spacers. The widening of the cavity with meso-4 to accommodate two anions not only requires a different puckering within the cyclopenteno rings than that used in binding phenyl phosphate but in addition requires that the bond between the spacer and the aminoimidazoline rotate. Both these adjustments within the receptor are shown by the crystal structure superposition given in Figure 11. As discussed above, the tetraphenylborate-meso forms of 3 and 4 better accommodate the phosphodiester guest than the d,l forms because the guanidinium groups are already preorganized on the same face of the spacer. Conversely, the dichloride d, l form of 3 is the better phosphodiester host relative to the meso form since a binding



Figure 12. Molecular mechanics-derived host-guest structures: (A) meso-4 and dimethyl phosphate and (B) meso-3 and dimethyl phosphate.

geometry involving both a phosphate and a chloride requires the cavity to be wider. This mode of complexation is most easily accommodated by the d,l form of 3, since the guanidiniums are already on different faces of the spacer and can more easily accommodate two anions.

X-ray quality crystals of dibenzyl phosphate bound in the *meso* forms of 3 and 4 with a tetraphenylborate counterion were not obtainable. We postulate on the basis of molecular mechanics calculations,³⁷ however, that the binding mode found for phenyl phosphate and *meso-*4 (Figure 10) represents the most probable structure for *meso-*3 and *meso-*4 with dibenzyl phosphate. The calculated structures are shown in Figure 12. The puckering in the cyclopenteno rings of 4 of these molecular mechanics-derived structures is the same as that found in the crystal structure with phenyl phosphate, and the hydrogen bond lengths vary by less than one tenth of an angstrom between the calculated and experimental structures. No crystal data are available for *meso-*3 as a comparison, but the molecular mechanics again predicts four hydrogen bonds near linearity with all heteroatom distances less than 3.0 Å.

Conclusion

A modular approach to the synthesis of preorganized bis-(guanidinium) receptors that complement a phosphodiester has been demonstrated. The syntheses allow for facile manipulation of the cavity size and flexibility and will allow for the future incorporation of RNA recognition units, general bases, and metalbinding sites. Both 1:1 and 2:1 binding of phosphodiesters to these bis(alkylguanidinium) hosts in highly competitive hydrogenbonding solvents have also been demonstrated. In agreement with predictions from the active site of SNase¹⁴ and simple guanidinium-phosphate salts,15,16 both guanidiniums in the receptors form two hydrogen bonds to a single bound phosphoester, as supported by the crystal structure of a phosphomonoester, molecular mechanics calculations, and comparison to the binding constants of a simple monoguanidinium receptor. The tetraphenylborate salts of the bis(guanidinium) hosts show stronger binding for the meso forms than for the d, l forms due to the preorganization of the guanidinium groups on the same face of the relatively rigid spacers. For the chloride salts, d,l-3 binds more strongly than meso-3 due to a special ion effect through a bridging chloride

⁽³⁷⁾ Still, C. MACROMODEL version 2.5. Columbia University, Amber Force Field.

anion. In addition, the puckering modes within the cyclohexeno and cyclopenteno rings, along with possible bond rotations between the aminoimidazoline groups and the spacers, allow the receptors to open or close and thus accommodate either one or two anionic guests.

meso-3 has been shown to enhance the imidazole-catalyzed cleavage of mRNA at micromolar concentrations.¹⁰ The current postulate is that catalysis derives from increased binding in the transition-state-receptor complex due to the formation of an additional ion pair that is absent in the substrate-receptor complex. Our current efforts are focused on increasing the rate enhancement and deciphering the details of the mechanism behind the energetic stabilization that bis(guanidinium) receptors impart to phosphodiester hydrolysis.

Experimental Section

General Information. ¹H and ¹³C NMR spectra were obtained in CDCl₃, CD₃CN, DMSO- d_6 , or D₂O used as purchased without further purification. Spectra were recorded on a General Electric QE-300 (300 MHz) or a Bruker AC-250 (250 MHz) spectrometer. Melting points were measured on a Thomas Hoover capillary melting-point apparatus. Data were not corrected. Elemental analyses were performed by Galbraith Laboratories and Atlantic Laboratories. Low-resolution and high-resolution mass spectra were measured with Finnigan TSQ70 and VG Analytical ZAB2-E instruments, respectively.

All ³¹P NMR binding studies were performed on a Nicolet NT-360 (146 MHz) spectrometer, and all ¹H NMR binding studies were carried out with a General Electric QE-300 (300 MHz) or a Bruker AC-250 (250 MHz) spectrometer. An internal capillary tube was used in some ³¹P NMR binding studies, which contained triphenylphosphine as a reference ($\delta = -6.00$ ppm). Other studies were referenced to phosphoric acid in D₂O. Syringes and other glassware were dried under vacuum. Exact guest concentrations were determined by UV spectra (Beckman DU-70 UV-vis spectrophotometer).

Solvents and reagents were of reagent grade quality, purchased from Aldrich, and used without further purification except for DME. DME was purified by distillation from sodium-benzophenone ketyl. Preparative flash chromatography was performed on Scientific Adsorbants Incorporated Silica Gel 40 μ m. Analytical TLC was performed on precoated Silica Gel 60 F-254 plates.

Syntheses. Ethyl 2,2-Bis(2'-oxocyclohexyl)acetate (7). To a reaction mixture of 7.6 g (75 mmol) of ethyl glyoxylate in 25 mL of benzene was added 18.9 g (130 mmol) of 1-pyrrolidinylcyclohexene, and the reaction was warmed to reflux for 16 h. Then, 35 mL of water was added and the reaction allowed to reflux for 6 h. The solvent was evaporated, and the residue was extracted three times with 25 mL of ether. The extract was dried over MgSO4 and evaporated to dryness on a rotary evaporator. The residue was flashed through a silica gel column with 1:1 ethyl acetate/ hexane to give an orange oil which was ready to proceed to the next step. Crude yield: 9.4 g, 55%. Further purification can be achieved by crystallization from the same solvent system to give white crystals. Mp: 158.5-159.5 °C. ¹H NMR (300 MHz, CDCl₃): δ 4.13 (q, OCH₂CH₃, 2 H), 2.92 (dd, EtO₂CCH, 1 H), 2.63 (s, O-CCH, 1 H), 2.50 (t, O=CCH, 1 H), 2.21-1.30 (m, CH₂CH₂CH₂CH₂, 16 H), 1.26 (t, OCH₂CH₃, 3H). ¹³C [¹H] NMR (75 MHz, CDCl₃): δ 206.4, 172.9, 60.9, 60.1, 50.4, 48.9, 41.7, 37.0, 31.9, 29.9, 26.8, 26.2, 21.3, 20.1, 14.7. MS-CI: $m/z 281 (M^+ + H)$, 263. Anal. Calcd for C₁₆H₂₄O₄: C, 68.43; H, 8.63. Found: C, 67.73; H, 8.64.

1,2,3,4,5,6,7,8-Octahydro-9-acridinecarboxylic Acid Ethyl Ester (8). Compound 7 (6.5 g, 23.3 mmol) and 3.2 g (41.5 mmol) of ammonium acetate were refluxed with 20 mL of glacial acetic acid for 3 h. Water (200 mL) was added, and the mixture was extracted with CH₂Cl₂ (10 mL × 3). The combined dichloromethane layer was washed with 5% NaHCO₃ and evaporated under reduced pressure. A ¹H NMR spectrum showed the cyclized product which can be purified by silica gel chromatography with 1:2 ethyl acetate/hexane eluent. Yield: 3.6 g, 60%. Mp: 88.5-89.0 °C. ¹H NMR (300 MHz, CDCl₃): δ 4.32 (q, OCH₂CH₃, 2 H), 2.80 (t, CCCH₂, 4 H), 2.60 (t, NCCH₂, 4 H), 1.75 (m, CCH₂CH₂, 8 H), 1.30 (t, OCH₂CH₃, 3 H). ¹³C [¹H] NMR (75 MHz, CDCl₃): δ 168.0, 154.7, 141.5, 125.0, 61.2, 32.6, 25.7, 22.9, 22.6, 14.2. HRMS-CI: m/z calcd 259.1572 (C₁₆H₂₁NO₂), found 259.1559.

