

Graphical Abstract

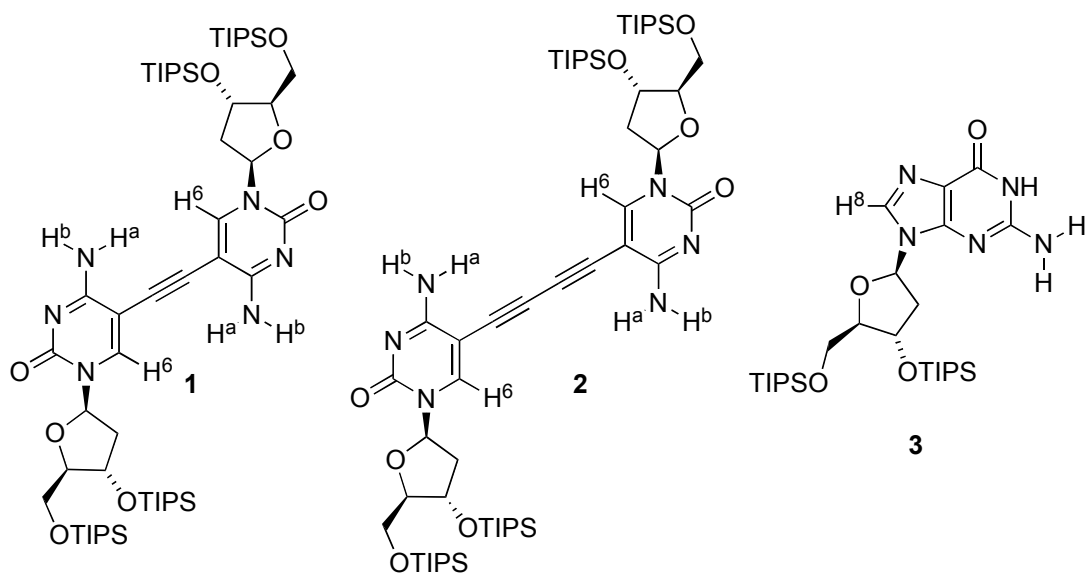
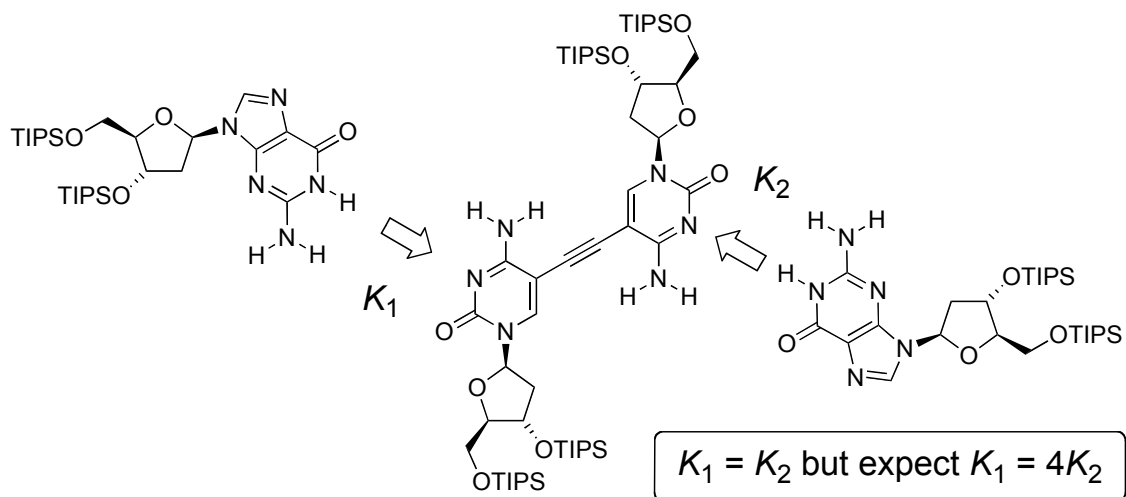


Figure 1. Receptors **1**, **2** and guanosine derivative, **3** used for binding studies; TIPS = triisopropylsilyl.

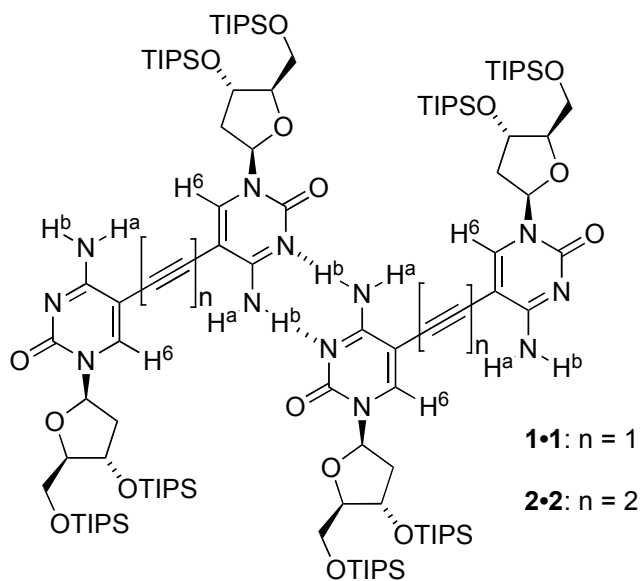


Figure 2. Dimers of **1** and **2**, **1•1** and **2•2**; respectively.

