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Cleft Type Receptors for Butenolides Based on Chromenone Derivatives

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Abstract: A chromenone building block is suitable for the preparation of H-bonding lactone receptors. Combination of this fragment with a second chromenone provides a new cleft suitable for strongly association of γ -hydroxymethyl- γ -lactones.

But-2-enolides display broad chemical reactivity in nucleophilic addition¹ or as dienophiles in electrocyclic reactions². Such molecules also present strong biological activity³. Butenolides are therefore interesting targets for Molecular Recognition. However, to our knowledge, so far very few receptors have been found to be able to associate lactones⁴.

Study of molecular models shows that chromenone fragments, as in receptor **5**, may be excellent candidates for lactones **1-3**, providing the necessary geometry to set three linear H-bonds (Fig. 1).

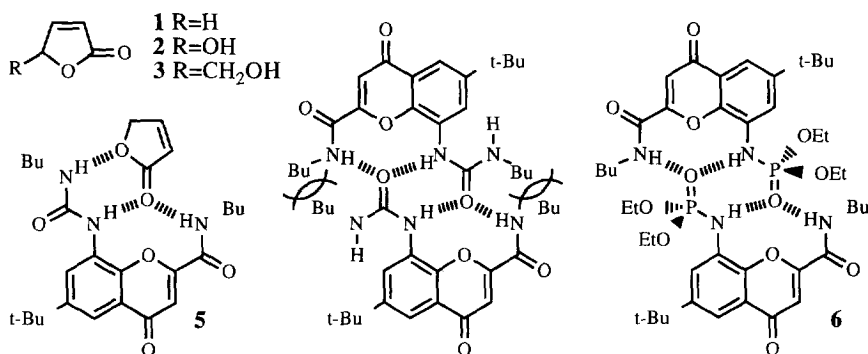
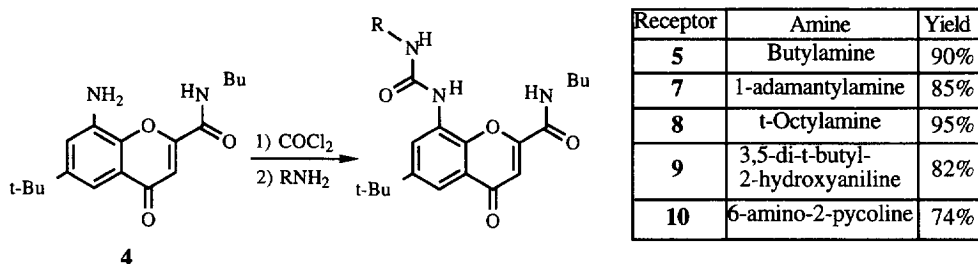


Figure 1. Guests, complex between receptor **5** and lactone **1** and proposed dimers for receptors **5** and **6**

Receptors **5** and **6** were prepared easily starting from the aminochromane **4**⁵ as shown in schemes 1 and 2, respectively. Receptor **5** (m.p.= 122-126°C) is essentially unable to bind butyrolactone in CDCl₃, probably because **5** shows a self-complementary structure which leads to dimerization in this solvent. A value of $K_d = 5.0 \times 10^2 \text{ M}^{-1}$ for the dimerization is afforded in ¹HNMR experiments⁶. Scrutiny of molecular models suggests a

certain degree of steric hindrance between the urea and amide butyl chains of different molecules (Fig. 1). This fact is also supported because a similar compound **6** (m.p.= 154-156°C), in which the urea has been changed into a phosphoramidate group, presents a stronger self-association, which ¹HNMR studies can only evaluate above 10⁴ M⁻¹. In this case, the tetrahedral geometry of the phosphorus atom prevents interference between the alkyl chains in the dimer units.



Scheme 1. Synthesis of urea-type receptors for lactones **1** and **2**

The presence of steric effects in the receptor **5** dimer should allow easy preparation of receptors with lower self-association (Scheme 1). The bulky adamantyl group in receptor **7** (Fig. 2, m.p.= 222-224°C) collides in its dimer with the butyl chain, reducing the self-association to $K_d=1.5 \times 10^2$ M⁻¹, and a similar effect is obtained for the t-octyl derivative **8** (m.p.= 178-180°C), which shows $K_d=1.0 \times 10^2$ M⁻¹.