4,5-Dibenzylidene-1,2,3,4,5,6,7,8-octahydro-9-acridinecarboxylic Acid Ethyl Ester (9). Compound 8 (2.9 g, 11.3 mmol), 12 g (113 mmol) of benzaldehyde, and 12 mL of acetic anhydride were heated to reflux under N₂ for 10 h. The resulting solution was distilled under vacuum to remove the benzaldehyde and the acetic anhydride. The residue, after distillation, was purified by crystallization from ethyl acetate to give 4.02 g of product, 85% yield. An alternative method was to cool the residue in an ice bath to cause precipitation followed by collecting the precipitate by filtration and crystallization from hexane/EtOAc. Mp: 133.5–134.5 °C. ¹HNMR (CDCl₃, 300 MHz): δ 8.04 (s, CCH, 2 H), 7.36–7.10 (m, C₆H₅, 10 H), 4.33 (q, J = 7.2 Hz, OCH₂CH₃, 2 H), 2.80 (t, J = 5.4 Hz, CCCH₂, 4 H), 2.70 (t, J = 6.0 Hz, CHCCH₂, 4 H), 1.32 (t, J = 7.2 Hz, OCH₂CH₃, 3 H). ¹³C [¹H] NMR (75 MHz, CDCl₃): δ 168.2, 150.5, 141.6, 138.0, 135.4, 129.6, 128.1, 127.3, 126.8, 126.7, 61.3, 27.7, 22.5, 14.3. MS-EI: m/z 435 (M⁺), 362, 343. Anal. Calcd for C₃₀H₂₉NO₂: C, 82.76; H, 6.67; N, 3.22. Found: C, 82.13; H, 6.77: N, 3.24.

4,5-Dioxo-1,2,3,4,5,6,7,8-octahydro-9-acridinecarboxylic Acid Ethyl Ester (11). Compound 9 (9.8 g, 23 mmol) was dissolved in 300 mL of CH₂Cl₂ in a three-neck round-bottom flask equipped with a drying tube and a gas inlet. The solution was cooled to -78 °C, and ozone was bubbled through the solution until a blue color persisted. The excess ozone was purged by oxygen. After the blue color disappeared, 5 mL of Me_2S (67) mmol) was added and the mixture allowed to slowly warm to room temperature. The excess dimethyl sulfide and solvent were removed by vacuum. The residue was purified by a silica gel column $(1.5 \times 7 \text{ in.}^2)$ with pure EtOAc. The third fraction $(R_f = 0.15)$ was identified as the product. Yield: 3.2 g, 48%. Mp: 164-165 °C. ¹H NMR (300 MHz, CDCl₃): δ 4.46 (q, J = 7.2 Hz, OCH₂CH₃, 2 H), 2.97 (t, CCH₂, 4 H), 2.80(t, O=CCH₂, 4H), 2.15(m, CH₂CH₂CH₂, 4H), 1.42(t, OCH₂CH₃, 3 H). ¹³C [¹H] NMR (75 MHz, CDCl₃): δ 193.9, 165.8, 147.7, 142.6, 138.9, 62.2, 39.2, 26.8, 21.7, 14.1. MS-EI: m/z 287 (M⁺), 230, 122, 105. Anal. Calcd for C₁₆H₁₇NO₄: C, 66.9; H, 5.92; N, 4.88. Found: C, 65.46; H, 5.78; N, 4.65.

4,5-Diamino-1,2,3,4,5,6,7,8-octahydro-9-acridinecarboxylic Acid Ethyl Ester (5). A flame-dried 500-mL round-bottom flask was charged with 51.4 g (667 mmol, 21.5 equiv) of ammonium acetate (dried for 1 day under vacuum), 2.89 g (43.79 mmol, 1.4 equiv) of sodium cyanoborohydride, about 50 molecular sieves, 8.90 g (31.02 mmol) of diketone 11, and 250 mL of absolute ethanol. The inhomogeneous reaction mixture was stirred under nitrogen at room temperature for 93 h and then evaporated. The black oily residue was extracted with 100 mL of dichloromethane and 100 mL of ammonium hydroxide (pH \simeq 12). Sodium chloride was added (~ 1.0 M) to the aqueous layer, and the water was extracted with 100 mL of dichloromethane three times. All organic portions were combined, dried with magnesium sulfate, and evaporated. Preparative silica gel chromatography with 90:10 dichloromethane/ ammonia-saturated methanol yielded 7.32 g (81.7%) of a cream solid. ¹H NMR (300 MHz, CDCl₃): δ 4.4 (q, CH₂CH₃, 2 H), 3.95 (t, CH, 2 H), 2.7 (m, PhCH₂, 4 H), 1.7-2.3 (m, CH₂CH₂, NH₂, 8 H + 4 H), 1.4 (t, CH_2CH_3 , 3H). MS: m/z 290 (MW + 1). The full characterization was performed after isomer separation; see below.

4,5-Bis[(1,1-dimethylethoxy)carbonyl]amino]-1,2,3,4,5,6,7,8-octahydro-9-acridinecarboxylic Acid Ethyl Ester (14). A 250-mL round-bottom flask was charged with 7.32 g (25.35 mmol) of diamine 5, 17.23 g (76.61 mmol, 3.02 equiv) of di-tert-butyl dicarbonate, and 150 mL of chloroform. The solution was stirred at room temperature for 21 h. After evaporation, the residue was flashed through a silica gel column with 90:10 dichloromethane/ethyl acetate. Fractions at $R_f 0.55-0.40$ and $R_f 0.35 0.20\,were\,collected\,separately.\,\,Evaporation\,of\,both\,fractions\,gave\,colorless$ solids. Yield: 8.16 g (65.9%); d,l isomer, 4.33 g (1st fraction); meso isomer, 3.83 g (2nd fraction). d,lisomer: mp 48-54 °C. ¹H NMR (300 MHz, CDCl₃): δ 5.57 (br, NH, 2 H), 4.58 (m, CH, 2 H), 4.39 (q, CH₂CH₃, 2 H), 2.68 (m, PhCH₂, 4 H), 1.62, 1.89, 2.48 (br, CH₂CH₂, 8 H), 1.49 (s, tBu, 18 H), 1.25 (t, CH₂CH₃, 3 H). ¹³C [¹H] NMR (75 CDCl₃ MHz): δ 167.61, 156.16, 153.22, 141.89, 127.12, 79.22, 61.61, 51.87, 29.58, 28.47, 25.33, 19.64, 14.21. MS: m/z 490 (MW + 1). HRMS (CI + mode): m/z calcd 489.2838, found 489.2829. Anal. Calcd for C₂₆H₃₉N₃O₆: C, 63.78; H, 8.03; N, 8.58. Found: C, 63.46; H, 8.10; N, 8.67. meso isomer: mp 129-130 °C. 1H NMR (300 MHz, CDCl3: δ 5.53 (br, NH, 2 H), 4.61 (m, CH, 2 H), 4.40 (q, CH₂CH₃, 2 H), 2.70 (m, PhCH₂, 4 H), 1.5-2.5, 1.88 (br, CH₂CH₂, 8 H), 1.48 (s, tBu, 9 H), 1.38 (t, CH₂CH₃, 3 H). ¹³C [¹H] NMR (75 MHz, CDCl₃): δ 167.52, 155.96, 153.28, 124.02, 127.22, 79.30, 61.63, 51.63, 29.45, 25.31, 19.53, 14.21. MS-FAB: m/z 490 (MW + 1). HRMS (CI + mode); m/zcalcd 489.2838, found 489.2833. Anal. Calcd for C₂₆H₃₉N₃O₆: C, 63.78; H, 8.03; N, 8.58. Found: C, 62.33; H, 8.01; N, 8.40.