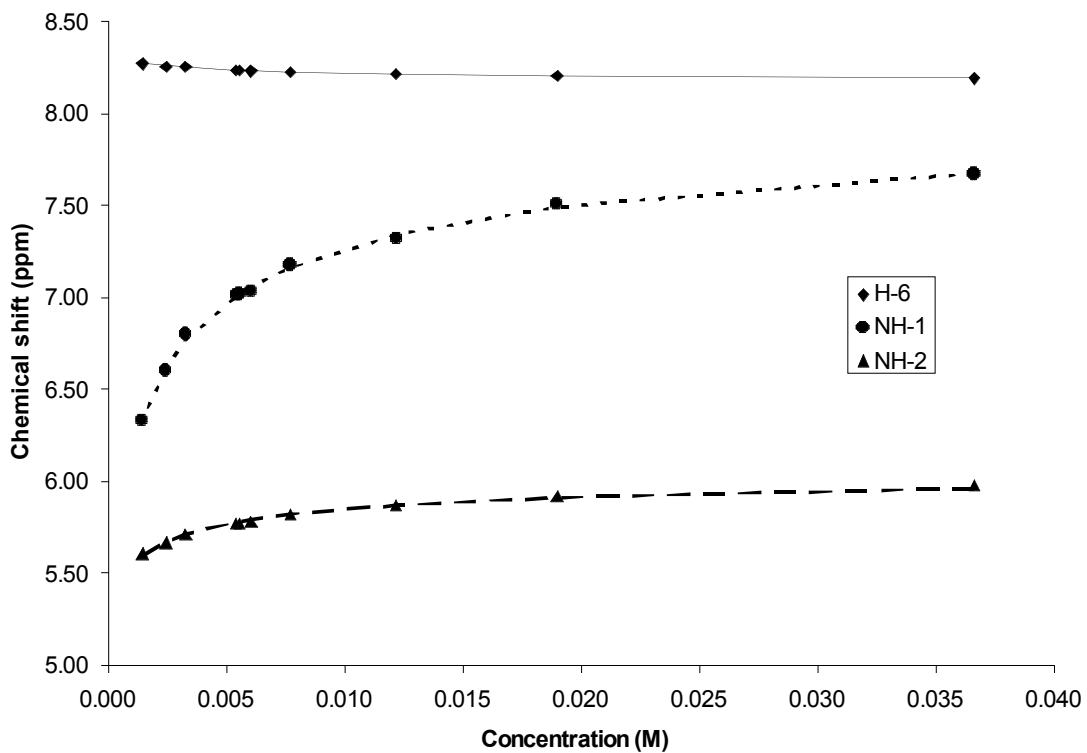


Figure 3. Curve fitting from NMR dilution of **2**; $K_{2,2} = 340 \pm 7 \text{ M}^{-1}$.

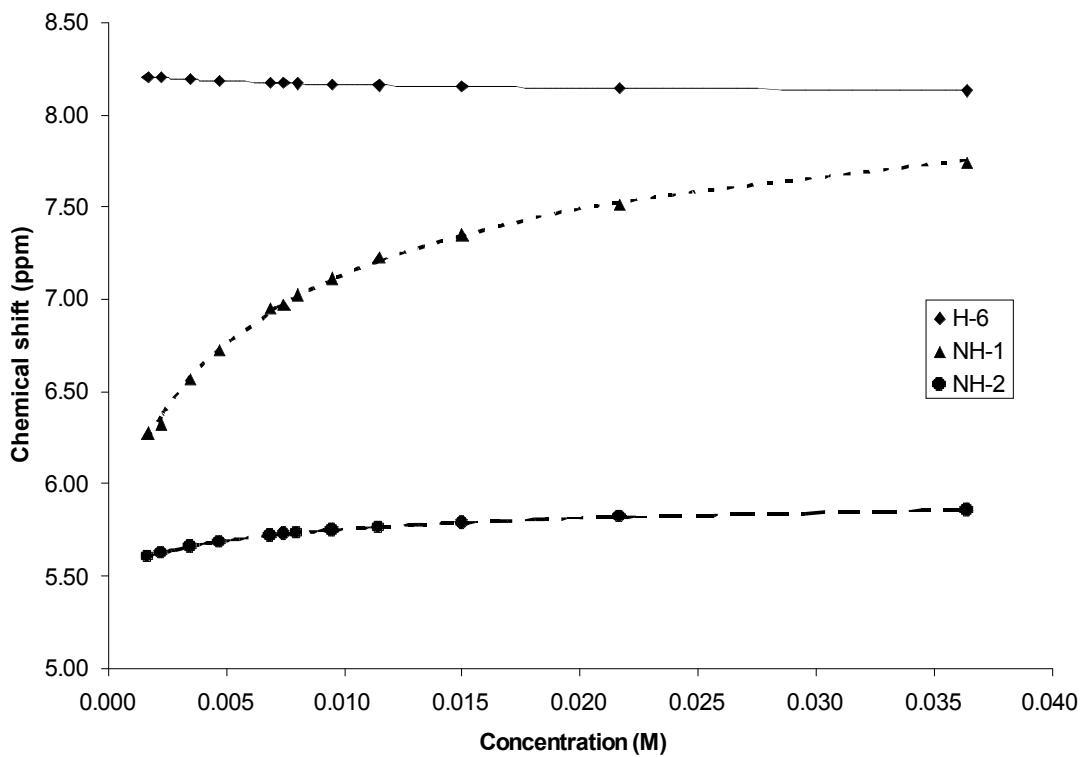


Figure 4. Curve fitting from NMR dilution of **1** in CDCl_3 ; $K_{1,1} = 83 \pm 3 \text{ M}^{-1}$.