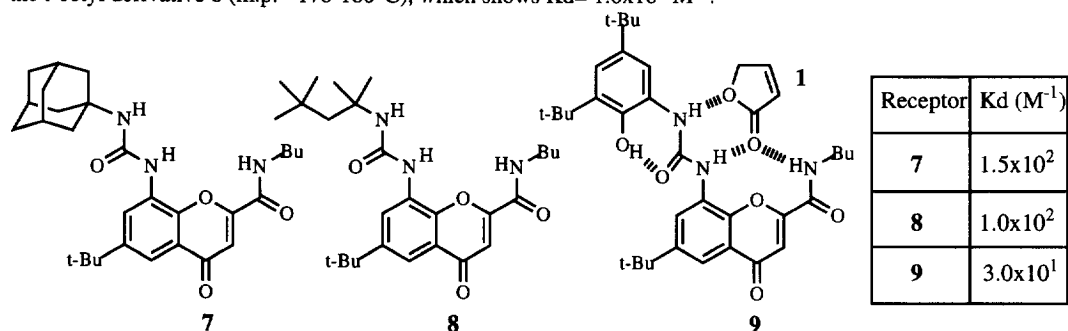


Figure 2. Structures for receptors **7** to **9** and their self-association constants

The best results to avoid self-association are obtained when a smaller steric effect is combined with an intramolecular H-bond, as in receptor **9** (m.p.= 154-156°C). In this case, self-association is reduced to $K_d=3.0 \times 10^1$ M⁻¹. This small dimerization constant allows one to study the association constant with 2-(5H)-furanone **1** (Fig. 2). The association constant⁷ was measured in diluted solution (3×10^{-3} M) with a large excess of the butenolide **1** to prevent a strong interference by the dimerization process. Under these conditions, a $K_{ass}=3.0 \times 10^1$ M⁻¹ was obtained. No change was obtained for this association constant when dimer formation was taken into account⁸.

Hydroxylactones such as 5-hydroxy-2-(5H)-furanone **2** offer the possibility of setting more H-bonds in the complex and therefore could provide higher association constants. Since aminopyridines are known to be good hydrogen bond acceptors⁹, one of these units was included in receptor **10** (m.p.= 216-218°C) to provide a fourth hydrogen bond with the hydroxyl group. However, both the dimer formation ($K_d=1.6 \times 10^2$ M⁻¹) and the lactone **2** association constant ($K_{ass}=3.2 \times 10^1$ M⁻¹) are surprisingly low for this receptor **10**, probably due to the presence of a competing six membered ring intramolecular hydrogen bond (Fig. 3).

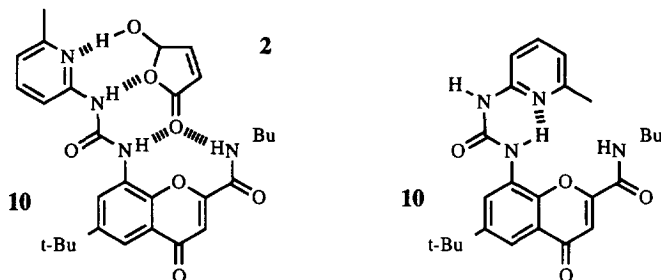
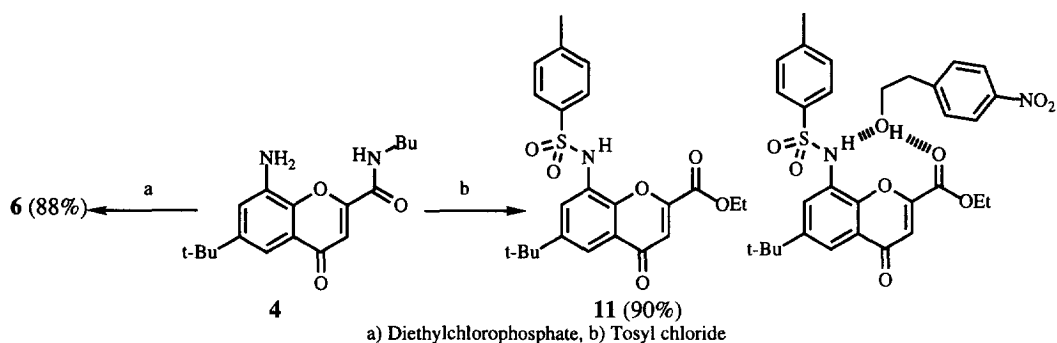