meso- and d,1-4,5-Diamino-1,2,3,4,5,6,7,8-octahydro-9-acridinecarboxylic Acid Ethyl Esters 5. A 250-mL round-bottom flask was charged

with 7.10 g (14.54 mmol) of 14 dissolved in 60 mL of trifluoroacetic acid and 20 mL of water. The solution was stirred at room temperature for 3.5 h and evaporated under vacuum. To the residue was added ammonium hydroxide solution (pH12 with 1 M NaCl), and the product was extracted with 100 mL of dichloromethane. The aqueous layer was washed with 100 mL of dichloromethane three times. All dichloromethane layers were combined, dried with magnesium sulfate, and evaporated. A white solid was obtained, which showed a single spot on the TLC plate with 90:10 dichloromethane/ammonia-saturated methanol, $R_f = 0.2$. Yield: 3.80 g (90.5%). d,l isomer: mp 93-94.5 °C. ¹H NMR (300 MHz, CDCl₃): δ 4.36 (m, CH₂CH₃, 2 H), 3.94 (m, CH, 2 H), 2.65 (m, PhCH₂, 4 H), 2.00 (s, NH₂, 4 H), 1.6–2.3 (m, CH₂CH₂, 8 H), 1.36 (t, CH₂CH₃, 3 H). ¹H NMR (500 MHz, CDCl₃): δ 4.32, 4.33, 4.34, 4.35, 4.36, 4.37, 4.38, 4.39. ¹³C [¹H] NMR (75 MHz, CDCl₃): δ 168.16, 157.46, 141.20, 125.27, 61.35, 51.58, 31.75, 26.16, 19.98, 14.25. MS-CI: m/z 290 (MW + 1). HRMS (CI + mode): m/z calcd for C₁₆H₂₄N₃O₂ (product + H⁺) 290.1868, found 290.1863. meso isomer: yield 99.8%; mp 73-75 °C. ¹H NMR (300 MHz, CDCl₃): δ 4.35 (q, CH₂CH₃, 2 H), 3.93 (m, CH, 2 H), 2.67 (m, PhC4H₂, 4 H), 2.19 (s, NH, 4 H), 1.6–2.1 (m, CH₂CH₂, 8 H), 1.36 (t. CH₂CH₃, 3 H), ¹H NMR (500 MHz, CDCl₃); δ 4.33, 4.34, 4.36, 4.37. ¹³C [¹H] NMR (75 MHz, CDCl₃): δ 168.06, 157.24, 141.30, 125.39, 61.37, 51.54, 31.61, 26.05, 19.87, 14.22, MS-CI: m/z 290 (MW + 1). HRMS (CI + mode): m/z calcd for C₁₆H₂₄N₃O₂ (product + H⁺) 290.1868, found 290.1871.

[[(1,1-Dimethylethoxy)carbonyl]amino]ethylIsothiocyanate (16). Ditert-butyl dicarbonate 17.98 g (79.91 mmol) was dissolved in 200 mL of chloroform and the solution added to a 30.0 mL (449 mmol, 5.62 equiv) solution of ethylenediamine dropwise for 2.5 h. The reaction mixture was stirred at room temperature for 19 h and then evaporated. The oily residue was extracted with 100 mL of Na₂CO₃ and 100 mL of dichloromethane. The aqueous layer was washed with 100 mL of dichloromethane. All dichloromethane layers were combined, dried with magnesium sulfate, and evaporated. The resulting oil was passed through a silica gel column with 90:10 dichloromethane/ammonia-saturated methanol eluent. The fraction at $R_10.3-0.2$ was collected. The product was a colorless oil. Yield: 10.72 g (83.9%). ^{1}H NMR (300 MHz, CDCl₃): δ 4.89 (br, NH, 1 H), 3.18 (q, CH₂CH₂NH, 2 H), 2.80 (t, CH₂CH₂NH, 2 H), 1.60 (br, NH, 2 H), 1.44 (s, tBu, 9 H). ¹³C [¹H] NMR (75 MHz, CDCl₃): δ 156.21, 79.26, 43.31, 41.83, 28.41. MS: m/z 161 (M + 1). A round-bottom flask was charged with 13.17 g (63.22 mmol, 1.0 equiv) of 1,3-dicyclohexylcarbodiimide, 26.0 mL (432 mmol, 6.82 equiv) of carbon disulfide, and 40 mL of THF and cooled to -7 to -10 °C with an NH₄Cl/ice (50 g/200 g) bath. The above oil (10.1 g, 63.2 mmol) was dissolved in 30 mL of dried THF and added dropwise over 30 min to the DCC solution at -7 to -10 °C. The reaction mixture was allowed to warm up to room temperature naturally and stirred at room temperature for 21 h. After evaporation, 400 mL of diethyl ether was added to the residue and a white insoluble solid (dicyclohexylthiourea) was removed by filtration. The filtrate was evaporated and passed through a silica gel column with 80:20 hexanes/ ethyl acetate. The fraction at $R_{f}0.4-0.3$ was collected. The product was a white fine solid. Mp: 63-64 °C. Yield: 11.89 g (93.2%). ¹H NMR (300 MHz, CDCl₃): δ 1.44 (s, tBu, 9 H), 3.37 (q, CH₂CH₂NH, 2 H), 3.63 (t, CH₂CH₂NH, 2 H), 4.90 (br, NH, 1 H). ¹³C [¹H] NMR (75 MHz, CDCl₃): § 155.54, 132.34, 80.13, 45.43, 40.57, 28.31. MS (CI +): m/z 203 (MW + 1). HRMS (EI + mode): m/z calcd for C8H14N2O2S 202.0776, found 202.0761.

meso- and d,I-4,5-Bis[[[[2-[[(1,1-dimethylethoxy)carbonyl]amino]ethyl]amino]thioxomethyl]amino]-1,2,3,4,5,6,7,8-octahydro-9-acridinecarboxylic Acid Ethyl Esters 17. A round-bottom flask was charged with 3.80 g (13.1 mmol) of 5, 6.7032 g (33.18 mmol, 2.52 equiv) of 16, and 250 mL of absolute ethanol. The reaction mixture was stirred at room temperature for 23 h during which time a precipitate formed. After evaporation, the residue was passed through a silica gel column eluted with 98:2 dichloromethane/methanol. The fraction at $R_f 0.1$ was collected. Yield: 7.23 g (78.7%). d,lisomer: mp 112.5-122.5 °C. ¹H NMR (300 MHz, DMSO-d₆): δ 7.80 (m, NH, 2 H), 7.41 (br, NH, 2 H), 6.83 (m, NH, 2 H), 5.17 (br, CH, 2 H), 4.36 (q, CH₂CH₃, 2 H), 3.42 (br, NHCH₂CH₂NH, 4H), 3.06 (m, NHCH₂CH₂NH, 4H), 2.62 (br, PhCH₂, 4 H), 1.74, 1.96 (br, CH₂CH₂, 8 H), 1.35 (s, tBu, 18 H), 1.29 (t, CH₂CH₃, 3 H). ¹³C [¹H] NMR (75 MHz, DMSO-*d*₆): δ 181.74, 166.91, 155.71, 153.35, 141.58, 127.25, 77.69, 61.48, 54.85, 53.50, 43.34, 28.49, 28.18, 25.09, 17.93, 13.96, MS-CI: m/z 694 (MW + 1). HRMS (CI + mode): m/z calcd for C₃₂H₅₂N₇O₆S₂ (M + H⁺) 694.3420, found 694.3410. Anal. Calcd for C₃₂H₅₁N₇O₆S₂: C, 55.39; H, 7.41; N, 14.13. Found: C, 54.81; H, 7.40; N, 14.00. meso isomer: yield 85.7%; mp 113-116.5 °C. ¹H

NMR (300 MHz, DMSO- d_6): δ 7.66 (br, NH, 4 H), 6.83 (m, NH, 2 H), 5.11 (br, NH, 2 H), 4.35 (q, CH₂CH₃, 2 H), 3.39 (br, NHCH₂CH₂NH, 4H), 3.08 (m, NHCH₂CH₂NH, 4H), 2.61 (m, PhCH₂, 4 H), 1.7–2.2 (br, CH₂CH₂, 8 H), 1.33 (s, tBu, 18 H), 1.32 (t, CH₂CH₃, 3 H). ¹³C [¹H] NMR (75 MHz, DMSO- d_6): δ 181.96, 166.81, 155.73, 153.25, 141.69, 127.05, 77.70, 61.51, 54.85, 53.61, 43.70, 28.71, 28.16, 24.86, 18.47, 13.96. MS (CI –): m/z 693 (MW). HRMS: m/z caled for C₃₂H₅₁N₇O₆S₂ 693.3342, found 693.3330. Anal. Calcd for C₃₂H₅₁N₇O₆S₂: C, 55.39; H, 7.41; N, 14.13. Found: C, 54.55; H, 7.39; N, 13.87.