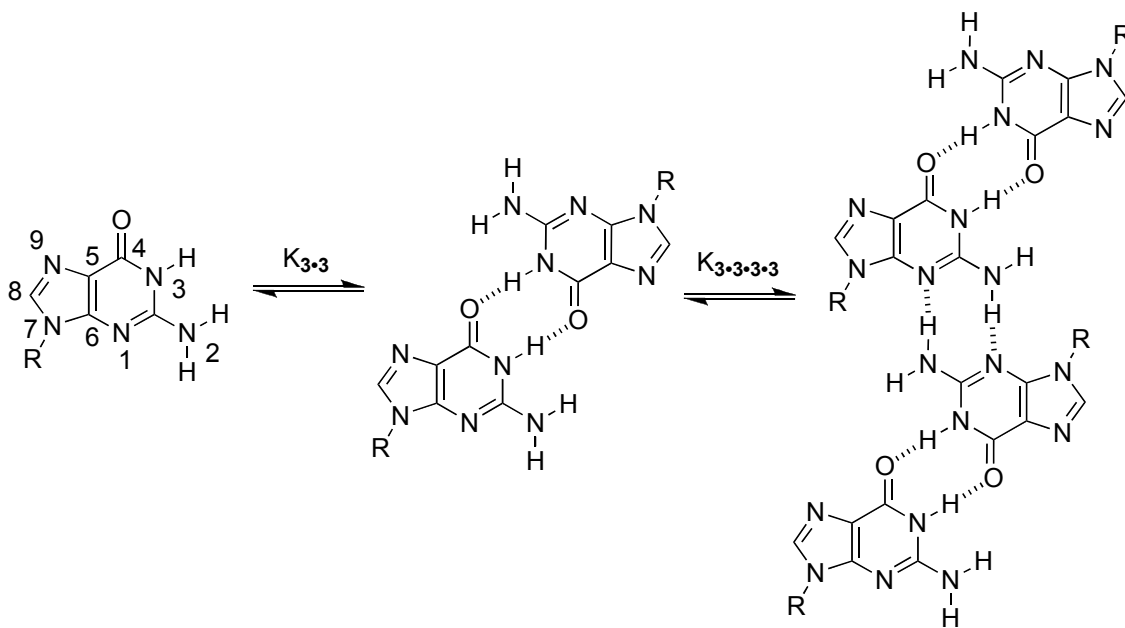
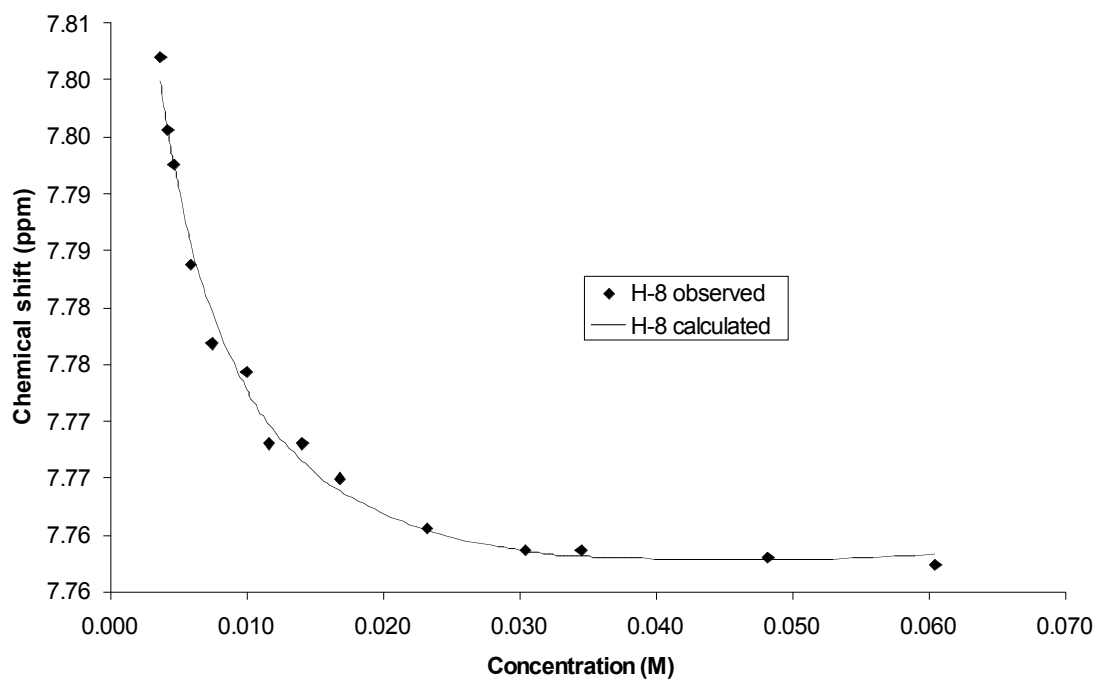
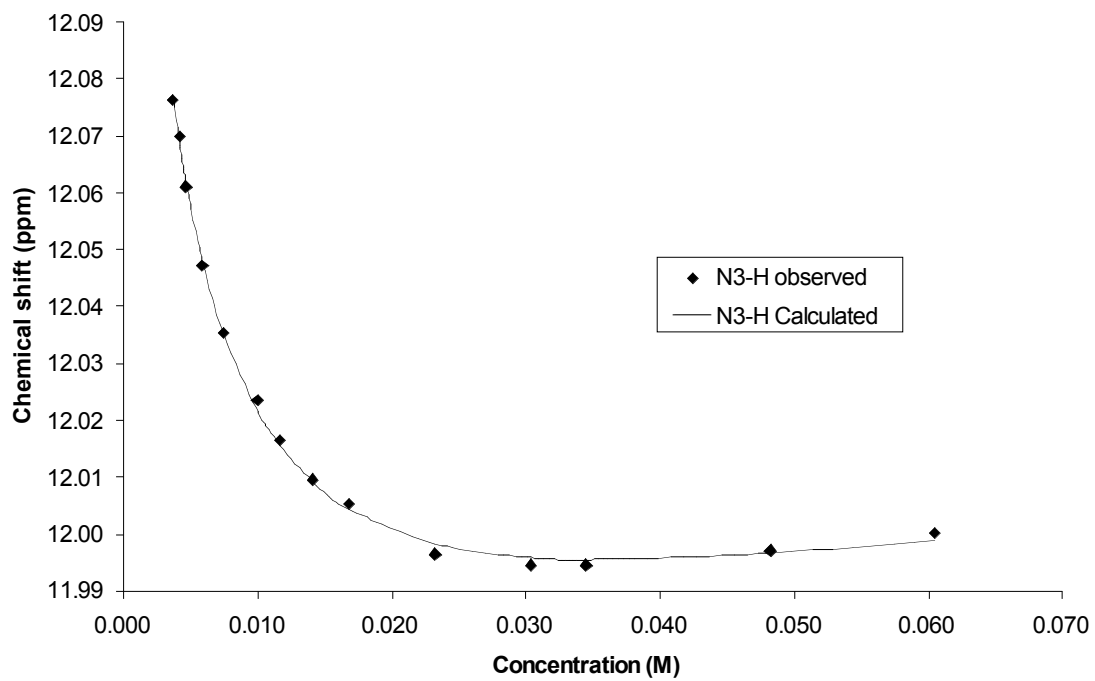


Figure 5. Self-association model of lipophilic guanosine **3** in chloroform.



