X-ray structure of 3d. We also thank Central Glass and Tosoh Akzo Co. for gifts of hexafluorocumyl alcohol and alkyllithiums, respectively.

Supplementary Material Available: Physical and spectral data of 3a-e and 6 and X-ray crystallographic data with tables of thermal and positional parameters, bond lengths, and bond angles for 3d (18 pages). Ordering information is given on any current masthead page.

New Triply Hydrogen Bonded Complexes with Highly Variable Stabilities

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The use of hydrogen bonds to confer binding strength and selectivity has become a dominant theme in host-guest complexation studies.¹ As the number of reports on hydrogen-bonded complexes grows, so will the opportunities to discern patterns and, in turn, formulate rules for predicting the properties of unknown systems.² A case in point is the insightful analysis of triply hydrogen bonded systems recently reported by Jorgensen.³ It notes that two complexes in which hydrogen bond donor (D) and acceptor (A) groups alternate (ADA DAD; 1.2, 3.4) have $K_{\text{assoc}} \approx 10^2 \text{ M}^{-1}$,^{4,5} while two DDA AAD complexes (5.6, 7.8) are significantly stronger with $K_{\rm assoc} \approx 10^4 \, {\rm M}^{-1.6.7}$ Since the primary hydrogen bonds were similar in each system, the discrepancy was proposed to result from the different arrangement of the hydrogen-bonding sites and, in turn, different secondary electrostatic interactions. To test the generality of this analysis, we have experimentally examined four new triply hydrogen bonded complexes (9.10, 9.11, 12.13, 14.15), of which one included the pre-

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viously unknown DDD-AAA hydrogen-bonding motif (12.13). This latter arrangement contained four attractive secondary interactions and was predicted computationally to lead to the strongest complex.³



Most of the compounds used in this study were commercially available or were readily prepared using known procedures.⁸⁻¹¹

⁽⁸⁾ Compounds 10 and 13 were prepared according to ref 10. Compounds 9 and 12 were prepared according to ref 9. Compound 11 was prepared according to ref 11. Compound 14 is available from the Aldrich Chemical Co. as the amine. All compounds used in this study gave correct elemental analyses and had spectral data in full accord with the assigned structures.

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Table I. Association Constants and Complexation Shifts for Various Triply Hydrogen Bonded Complexes in Chloroform- d^a

complex	type	method used	proton(s) monitored	$\Delta \delta_{max}$ (ppm)	K_{assoc} (M ⁻¹)	$-\Delta G^{\circ}_{298}$ (kcal mol ⁻¹)	ref
1.2	DADADA				90	2.7	5
3.4	DADIADA				140	2.9	6
9.10	DAD·ADA	Α	N9-H (9)	3.98	78	2.6	this work
9.11	DAD·ADA	Α	N9-H (9)	4.10	70	2.5	this work
7.8	DDA·AAD				1.7×10^{4}	5.8	7
16-17 (5-6)	DDA ·AAD	В			1×10^{4}	5.5	this work
14.15	DDA·AAD	В	N1-H (15)	1.63	9.3×10^{3}	5.4	this work
12·13 ^b	DDD-AAA	В	NH ₂ (13)	1.80	≥10 ⁵	≥7	this work

"At 298 K. Duplicate runs gave Kassoc values that agreed within 5%. Method A: Titration with 9 at fixed concentration. Method B: Dilution of 1:1 complex. ^bCarried out in the presence of 2 molar equiv of 1,8-bis(dimethylamino)naphthalene (see ref 18).

Compound 15 was prepared by alkylating 9-deazaguanosine¹² with dioctyl 4-(bromomethyl)phthalate,¹³ the long-chain ester groups providing high solubility in chloroform. Only an approximate association constant was reported⁶ for the GC base pair (5.6) in chloroform, so a more accurate determination was made using 2',3',5'-tripentanoylguanosine (16)¹⁴ and 4-ethylcytosine (17).¹¹

The 4-aryl group in dihydropyridines 10 and 13 serves as a convenient tautomeric "switch". Thus, in dimethylformamide- d_7 , 13 exists exclusively in the 1,4-dihydro form, while 10 is entirely in the 3,4-dihydro form, presumably due to a steric effect.¹⁰ In chloroform-d, the ¹H NMR spectrum of 10 showed it to be in the 3,4-dihydro form, while that for 13 indicated a solvent-induced shift in the equilibrium to a ca. 67:33 mixture of 1,4-dihydro and 3,4-dihydro forms.¹⁶ Interestingly, 10 equiv of 12 converted 10 (ca. 1 mM, CDCl₃) entirely into the 1,4-dihydro form. Likewise, 4 equiv of 9 converted 13 (ca. 5 mM, CDCl₃) from a 67:33 mixture to a 44:56 mixture of 1,4-dihydro and 3,4-dihydro forms.