meso- and d,I-N,N'-Bis(4,5-dihydro-1H-imidazol-2-yl)-1,2,3,4,5,6,7,8octahydro-9-acridinecarboxylic Acid Ethyl Esters 3. A round-bottom flask was charged with 7.01 g (10.12 mmol) of 17, 100 mL (1.34 mol, 132 equiv) of bromoethane, and 100 mL of absolute ethanol. The reaction mixture was allowed to reflux gently (oil bath, 50-60 °C) for 18 h. After reaction, the solution was evaporated and dried under vacuum. The product was an oily solid and used in the next step without purification. The oil (9.21 g, 10.12 mmol) was dissolved in 60.0 mL (779 mmol, 77.0 equiv) of trifluoroacetic acid and 20 mL of water, and the solution was stirred at room temperature for 2.5 h. The solvent was evaporated under vacuum to yield a brown solid which was used in the next step without further purification. Sodium (1.47 g, 64.08 mmol, 6.33 equiv) was dissolved in 500 mL of absolute ethanol. The brown solid (10.17 g, 10.12 mmol) was dissolved in 100 mL of absolute ethanol and added dropwise to the sodium ethoxide solution at room temperature over 30 min. The reaction mixture was stirred for an additional 20 min. After evaporation. the residue was passed through a silica gel column with 85:15 dichloromethane/methanol as eluent. The product was a slightly hygroscopic solid. Yield: 4.07 g (94.7%) from 9. d,l isomer. ¹H NMR (300 MHz, DMSO-d₆): δ 7.5-9.0 (br, NH, 4 H), 4.65 (m, CH, 2 H), 4.37 (q, CH2CH3, 2 H), 3.63 (s, NHCH2CH2NH, 8 H), 2.65 (m, PhCH2, 4 H), 1.7-2.2 (m, CH₂CH₂, 8 H), 1.30 (t, CH₂CH₃, 3 H). ¹³C [¹H] NMR (75 MHz, DMSO-d₆): δ166.48, 159.09, 151.92, 141.80, 127.75, 61.70, 52.68, 42.50, 28.76, 24.79, 18.18, 13.92, MS (CI-): m/z 425.5 (MW). HRMS (CI - mode) calcd for C₂₂H₃₁N₇O₂ 425.2539, found 425.2539. meso isomer: yield 85.3% (from 9). ¹H NMR (300 MHz, DMSO-d₆): δ 7.8-9.0 (br, NH, 4 H), 4.65 (m, CH, 2 H), 4.36 (q, CH₂CH₃, 2 H), 3.62 (s, NHCH2CH2NH, 8 H), 2.64 (m, PhCH2, 4 H), 1.7-2.2 (m, CH2CH2, 8 H), 1.29 (t, CH₂CH₃, 3 H). ¹³C [¹H] NMR (75 MHz, DMSO-d₆): δ 166.68, 159.65, 152.29, 142.06, 127.74, 61.91, 52.88, 42.76, 28.87, 24.93, 18.63, 14.13. HRMS (CI – mode): m/z calcd for C₂₂H₃₁N₇O₂ 425.2539, found 425.2513.

meso- and d,I-N,N'-Bis(4,5-dihydro-1H-imidazol-2-yl)-1,2,3,4,5,6,7,8octahydro-9-acridinecarboxylic Acid Ethyl Ester Dihydropicrates 3. Picric acid (0.27 g, 1.07 mmol, 4.1 equiv) was dissolved in 50 mL of hot water, and to this solution was added 0.1125 g (0.2647 mmol) of meso-3 or d,l-3 in 7 mL of water. After the solution was cooled to room temperature, the yellow precipitates were collected by filtration and washed with water. The precipitates were purified by crystallization from acetonitrile and ethyl acetate. d,l isomer: mp 218-219 °C. ¹H NMR (300 MHz, DMSO-d₆): δ 8.59 (d, NH, 2 H), 8.57 (s, PicH, 4 H), 7.4-9.0 (br, NH, 4H), 4.63 (m, CH, 2H), 4.36 (q, CH₂CH₃, 2H), 3.63 (s, NH(CH₂)₂NH, 8 H), 2.64 (m, PhCH₂, 4 H), 1.7-2.2 (br, CH₂CH₂, 8 H), 1.29 (t, CH₂CH₃, 3 H). ¹³C [¹H] (72.5 MHz, DMSO): δ 166.46, 160.77, 159.08, 151.97, 141.88, 141.79, 127.86, 125.21, 124.34, 61.71, 52.68, 42.57, 28.73, 24.82, 18.13, 13.91. Anal. Calcd for C₃₄H₃₇N₁₃O₁₆·H₂O: C, 45.28; H, 4.35; N, 20.18. Found: C, 45.53; H, 4.13; N, 20.38. meso-isomer: mp 247.5-248.5 °C dec. ¹H NMR (300 MHz, DMSO- d_6): δ 8.57 (s, NH, 4 H), 8.51 (d, PicH, 2 H), 7.5-9.0 (br, NH, 4 H), 4.62 (m, CH, 2 H), 4.36 (q, CH₂CH₃, 2 H), 3.62 (s, NH(CH₂)₂NH, 8 H), 2.64 (m, PhCH₂, 4 H), 1.7-2.2 (br, CH₂CH₂, 8 H), 1.29 (t, CH₂CH₃, 3 H). ¹³C [¹H] (72.5 MHz, DMSO- d_6): δ 166.50, 159.44, 152.09, 141.93, 141.84, 127.63, 125.20, 61.77, 56.04, 52.67, 42.63, 28.63, 24.76, 18.53, 18.42, 13.96. Anal. Calcd for C₃₄H₃₇N₁₃O₁₆·2H₂O: C, 45.28; H, 4.35; N, 20.18. Found: C, 45.81; H, 4.15; N, 20.36.

meso- and d, I-N, N-Bis(4,5-dihydro-1H-imidazol-2-yl)-1,2,3,4,5,6,7,8octahydro-9-acridinecarboxylic Acid Ethyl Ester Dihydrochlorides 3. The picrate crystals were dissolved in a minimum amount of THF/H₂O with 1-2 drops of concentrated aqueous HCl and then eluted on Amberlite IRA-400 (Cl⁻) resin. The resin was washed with water, and the effluent was collected fractionally. The fractions were spotted on a silica gel plate in the presence of iodine vapor. Iodine-active fractions were collected and lyophilized twice. A fine pale yellow solid was obtained. *d*, *isomer*: Yield 0.056 g (from free base) 42.8%. ¹H NMR (300 MHz, DMSO-d_6): δ 8.81 (d, NH, 2 H), 7.5-9.0 (br, NH, 4 H), 4.72 (t, CH, 2 H), 4.36 (q, CH₂CH₃, 2 H), 3.62 (s, NHCH₂CH₂NH, 8 H), 2.64 (m, PhCH₂, 4 H), 1.7–2.2 (m, CH₂CH₂, 8 H), 1.29 (t, CH₂CH₃, 3 H). ¹³C [¹H] NMR (75 MHz, DMSO- d_6): δ 166.49, 159.24, 152.07, 141.71, 127.54, 61.65, 52.69, 42.49, 28.82, 24.77, 18.36, 13.91. Anal. Calcd for C₂₂H₃N₇-O₂Cl₂·2H₂O: C, 49.44; H, 6.22; N, 18.34. Found: C, 50.62; H, 6.56; N, 18.23. *meso* isomer: yield 45.7% (from free base). ¹H NMR (300 MHz, DMSO- d_6): δ 8.78 (d, NH, 2 H), 7.6–9.0 (br, NH, 4 H), 4.85 (m, CH, 2 H), 4.35 (q, CH₂CH₃, 2 H), 3.63 (s, NHCH₂CH₂NH, 8 H), 2.64 (m, PhCH₂, 4 H), 1.7–2.3 (m, CH₂CH₂, 8 H), 1.29 (t, CH₂CH₃, 3 H). ¹³C [¹H] NMR (75 MHz, DMSO- d_6): δ 166.72, 159.87, 152.13, 142.07, 127.52, 53.27, 61.86, 42.70, 29.19, 24.77, 18.99, 14.13. Anal. Calcd for C₂₂H₃₃N₇O₂Cl₂·4H₂O): C, 46.30; H, 7.17; N, 16.99. Found: C, 46.31; H, 7.24; N, 17.19.