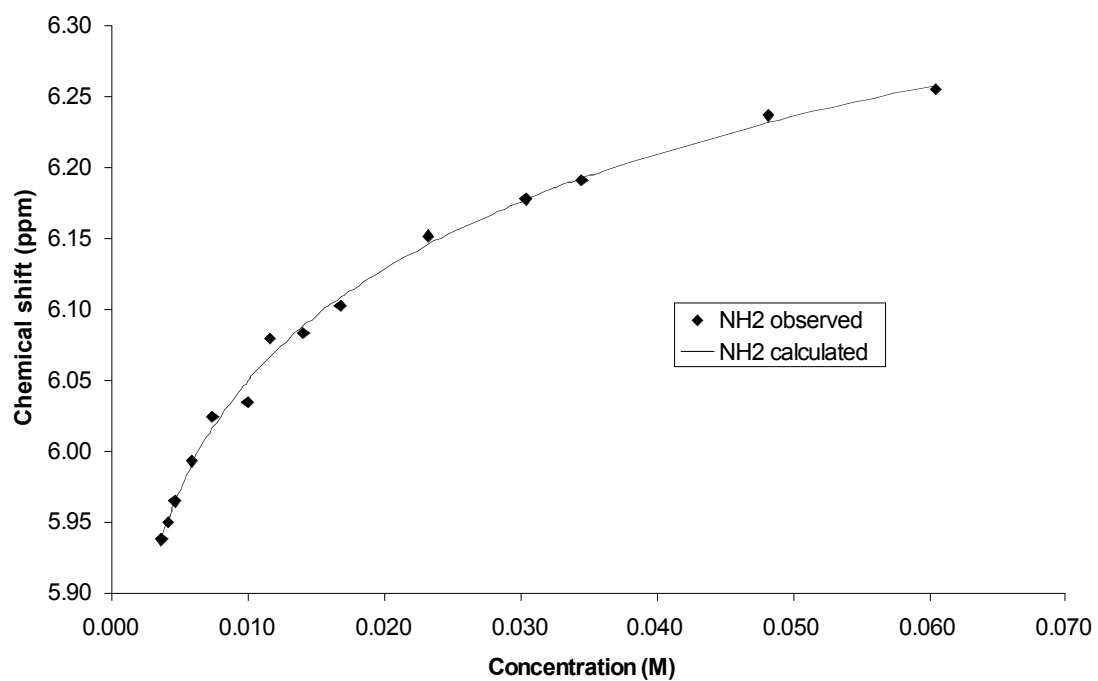


Figure 6. Curve fitting for NMR dilution of **3** in CDCl_3 . The three titration curves (A) N3-H (B) H-8 and (C) NH_2 are fitted simultaneously to give $K_{3,3}$ of 370 ± 72 and $K_{3,3,3}$ of $15 \pm 1 \text{ M}^{-1}$.

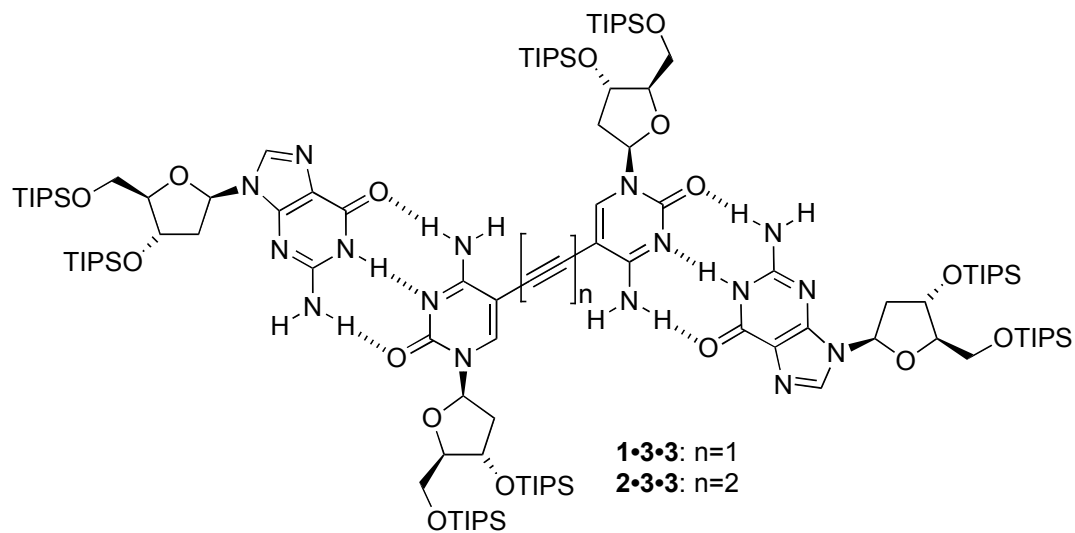
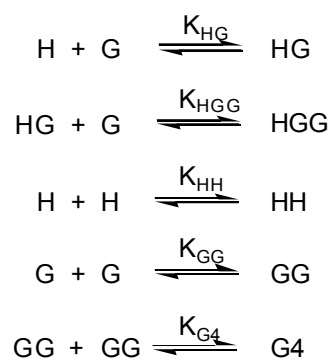


Figure 7. Complexation of receptors **1** or **2** with **3**; TIPS = triisopropylsilyl.



Scheme 1. Equilibria in the binding studies between **2** and **3**.