Complexation studies were performed by ¹H NMR spectroscopy in chloroform-d under conditions where self-association of 9-17was negligible. The association constants were determined using standard methods;17 they are compiled in Table I. The alternating hydrogen-bonding motif (DAD ADA) in 9.10 and 9.11 was weak with K_{assoc} values of 78 and 70 M⁻¹, respectively. The DDA AAD complexes of 1415 and the GC base pair (1617) were significantly more stable with $K_{\rm assoc}$ values of 9.3 \times 10³ and 1 \times 10⁴ M⁻¹ respectively. The 12.13 (DDD AAA) complex was by far the tightest examined. Indeed, only ca. 15% of uncomplexed 13 was observed when the concentration of the 1:1 complex was ca. 2 \times 10⁻⁴ M. By accounting for the tautomeric equilibrium constant, the association constant can be estimated to be larger than or equal to $10^5 \text{ M}^{-1.18}$ The difference in stability between complexes 9.10 and 12.13 is striking. Despite their structural similarity and identical number of hydrogen bonds, the latter complex is more stable by over 4 kcal mol⁻¹.

These results are consistent with Jorgensen's proposal that the variable stabilities of triply hydrogen bonded complexes originate in the arrangement of the hydrogen-bonding sites.^{3,19} Combined with previous results,³⁻⁷ the variations in K_{assoc} seen in Table I, and in particular the large difference in stability of complexes 9.10 and 12.13, indicate that the number of hydrogen bonds alone will

be a poor predictor of complex stability. For triply hydrogen bonded complexes the arrangement of donor and acceptor groups appears to correlate well with K_{assoc} , although not all triply hydrogen bonded complex stabilities are expected to fall in the narrow ranges found for the complexes in Table I.¹⁹⁻²¹ Efforts are underway to incorporate some of these new complexes into supramolecular assemblies.

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Molecular Harpoons: Membrane-Disrupting Surfactants That Recognize Osmotic Stress¹

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In this paper we introduce a class of membrane-disrupting surfactants that has been designed to recognize osmotic stress in lipid bilayers. On the basis of their molecular shape and their pressure-sensitivity, we term such amphiphiles molecular harpoons. Results presented herein show that the ability of a harpoon to recognize a stressed bilayer is a function of its structure, its oligomeric state, and the strength of the osmotic gradient. The relevance of these findings to the creation of novel antimicrobial agents is briefly discussed.

When large unilamellar vesicles are first prepared via freezethawing and extrusion, the concentration of ionic solute within their aqueous interior equals that of the external phase.^{2,3} The membrane is thus formed under isotonic conditions, and its compactness is maximized through hydrophobic interactions.⁴

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 (16) The tautomers were in slow exchange at 300 MHz. The ester resonances in the ¹H NMR spectrum of 10 are doubled, with $\delta_{H-3} = 3.61$ ppm and $\delta_{\rm H.4}$ = 4.99 ppm (J \approx 0 Hz). Compound 10 appears to be a single diasteof 13 is highly symmetrical, with $\delta_{H,4} = 4.88$ ppm. (17) For an excellent overview, see: Wilcox, C. S. In *Frontiers in Su*-

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⁽¹⁸⁾ In the presence of acid this complex is unstable, undergoing clean hydride transfer from C-4 of 13 to C-10 of 12. The binding constants were determined in the presence of 1,8-bis(dimethylamino)naphthalene.

⁽¹⁹⁾ The present results do not indicate the origin of the variable stabilities, and other possibilities such as changes in the strength of the primary hydrogen bonds, differential solvation, alignment of molecular dipole moments, etc. cannot be ruled out.

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