meso- and d, I-N, N'-Bis (4,5-dihydro-1H-imidazol-2-yl)-1,2,3,4,5,6,7,8octahydro-9-acridinecarboxylic Acid Ethyl Ester Dihydrotetraphenylborates 3. 3 dichloride (112 mg, 0.226 mmol) was dissolved in 20 mL of water, and an aqueous solution of sodium tetraphenylborate (170 mg, 0.497 mmol) was added. The solution was lyophilized and dissolved in acetonitrile. The insoluble NaCl was removed by filtration through Celite and the acetonitrile removed under vacuum. Yield: 249.8 mg, 88%. *meso* isomer: mp 117–118 °C. ¹H NMR (250 MHz, CD₃CN): δ 6.83, 6.99, 7.29 (tts, TPB, 40 H), 4.49 (t, CH, 2 H), 4.41 (q, CH₂CH₃, 2 H), 3.59 (s, NHCH2NH, 8 H), 2.73 (m, PhCH2, 4 H), 1.34 (t, CH2CH3, 3 H), 1.85, 2.17 (m, CH₂CH₂, 8 H). ¹³C NMR [¹H] (72.5 MHz, CD₃CN): δ 167.55, 164.5 (q), 161.01, 152.43, 136.64, 129.64, 126.55, 126.50, 122.68, 62.98, 54.44, 43.83, 29.44, 25.79, 19.39, 14.45. Anal. Calcd for C₇₀H₇₃N₇O₂B₂·3H₂O: C, 75.06; H, 7.10; N, 8.75. Found: C, 76.64; H, 6.94; N, 9.48. d,lisomer: mp 121.5-122.5 °C. ¹H NMR (250 MHz, CDCl₃): δ 6.87, 7.00, 7.26 (ttd, TPB, 40 H), 4.48 (m, CH, 2 H), 4.39 (q, CH₂CH₃, 2 H), 3.59 (s, NHCH₂CH₂NH, 8 H), 1.90, 2.08, 2.71 (m, CH₂CH₂, 8 H), 1.34 (t, CH₂CH₃, 3 H). ¹³C NMR [¹H] (72.5 MHz, CDCl₃): 8 170.18, 164.60 (q), 164.52, 160.50, 152.60, 144.17, 136.74, 126.66, 125.45, 62.90, 54.56, 43.93, 29.48, 25.87, 19.50, 14.45. Anal. Calcd for C₇₀H₇₃N₇O₂B₂·2H₂O: C, 76.29; H, 7.04; N, 8.89. Found: C, 76.64; H, 6.94; N, 9.48.

N-(4,5-Dihydro-1H-imidazol-2-yl)-cyclohexylamine Hydropicrate, Hydrochloride, and Hydrotetraphenylborate 19. 2-(Methylthio)-2-imidazoline hydroiodide (1.00 g, 4.10 mmol), 0.262 g (3.85 mmol) of sodium ethoxide, and 3.0 mL (26.23 mmol) of cyclohexylamine were combined in a round-bottom flask. The reaction mixture was heated at 90-100 °C, and equimolar amounts of 2-(methylthio)-2-imidazoline hydroiodide and sodium ethoxide along with a minimum amount of DMF were added portionwise over 27.5 h. The solvent was removed under vacuum, and the resulting orange oily solid was passed through a silica gel column with a 95:5 dichloromethane/ammonia-saturated methanol eluent. The fraction at $R_f 0.2-0.1$ was collected as the product. Further purification was performed by adding 6.70 g (26.37 mmol, 1.0 equiv) of picric acid in hot water. After the solution was cooled to room temperature, the yellow precipitate was collected by filtration. The yellow precipitate was crystallized from 100 mL of acetonitrile, which gave fine needle crystals. Yield: 4.22 g (40.6%). The monoguanidinium picrate was converted to the chloride by the same procedure as that of diguanidine compounds. Yield: 63.2%. Picrate 19: mp 220.5-221.5 °C. ¹H NMR (300 MHz, DMSO- d_6): δ 8.57 (s, PicH, 2 H), 8.11 (d, NH, 1 H), 7.3–8.5 (br, NH, 2 H), 3.55 (s, CH₂CH₂, 4 H), 3.24 (m, CH, 1 H), 1.19, 1.55, 1.65, 1.80 (m, (CH₂)₅, 10 H). ¹³C [¹H] NMR (75 MHz, DMSO-d₆): δ 160.79, 158.25, 147.80, 125.15, 124.21, 51.40, 42.34, 32.28, 24.63, 24.16. Anal. Calcd for C₁₅H₂₀N₆O₇: C, 45.45; H, 5.09; N, 21.20. Found: C, 45.45; H, 5.02; N, 21.38. Chloride 19: mp 186.5-187 °C. ¹H NMR (300 MHz, DMSO-d₆): δ 8.42 (d, PicH, 1 H), 7.3-8.8 (br, NH, 2 H), 3.54 (s, CH₂CH₂, 4 H), 3.33 (m, 1 H), 1.21, 1.50, 1.64, 1.80 (m, (CH₂)₅, 10 H). ¹³C [¹H] NMR (75 MHz, DMSO- d_6): δ 158.36, 51.18, 42.28, 32.25, 24.65, 23.97. Anal. Calcd for C9H18N3Cl: C, 53.06; H, 8.91; N, 20.62. Found: C, 52.08; H, 8.89; N, 20.05. Tetraphenylborate Salt 19: mp 151-152 °C. ¹H NMR (250 MHz, CDCl₃): δ 6.92, 7.05, 7.41 (tts, TPB, 20 H), 3.51 (d, NH, 1 H), 2.98 (s, CH, 2 H), 2.41 (s, CH₂CH₂, 4 H), 2.18 (m, CH, 1 H), 0.85, 1.10, 1.49 (m, (CH₂)₅, 10 H). Anal. Calcd for C33H38N3B: C, 81.24; H, 7.86; N, 8.62. Found: C, 80.27; H, 7.95; N. 8.66.

3,5-Dibenzylidene-1,2,3,5,6,7-hexahydro-8-phenyldicyclopenta[b,e]**pyridine (10).** Cyclopentanone (0.25 mol, 18.4 mL), benzaldehyde (0.5 mol, 50.8 mL), and ammonium acetate (0.25 mol, 19.3 g) were dissolved in 250 mL of absolute ethanol in an Erlenmeyer flask and warmed to reflux with stirring for 1 h. During this time, the solution changed from colorless to orange and a yellow-orange precipitate formed. After the solution was cooled on ice, the supernatant was decanted, leaving behind a pasty mass, which was washed with methylene chloride, leaving 8 g of

a light yellow powder (15%). Mp: 213–215 °C. ¹H NMR (250 MHz, CDCl₃): δ 7.9–7.0 (m, C₆H₅, CCH, 17 H), 3.0 (dt, CH₂CH₂, 8 H). ¹³C [¹H] NMR (72.5 MHz, CDCl₃): δ 160.88, 141.63, 137.97, 136.78, 129.05, 128.57, 128.39, 128.08, 126.67, 121.93, 29.37, 27.68. HRMS: m/z calcd for C₃₁H₂₅N 411.1987, found 411.1975.

3,5-Dioxo-1,2,3,5,6,7-hexahydro-8-phenyldicyclopenta[b,e]pyridine (12). Compound 10 (0.043 mol, 17.5 g) was dissolved in 1000 mL of methylene chloride in a 2000-mL three-necked round-bottom flask fitted with a drying tube, a gas inlet adapter, and a stopper. The solution was cooled in a dry ice/acetone bath (10 precipitates with cooling). Ozone was bubbled through the solution via the gas inlet for approximately 20 min. The flask was raised out of the cooling bath until the precipitated 10 was redissolved, about 10 min. The flask was returned to the dry ice/acetone bath for the remainder of the ozonolysis time, another 10-15 min. Completion of reaction was signaled by a color change from yellow to pale blue. The solution was purged with oxygen for about 10 min to remove excess ozone, and 3 equiv of dimethyl sulfide was added to the solution. The mixture was gradually warmed to room temperature and stirred for 12 h. The solvent was removed by rotary evaporation, leaving behind a dark brown oil. Purification was done on a silica gel flash column with 70:30 ethyl acetate/hexane as eluent ($R_f = 0.5$). Yield: 7.4 g, (66%). Mp: 215–217 °C. ¹H NMR (250 MHz, CDCl₃): δ7.6–7.35 (m, C₆H₅, 5 H); 3.12 (t, CH₂CH₂, 4 H), 2.83 (t, CH₂CH₂, 4 H). ¹³C [¹H] NMR (72.5 MHz, CDCl₃): δ203.96, 155.81, 150.46, 148.20, 133.42, 129.42, 129.26, 127.85, 35.67, 23.05. HRMS: calcd for C17H14NO2 264.1024, found 264.1019.