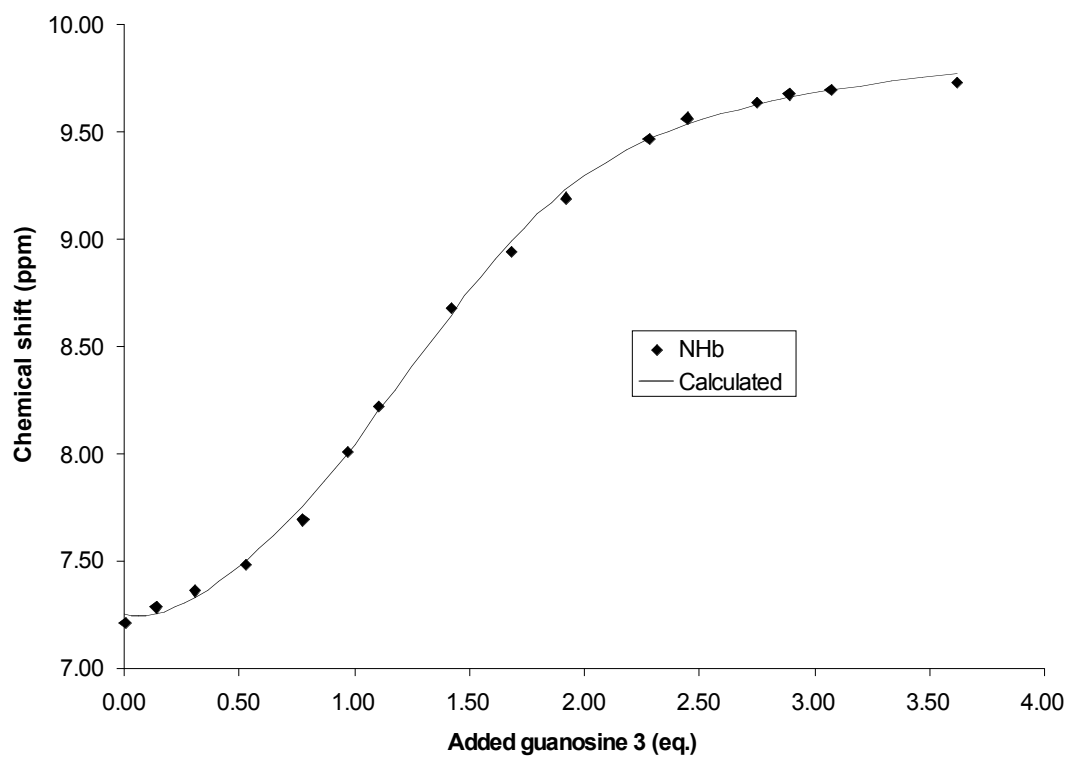


Figure 8. Curve fitting of NMR titration in CDCl_3 between **2** and **3**. $K_{2\cdot3}$ 8100 ± 380 ; $K_{2\cdot3\cdot3}$ $1170 \pm 80 \text{ M}^{-1}$.

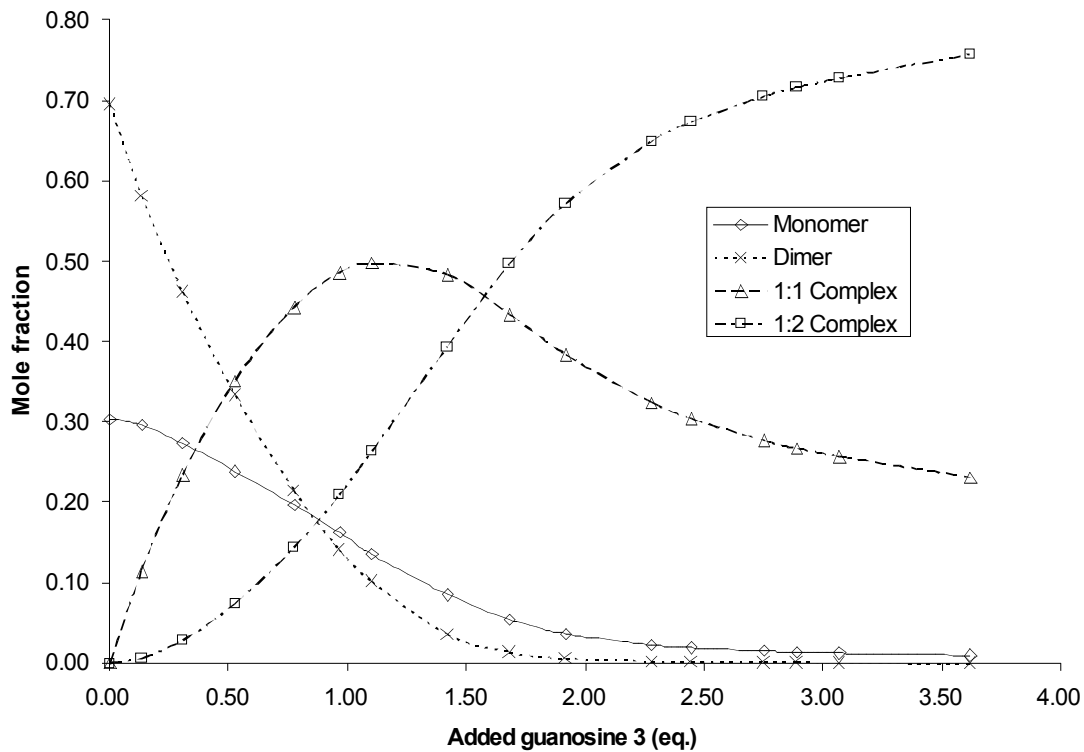


Figure 9. Speciation curve for receptor **2** upon addition of **3** in CDCl_3 , showing monomeric **2**, dimeric **2**, 1:1 complex (**2**•**3**) and 1:2 complex (**2**•**3**•**3**).