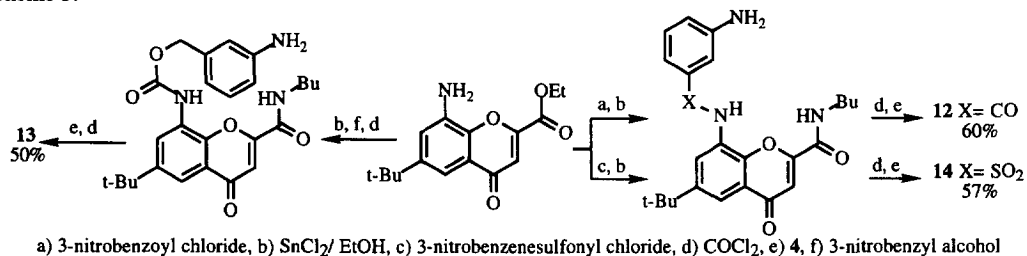
Figure 3. Proposed structures for the complex of receptor **10** with hydroxylactone **2** and its possible intramolecular H-bond

A chromenone unit is also able to associate alcohols because two strong linear hydrogen bonds can be set with the hydroxyl group. For example, the association constant for the sulfonamide **11** (prepared as shown in scheme 2, m.p.= 144-146°C) and the crystalline 2-(4-nitrophenyl)-ethanol is $K_{ass}=9.0\text{ M}^{-1}$.



Scheme 2. Synthesis of receptors **6** and **11** and the proposed complex for receptor **11** and 2-(4-nitrophenyl)-ethanol

CPK molecular models suggest three readily available spacers to link both lactone and hydroxyl binding sites of dichromenone receptors **12-14** in order to fit 5-hydroxymethyl-2-(5*H*)-furanone **3** (Fig. 4). The greater distance between the OH group and the lactone compared to lactone **2** in this guest allows an easier design of receptors lacking intramolecular hydrogen bonds. Easy preparation of these receptors was carried out as shown in scheme 3.



Scheme 3. Preparation of compounds **12** to **14**

The benzoic unit of receptor **12** (m.p.= 186-190°C) is the most rigid spacer. However, it shows the worst fit with hydroxylactone **3** and consequently the complex has poor stability, with $K_{ass}=8.5 \times 10^1\text{ M}^{-1}$. The flexible spacer in receptor **13** (m.p.= 224-226°C) allows a good model fit with guest **3**, although many rotational degrees of freedom must be frozen in the complex, leading to poor cooperativity between both binding sites. In this case,

an apparent association constant can be measured $K_{\text{ass}} = 1.3 \times 10^2 \text{ M}^{-1}$. Correction due to dimer formation ($K_{\text{d}} = 7.0 \times 10^2 \text{ M}^{-1}$) leads to a small increase up to $K_{\text{ass}} = 1.7 \times 10^2 \text{ M}^{-1}$.

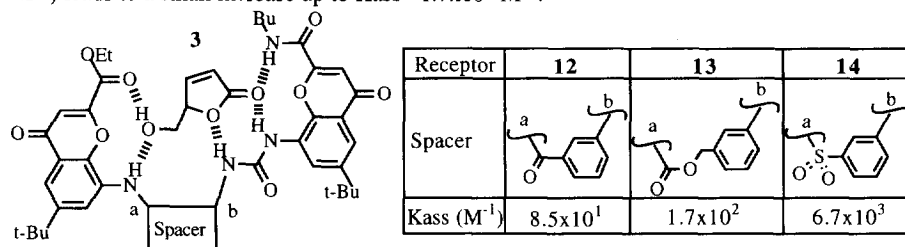


Figure 4. Association constants and proposed structure for the complexes of receptors **12** to **14** with hydroxylactone **3**

A phenylsulfonyl spacer provides the best compromise between rigidity and fit in receptor **14** (m.p. = 178–182°C). This compound again shows self-association with a $K_{\text{d}} = 5.0 \times 10^2 \text{ M}^{-1}$. The association constant with hydroxylactone **3** measured in a diluted solution ($3 \times 10^{-4} \text{ M}$) to prevent strong dimer interference, showed $K_{\text{ass}} = 4.2 \times 10^3 \text{ M}^{-1}$. Correction of the data due to the small amount of dimer leads to $K_{\text{ass}} = 6.7 \times 10^3 \text{ M}^{-1}$. This high association constant indicates that cooperativity between both binding sites is good, making further work on these lactone receptors promising.

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- 6 Dimerization constants were measured following proton shifts in dilution studies (concentrations from 10^{-2} M to 10^{-4} M) in CDCl_3 at 293K. A Monte-Carlo non-linear curve-fitting program was used to evaluate the self-association.
- 7 Association constants were measured in CDCl_3 at 293K. Samples a with constant concentration of host (variable from 5×10^{-3} to $3 \times 10^{-4} \text{ M}$, depending on the association constant range) and an increasing concentration of guest were prepared for ^1H NMR spectra. A Monte-Carlo non-linear curve-fitting program was used to calculate the value of the constant. A similar program allows correction due to self association.
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