3,5-Bis(hydroxyimino)-1,2,3,5,6,7-hexahydro-8-phenyldicyclopenta-[b,e]pyridine (13). Compound 12 (0.019 mol, 5 g) was dissolved in 40 mL of dimethylformamide in a 250-mL round-bottom flask fitted with a reflux condenser. Five equivalents (0.095 mol, 10 g) of sodium carbonate and 4 equiv (0.072 mol, 5 g) of hydroxylamine hydrochloride were added with stirring. The mixture was heated on an oil bath to 80 °C for 3 h, during which time a cream-colored precipitate formed. The mixture was allowed to cool, and dimethyl formamide was removed under vacuum overnight. The precipitate was suspended in cold water, filtered, and washed with cold ethanol and ether. Yield: 4.4 g (79%). Mp: 285 °C dec. ¹H NMR (250 MHz, DMSO- d_6): δ 11.31 (s, NOH, 2 H), 7.48 (m, C₆H₅, 5 H), 2.81 (m, CH₂CH₂, 8 H). ¹³C [¹H] NMR (72.5 MHz, DMSO- d_6): δ 159.47, 155.75, 143.86, 140.46, 135.36, 128.65, 128.41, 128.13, 24.90, 24.78. HRMS: m/z calcd for C₁₇H₁₅N₃O₂ 293.1164, found 293.1161.

3,5-Diamino-1,2,3,5,6,7-hexahydro-8-phenyldicyclopenta[b,e]pyridine (6). Sodium borohydride (0.112 mol, 4.1 g, 8 equiv) and titanium(IV) tetrachloride (0.056, 6 mL, 4 equiv) were combined in a 1000-mL roundbottom flask in a drybox. The flask was fitted with a rubber septum, removed from the drybox, equipped with a nitrogen balloon, and cooled on an ice bath. Freshly distilled dimethoxyethane (100 mL) was added slowly while the flask was vented with a large bore needle. Compound 13 (0.014 mol, 4 g) (dried over phosphorus pentoxide under vacuum) was added as a powder over 5 min. The reaction mixture changed color over half an hour, from bright green to green-brown. The solution was allowed to warm to room temperature and stirred for 24 h. The mixture was cooled on ice, and water (200 mL) was added slowly to quench the reaction. The solution was basified with aqueous ammonia to pH 13, at which point it became gelatinous and turned dark blue. The aqueous layer was extracted several times with methylene chloride (total 1000 mL). The organic layer was dried over sodium sulfate, filtered, and evaporated to dryness on a rotary evaporator. Yield: 1.7 g (60%). ¹H NMR (250 MHz, CDCl₃): δ 7.4–7.2 (m, C₆H₅, 5 H), 4.35 (t, CH₂CH₂CH, 2 H), 3.0-2.0 (m, CH₂CH₂, 8 H). The compound was fully characterized after isomer separation.

meso- and d,I-3,5-Bis[[(1,1-dimethylethoxy)carbonyl]amino]-1,2,3,5,6,7hexahydro-8-phenyldicyclopenta[b,e]pyridines 15. Compound 6 (0.011 mol, 3 g) was dissolved in chloroform, and di-*tert*-butyl dicarbonate (0.033 mol, 7.2 g, 3 equiv) was added with stirring over a period of 15 min. The flask was fitted with a septum and a nitrogen balloon and allowed to stir for another 45 min. The solvent was removed on a rotary evaporator. TLC of the reaction mixture in 70:30 hexane/ethyl acetate gave R_f values for the d,l and meso isomers of 0.38 and 0.23, respectively. Separation of the d,l and meso isomers was achieved on a flash column with 70:30 hexane/ethyl acetate eluent. The appearance of the d,l isomers was observed as a white precipitate forming on the tip of the column. The flow was then reduced to gravity flow until the first fraction was finished and then the flow rate was increased again. Yield of each isomer: 50%, 1 g each. meso lsomer: mp 162–165 °C. ¹H NMR (250 MHz, CDCl₃): δ 7.5–7.3 (m, C₆H₅, 5 H), 5.2 (s, NH, 2 H), 4.95 (q, CH₂CHNH, 2 H), 3.5-3.0 (m, CH2C*H*H, 6 H), 1.95 (m, CH*H*, 2 H), 1.48 (s, COC(CH₃)₃, 18 H). ¹³C [¹H] NMR (72.5 MHz, CDCl₃): δ 161.99, 155.95, 143.74, 136.49, 134.06, 128.21, 79.45, 56.31, 33.82, 28.44, 26.99. HRMS: *m/z* calcd for C₂₇H₃₅N₃O₄ 465.2627, found 465.2634. *d,l* isomer: mp 179– 182 °C. All the ¹H NMR resonances are the same for the *d,l* isomer as for the *meso* except the peak at 5.2 is shifted to 5.35. ¹³C [¹H] NMR (72.5 MHz, CDCl₃): δ 161.99, 156.09, 143.25, 136.27, 133.75, 128.37, 79.38, 56.47, 34.55, 28.31, 27.31. HRMS: *m/z* calcd for C₂₇H₃₅N₃O₄ 465.2627, found 465.2632.

meso- and d,1-3,5-Diamino-1,2,3,5,6,7-hexahydro-8-phenyldicyclopenta[b,e]pyridines 6. Compound 15 (0.0064 mol, 3 g) was dissolved in a mixture of 10 mL of trifluoroacetic acid and 10 mL of H₂O. The solution was stirred for 4 h and the solvent removed under vacuum. The resulting paste was dissolved in 200 mL of 1 M sodium chloride in aqueous ammonia (pH 13). The amine was extracted from the aqueous layer with four 75-mL fractions of dichloromethane. After the solution was dried over sodium sulfate, the dichloromethane was removed by rotary evaporation. Yield: 1.15 g (68%). meso isomer: mp 134-139 °C. 1H NMR (250 MHz, CDCl₃): δ 7.43-7.26 (m, C₆H₅, 5 H), 4.37 (t, CH, 2 H), 2.85-2.49 (m, PhCH₂CH₂, NH₂, 10 H), 1.76 (m, CH₂, 2 H). ¹³C [¹H] NMR (72.5 MHz, CDCL₃): δ 165.82, 143.37, 137.05, 132.72, 128.85, 128.42, 128.24, 127.95, 53.37, 32.41, 26.05. HRMS: m/z calcd for C17H19N3 265.1579, found 265.1569. d,l isomer: mp 163-165 °C. ¹H NMR (CDCl₃): δ 7.4–7.2 (m, C₆H₅, 5 H), 4.37 (t, CH, 2 H), 2.79– 2.60 (m, PhCH₂CH₂, NH₂, 10 H), 1.68 (m, CH₂, 2 H). ¹³C [¹H] NMR (72.5 MHz, CDCl₃) δ 166.14, 143.02, 136.98, 132.45, 128.34, 127.94, 56.74, 34.91, 27.06. HRMS: m/z calcd for C17H19N3 265.1579, found 265.1570.

meso- and d,l-3,5-Bis[[[[2-[[(1,1-dimethylethoxy)carbonyl]amino]ethyl]amino]thioxomethyl]amino]-1,2,3,5,6,7-hexahydro-8-phenyldicyclopenta[b,e]pyridines 18. Compound 16 (0.005 mol, 1.05 g, 2.5 equiv) in 50 mL of absolute ethanol was added dropwise to compound 6 (0.002 mol, 550 mg) dissolved in 25 mL of absolute ethanol. The solution was stirred for 20 h under a positive pressure of nitrogen. TLC in 98:2 methylene chloride/methanol (developed with iodine) revealed the product at an R_f of 0.25. Flash silica gel chromatography in the same eluent gave 2.3 g (67%) for the d,l isomer and 3.4 g (quantitative) for the meso isomer. meso isomer: mp 73-75 °C. ¹H NMR (250 MHz, CDCl₃): δ 7.7-7.3 (m, C₆H₅, 5 H), 5.3 (br t, CH₂CH₂NH, 2 H), 3.7 (br, NHCH₂, 4 H), 3.3 (br, NHCH₂, 4 H), 2.75 (br, CH₂CH₂, 4 H), 1.95 (br, CH₂CH₂, 4 H), 1.3 (s, C(CH₃)₃, 18 H). ¹³C [¹H] NMR (72.5 MHz, CDCl₃): δ 180.84, 161.36, 150.09, 144.43, 135.82, 134.75, 128.44, 128.33, 128.00, 83.13, 79.38, 59.30, 52.66, 47.57, 45.22, 40.89, 40.43, 32.66, 28.16, 27.88, 27.14, 25.24, 24.58. HRMS: m/z calcd for C₃₃H₄₇N₇O₄S₂ 669.3131, found 699.3130. d,l isomer: mp 104-112 °C. ¹H NMR (250 MHz, CDCl₃): δ 7.47, 7.24 (m, C₆H₅, 5 H), 5.65 (br s, NH, 2 H), 3.84 (br s, NH, 2 H), 3.4 (br s, NH, 2 H), 2.81 (m, CH₂CH₂), 2.2-1.7 (m, CH₂, 4 H), 1.26 (s, tBu, 18 H). ¹³C [¹H] NMR (72.5 MHz, CDCl₃): δ183.12, 162.28, 157.19, 144.67, 135.68, 134.68, 128.72, 128.06, 79.68, 77.21, 59.31, 52.84, 40.86, 40.61, 33.61, 32.84, 28.31, 28.24, 28.06, 27.23, 25.40, 24.67. HRMS: m/z calcd for C33H47N7O4S2699.3131, found 699.3120.