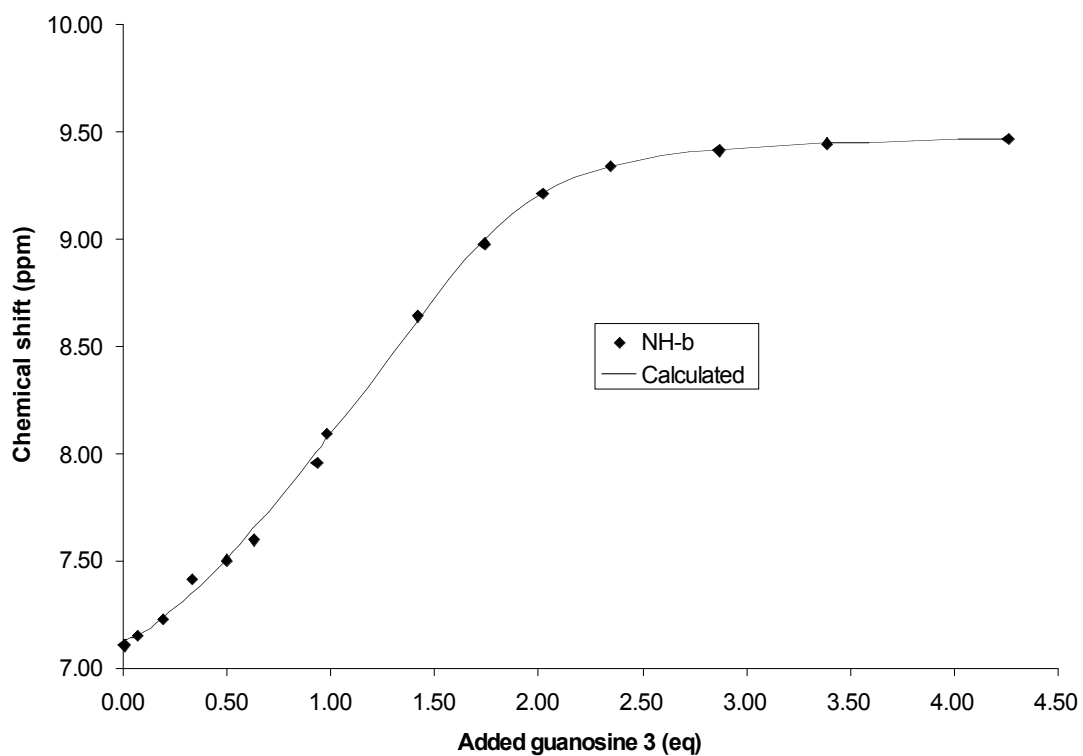


Figure 10. Curve fitting of NMR titration in CDCl_3 between **1** and **3**. $K_{1,3} = 5180 \pm 210$, $K_{1+3,3} = 4800 \pm 170 \text{ M}^{-1}$.

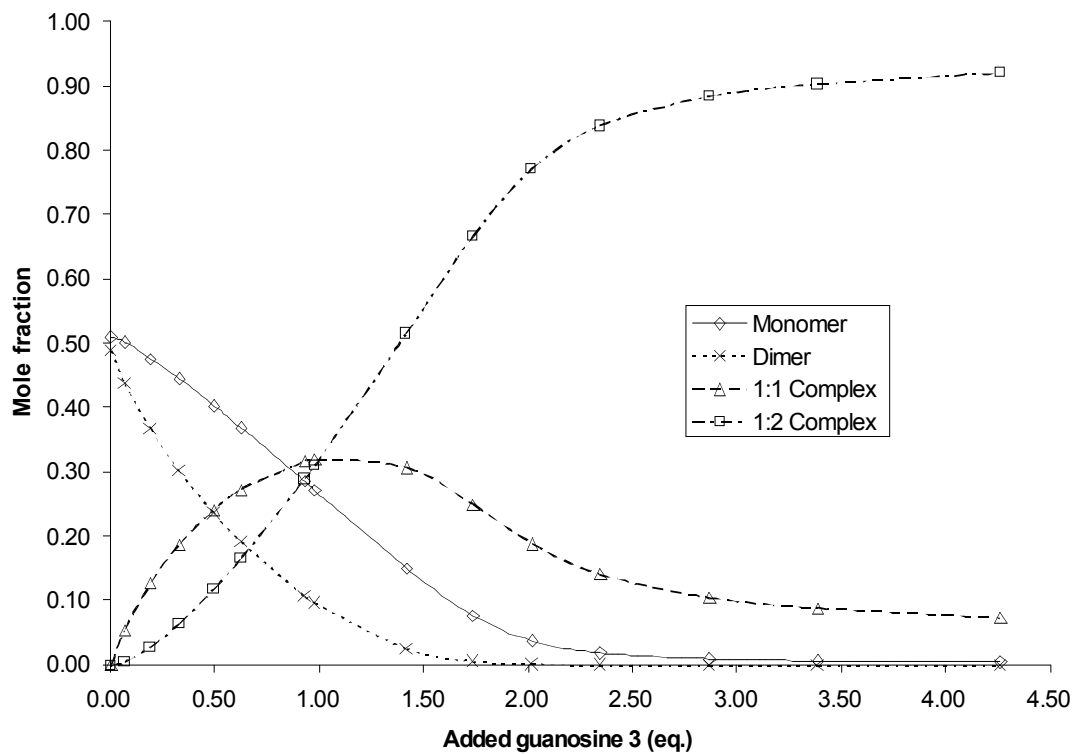


Figure 11. Speciation curve for **1** upon addition of **3** in CDCl_3 .