meso- and d,I-N,N'-Bis(4,5-dihydro-1H-imidazol-2-yl)-1,2,3,5,6,7hexahydro-8-phenyldicyclopenta[b,e]pyridine-3,5-diamines 4. Compound meso-18 (0.0015 mol, 0.9 g) and compound d,l-18 (0.0025 mol, 1.5 g) were each dissolved in a solution of 10 mL of bromoethane and 10 mL of absolute ethanol. Each flask was fitted with a reflux condenser and a nitrogen ballon, and the mixtures were refluxed for 20 h. A TLC in 90:10 methylene chloride/methanol confirmed formation of the product, with an R_f of 0.72 for the meso isomer and 0.61 for the d,l isomer. The solvent was removed by rotary evaporation. This product was not isolated. The resulting oils were each dissolved in about 6 mL of trifluoroacetic acid and 3 mL of H₂O and stirred at room temperature for 3-4 h. TLC in 90:10 methylene chloride/methanol saturated with NH₃ showed a product with an R_f of 0.37 for the d, l isomer and 0.44 for the meso isomer. The TFA was removed under vacuum overnight and the product lyophilized. This product was not isolated but taken directly to the next step. Approximately 100 mg of sodium was dissolved in 10 mL of absolute ethanol. The resulting solid from the previous step was dissolved in 20 mL of absolute ethanol and added dropwise to the sodium ethoxide solution over 2 h. The system was sealed with a balloon and stirred for 24-36 h, and the solvent was removed by rotary evaporation. meso isomer: yield 522 mg (90%). ¹H NMR (250 MHz, CDCl₃): 88.89 (br s, NH, 2 H), 8.61 (d, NH, 2 H), 7.96 (br s, NH, 2 H), 7.42, 7.25 (m, C₆H₅, 5 H), 5.11 (q, CH₂CHNH, 2 H), 3.67 (s, NHCH₂CH₂NH, 8 H), 2.67, 1.88 (m, CCH₂CH₂CH, 8 H). ¹³C [¹H] NMR (72.5 MHz, CDCl₃): δ 162.72, 162.18, 161.63, 161.88, 160.78, 160.37, 160.22, 144.52, 135.59,

135.75, 128.64, 128.03, 57.72, 45.19, 32.73, 27.08. *d*,*l*isomer: yield 600 mg (62%). ¹H NMR (250 MHz, CDCl₃): δ 9.43 (d, NH, 2 H), 9.18 (br s, NH, 2 H), 7.96 (br s, NH, 2 H), 7.43, 7.28 (m, 2 H, 5 H), 4.82 (q, CH₂CH₁NH, 2 H), 3.66, (s, NHCH₂CH₂NH, 8 H), 2.9–2.7, 1.92 (m, CCH₂CH₂CH, 8 H). ¹³C [¹H] NMR (72.5 MHz, CDCl₃) 163.52, 162.96, 162.41, 161.85, 160.87, 160.83, 144.32, 135.99, 134.14, 128.81, 128.72, 128.14, 118.93, 114.27, 57.66, 43.05, 32.17, 27.36.

meso- and d,I-N,N'-Bis(4,5-dihydro-1H-imidazol-2-yl)-1,2,3,5,6,7hexahydro-8-phenyldicyclopenta[b,e]pyridine-3,5-diamine Dihydropicrates 4. Compounds meso-4 and d,l-4 were purified by conversion to picrate salts in the following manner. Each compound (3.62 mmol, 1.4 g) was dissolved in 80 mL of water. Picric acid (8 mmol, 1.82 g, 2.2 equiv) was dissolved in 100 mL of water. The solutions were heated to boiling and then combined, at which time a precipitate formed. The mixture was allowed to stand overnight and then filtered and washed with cold water, ethanol, and ether. The same procedure was used for the d,l compound except it was found that the solutions should not be heated before mixing as this induces an oil to form. meso-isomer: mp 137-139 °C. ¹H NMR (250 MHz, DMSO-d₆): δ 8.69 (d, NH, 2 H), 8.58 (s, C₆H₂O(NO₂)₃, 4 H), 7.45 (m, C₆H₅, 5 H), 4.99 (q, CH₂CHNH, 2 H), 3.67 (s, NHCH₂CH₂NH, 8 H), 2.80, 1.98 (m, CCH₂CH₂CH, 8 H). ¹³C [¹H] NMR (62.5 MHz, DMSO-d₆): δ160.87, 159.40, 143.48, 141.81, 135.45, 134.30, 128.70, 128.22, 125.17, 124.29, 57.35, 42.61, 31.31, 26.74. Anal. Calcd for C₃₅H₃₃N₁₃O₁₄·4H₂O: C, 45.12; H, 4.43; N, 19.53. Found: C, 45.13; H, 3.56; N, 19.70. *d*,*l* isomer: mp 179–181 °C. ¹H NMR (250 MHz, D₂O): δ 8.69 (d, NH, 2 H), 8.58 (s, C₆H₂O(NO₂)₃, 4 H), 7.49 (m, C₆H₅, 5 H), 5.03 (q, CH₂CHNH, 2 H), 3.60 (s, NHCH₂CH₂NH, 8 H), 2.85, 2.60, 1.85 (m, CH₂CH₂, 8 H). ¹³C [¹H] NMR (72.5 MHz, D_2O): δ 160.71, 159.21, 143.21, 141.81, 135.37, 134.02, 128.66, 128.33, 125.15, 124.23, 57.47, 42.59, 32.00, 26.67. Anal. Calcd for C35H33N13O14.4H2O: C, 45.12; H, 4.43; N, 19.53. Found: C, 45.10; H, 3.53; N, 19.84.

meso- and d, I-N, N'-Bis(4,5-dihydro-1H-imidazol-2-yl)-1,2,3,5,6,7hexahydro-8-phenyldicyclopenta[b,e]pyridine-3,5-diamine Dihydrochlorides 4. IRA-100 ion-exchange resin in the OH⁻ form was equilibrated in 0.2 M HCl overnight. Approximately 4 mL of resin was packed in a 1-cm-diameter column and washed to neutrality with water. The dipicrate of 4 (300 mg) was dissolved in 300 mL of 50% acetonitrile in water. The solution was eluted slowly, and the effluent was collected in a 1-L round-bottom flask, evaporated to dryness, redissolved in a minimum amount of water, transferred to a smaller flask, and lyophilized. Yield: 135.1 mg (83.3%). meso isomer: mp 186-187 °C. ¹H NMR (250 MHz, D_2O): δ 7.45 (m, C₆H₅, 5 H), 4.9 (t, CH₂CHNH, 2 H), 3.62 (s, NHCH₂CH₂NH, 8 H), 2.75, 2.5, 1.9 (m, CH₂CH₂, 8 H). ¹³C [¹H] NMR (72.5 MHz, D₂O): δ 160.50, 159.06, 148.76, 138.05, 136.06, 130.04, 129.60, 129.15, 58.14, 43.58, 32.49, 27.56. Anal. Calcd for C23H29N7Cl2O: C, 50.55; H, 6.81; N, 17.95. Found: C. 50.45; H, 6.20; N, 16.84. *d*,*l* isomer: mp 146–147 °C. ¹H NMR (250 MHz, D₂O): δ 7.33 (m, C₆H₅, 5 H), 4.97 (t, CH₂CHNH, 2 H), 3.59 (s, NHCH₂CH₂NH, 8 H), 2.74, 2.50, 1.83 (m, CH₂CH₂, 8 H). ¹³C [¹H] NMR (72.5 MHz, D_2O): δ 160.89, 160.68, 145.77, 136.67, 136.43, 129.67, 129.60, 129.31, 119.99, 58.68, 45.94, 43.64, 32.55, 27.42. Anal. Calcd for C23H29N7Cl2O·3H2O: C, 52.28; H. 6.66; N, 18.56. Found: C, 52.63; H, 6.33; N, 18.00.

meso- and d,I-N,N'-Bis(4,5-dihydro-1H-imidazol-2-yl)-1,2,3,5,6,7hexahydro-8-phenyldicyclopenta[b,e]pyridine-3,5-diamine Dihydrotetraphenylborates 4. Compound 4 dichloride (0.285 mmol, 135 mg) was dissolved in water and then combined with sodium tetraphenylboron (0.57 mmol, 222 mg, 2 equiv) also dissolved in water. A white precipitate formed immediately. The solution was stirred overnight and then lyophilized. A white powder was formed, which was dissolved in acetonitrile. Any additional sodium chloride was left undissolved and filtered off through Celite. meso isomer: mp 108-109 °C. ¹H NMR (250 MHz, D₂O): § 7.43 (m, C₆H₅, 5 H), 7.30, 6.99, 6.83 (stt, TPB, 40 H), 4.85 (t, CH₂CHNH, 2 H), 3.62 (s, NHCH₂CH₂NH, 8 H), 2.86, 2.60, 1.96 (m, CH₂CH₂, 8 H). ¹³C [¹H] NMR (72.5 MHz, D₂O): δ 164.65 (q), 161.18, 146.0, 136.6, 135.47, 129.77, 129.15, 125.57, 126.50, 122.70, 125.50. Anal. Calcd for C71H69N7OB: C, 79.10; H, 6.82; N, 9.10. Found: C, 79.72; H, 6.72; N, 9.31. d,l isomer: mp 220-225 °C. ¹H NMR (250 MHz, CD₃CN): δ 7.63 (d, NH, 2 H), 7.42 (m, C₆H₅, 5 H), 7.28, 7.01, 6.82 (stt, TPB, 40 H), 4.88 (t, CH₂CHNH, 2 H), 3.65 (s, NHCH₂CH₂NH, 8 H), 2.75, 2.61, 1.9 (m, CH₂CH₂, 8 H). ¹³C [¹H] NMR (72.5 MHz, CD₃CN): δ 165.17, 160.5, 146.00, 135.47, 129.71, 129.15, 126.56, 126.50, 122.76, 122.70, 125.5.

Determination of Binding Constants. All solutions were prepared in a drybox. A solution of sodium dibenzyl phosphate, henceforth called

the stock solution, of about 10 mM concentration was made, and the concentration was determined by UV-vis absorbance. The extinction coefficients at 255, 256, and 260 nm were 404, 437, and 377 M⁻¹ cm⁻¹, respectively (determined by Beer's law plots). Approximately 80 mM host solution ($550 \ \mu$ L) was made using the phosphate stock solution, and exactly 500 $\ \mu$ L was transferred to a 5-mm NMR tube sealed with a septum. The dibenzyl phosphate resonance was observed as a single peak in the ³¹P NMR. The chemical shift of pure dibenzyl phosphate in DMSO is δ 0.52 as referenced to phosphoric acid in D₂O. Aliquots of the host-guest solution were removed from the NMR tube via syringe through the septum and replaced by an equal volume of the guest stock solution. In this manner, the host concentration was systematically reduced while the guest concentration remained constant. The ratio of host to guest was determined by ¹H NMR integration.

The ¹H NMR binding studies followed the chemical shift of the amine protons of the host compounds when the solvent was pure DMSO. In DMSO/water mixtures, the methylenes of the imidazoline groups were monitored. A 5-mm NMR tube was fitted with a septum and charged with 500 μ L of an approximately 3 mM solution of host, and small aliquots (1-5 μ L) of a guest stock solution of approximately 100 mM were added. The binding constants were obtained by using a curve-fitting program provided by Dr. Howard Whitlock of The University of Wisconsin.

X-ray Crystallography. Crystals of (C23H29N7)2+(C6H5PO4)2-H2O [meso-4(C₆H₅PO₄)²- H_2O] were grown from H₂O-DMF solution as long, colorless crystals, while crystals of (C23H29N7)2+(C14H14OPO3)-H2O [meso-4($C_{14}H_{14}OPO_3$)⁻(Cl)⁻· H_2O] were grown from H_2O -DMSO solution, also as long, colorless needles. Data for both were collected on a Nicolet P3 diffractometer with a graphite monochromator, using Mo $K\alpha$ radiation ($\lambda = 0.71073$ Å). Details of the crystal data, data collection, and structure refinement are listed in the supplementary material, Table 1. The data for meso-4(C₆H₅PO₄)²- \cdot H₂O were collected from 4.0° to 47.5° in 2 θ using the ω scan technique, with a 1.2° scan range in ω at 2.5-5° min⁻¹. A total of 6680 reflections were measured, of which 4565 were unique (R for averaging equivalent reflections = 0.019). It was apparent from the initial photographs of the data crystal that it was limited in scattering power, and therefore, the data were collected in two stages. For the first stage, a quadrant of the reciprocal lattice was collected out to a 2θ maximum of 47.5° where h ranged from 0 to 12, k ranged from 0 to 10, and I ranged from -36 to 36. A second stage was collected out to a 2θ maximum of 30° to give some redundancy to that part of the data set most likely to be observable. Three reflections (1, 0, 8; 2, 1, -3;and -2, -1, -3) were remeasured every 97 reflections to monitor instrument and crystal stability. A smoothed curve of the intensities of these check reflections was used to scale the data. The scaling factor ranged from 0.982 to 1.01. The data for meso-4($C_{14}H_{14}OPO_{3}$)-(Cl)- $H_{2}O$ were collected from 4.0° to 50° in 2θ using ω scan technique, with a 1.2° scan range in ω at 3-6° min⁻¹. Two complete equivalent sets of data were collected. A total of 10 516 reflections were measured, of which 6545 were unique (R for averaging equivalent reflections = 0.026) where h ranged from -21 to 20, k ranged from 0 to 12, and l ranged from -23 to 23. Four reflections (-2, 2, 4; 3, 0, 4; 1, -3, -1; and -1, 2, -3) were remeasured every 96 reflections to monitor instrument and crystal stability. A smoothed curve of the intensities of these check reflections were used to scale the data. The scaling factor ranged from 0.991 to 1.01. The data for both samples were corrected for L_p effects and absorption. The absorption correction was based on measured crystal faces. Data reduction and decay correction were performed using the SHELXTL-Plus software package.³⁸ The structures were solved by direct methods and refined by full-matrix least squares³⁸ with anisotropic thermal parameters for the non-H atoms. For meso-4($C_6H_5PO_4$)^{2-,} H_2O , the hydrogen atoms were located from a ΔF map and refined with isotropic thermal parameters. For meso-4(C14H14OPO3)-(Cl)-H2O, hydrogen atoms, except for those on the water molecules, were calculated in ideal positions (C-H, 0.96 Å and N-H, 0.90 Å) with U_{iso} fixed at $1.2U_{eq}$ of the relevant atom. Positions for the hydrogen atoms on the water molecules were located from a ΔF map and were tied to the water oxygen atom during least squares. The function $\sum w(|F_0| - |F_c|)^2$ was minimized, where $w = 1/(\sigma(F_0))^2$ and $\sigma(F_0)$ = $0.5kI^{-1/2}[(\sigma(I))^2 + (0.02I)^2]^{1/2}$. The intensity I is given by $(I_{\text{peak}} I_{\text{background}}$ (scan rate), where 0.02 is a factor to down-weight intense reflections and to account for instrument instability and k is the correction due to Lp effects, absorption, and decay. $\sigma(I)$ was estimated from counting statistics; $\sigma(I) = (I_{\text{peak}} + I_{\text{background}})^{1/2}$ (scan rate). Neutral atom scattering factors for the non-H atoms were taken from Cromer and Mann,³⁹ with the anomalous-dispersion corrections taken from the work of Cromer and Liberman.⁴⁰ The scattering factors for the H atoms were obtained from Stewart, Davidson, and Simpson.⁴¹ Values used to calculate the linear absorption coefficients are from International Tables for X-ray Crystallography (1974).42 All figures were generated using SHELXTL-Plus.¹ Other computer programs used in this work are listed elsewhere.⁴³

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Supplementary Material Available: X-ray experimental summary, positional and thermal parameters, bond lengths and angles, listing of the H-bonding interactions, view showing the atomlabeling schemes, and unit cell packing diagrams along with some additional superposition plots of the tricyclic ring system of *meso*- $4(C_6H_5PO_4)^2$ ·H₂O fitted by *meso*- $4(C_{14}H_{14}OPO_3)^-(Cl)^-$ ·H₂O (48 pages); tables of observed and calculated structure factor amplitudes (41 pages). Ordering information is given on any current masthead page.

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