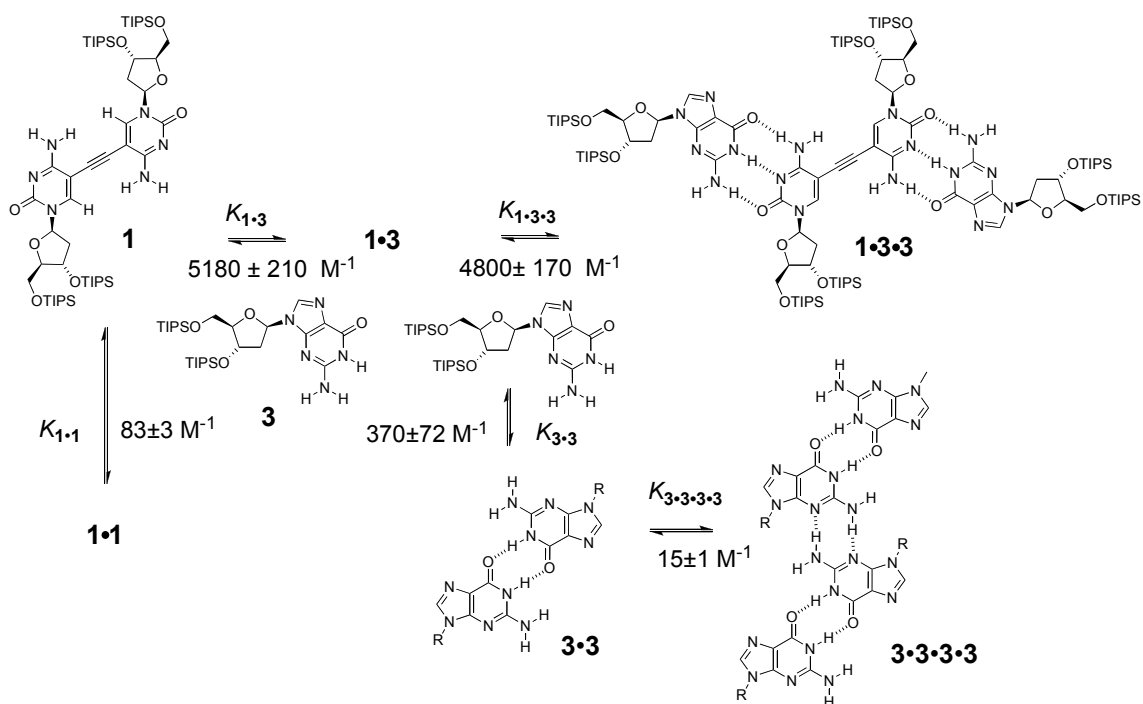


Figure 12. Summary of equilibria in deuteriochloroform for monoalkyne receptor **1**.

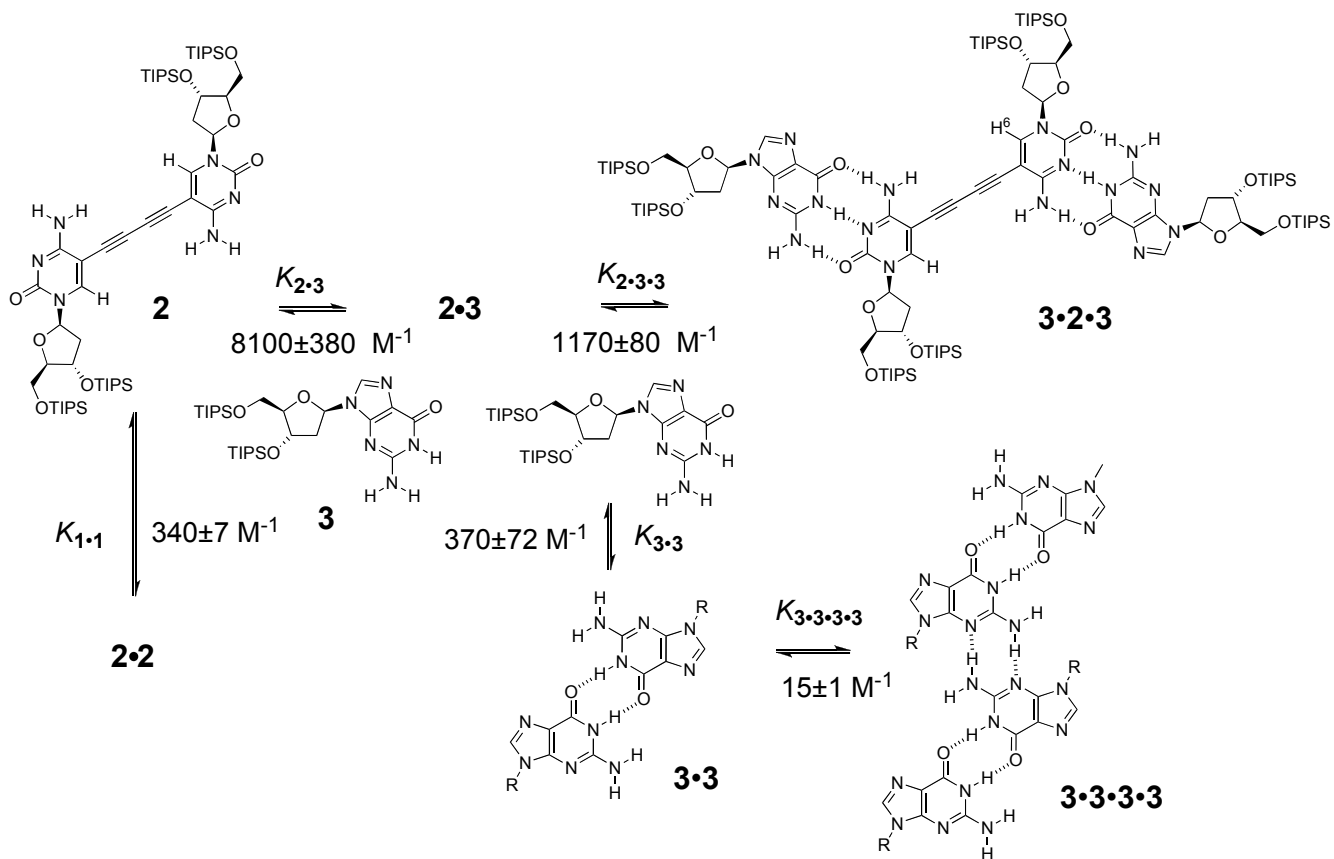


Figure 13. Summary of equilibria in deuteriochloroform for dialkyne receptor **2**.

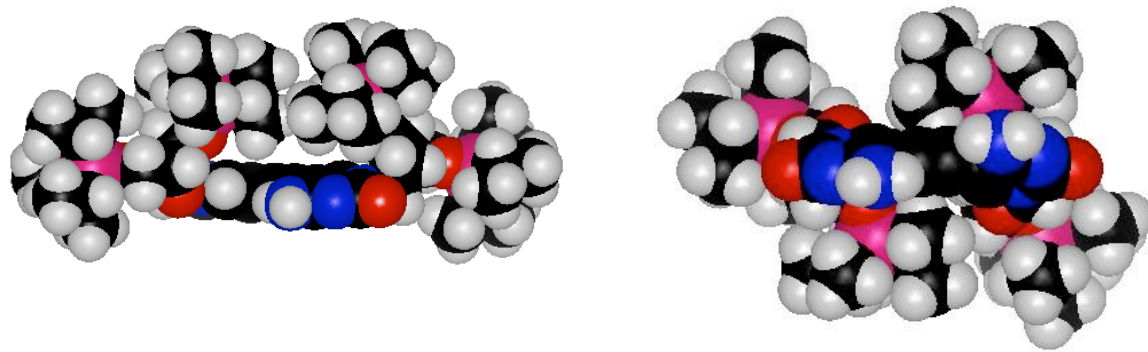


Figure 14. C-P-K models of **1** at E_{\min} (left) dihedral $\approx 30^\circ$ showing interactions between TIPS groups and E_{\max} (right) dihedral $\approx 165^\circ$.

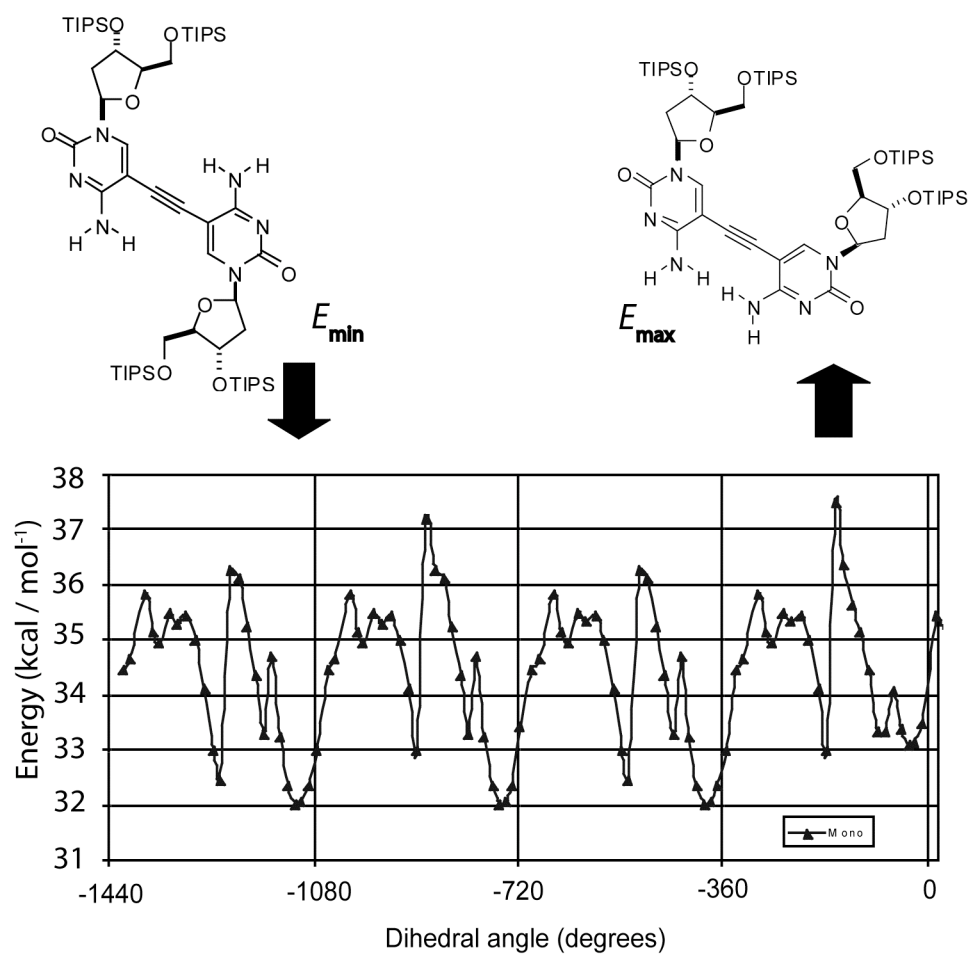


Figure 15. Torsional energy vs. dihedral angle for monoalkyne 1

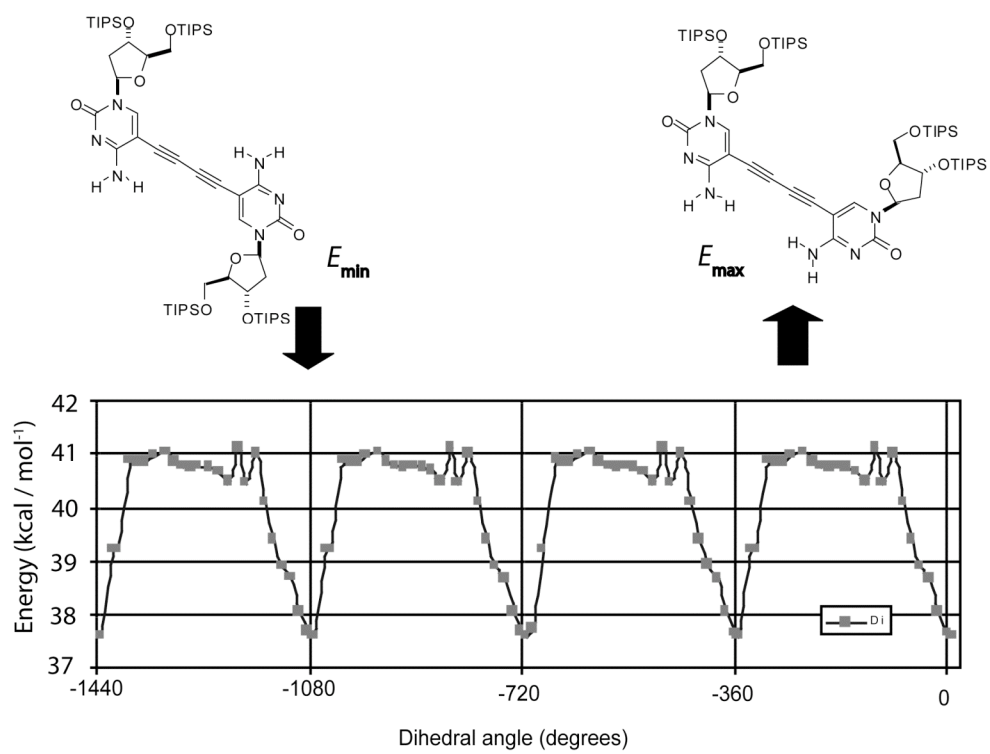


Figure 16. Torsional energy vs. dihedral angle for dialkyne 2