

Dopamine Interaction in the Absence and in the Presence of Cu²⁺ Ions with Macrocyclic and Macrobicyclic Polyamines Containing Pyrazole Units. Crystal Structures of [Cu₂(L₁)(H₂O)₂](ClO₄)₄ and [Cu₂(H₋₁L₃)](ClO₄)₃·2H₂O

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Abstract: The interaction with Cu²⁺ and dopamine of three polyazacyclophanes containing pyrazole fragments as spacers is described. Formation of mixed complexes Cu²⁺–macrocyclic–dopamine has been studied by potentiometric methods in aqueous solution. The crystal structures of the complexes [Cu₂(L₁)(H₂O)₂](ClO₄)₄·2H₂O (**4**) (L₁ = 13,26-dibenzyl-3,6,9,12,13,16,19,22,25,26-decaazatricyclo[22.2.1.1^{11,14}]octacos-1(27),11,14(28),24-tetraene) and [Cu₂(H₋₁L₃)](HClO₄)(ClO₄)₂·2H₂O (**6**) (L₃ = 1,4,7,8,11,14,17,20,21,24,29,32,33,36-tetradecaazapentacyclo[12.12.12.1^{6,9}.1^{19,22}.1^{31,34}]hentetraconta-6,9(41),19(40),21,31,34(39)-hexaene) are presented. In the first one (**4**), each Cu²⁺ coordination site is made up by the three nitrogens of the polyamine bridge, a sp² pyrazole nitrogen and one water molecule that occupies the axial position of a square pyramid. The distance between the copper ions is 6.788(2) Å. In the crystal structure of **6**, the coordination geometry around each Cu²⁺ is square pyramidal with its base being formed by two secondary nitrogens of the bridge and two nitrogen atoms of two different pyrazolate units which act as exobidentate ligands. The axial positions are occupied by the bridgehead nitrogen atoms; the elongation is more pronounced in one of the two sites [Cu(1)–N(1), 2.29(2) Å; Cu(2)–N(6), 2.40(1) Å]. The Cu–N distances involving the deprotonated pyrazole moieties are significantly shorter than those of the secondary nitrogens. The Cu(1)···Cu(2) distance is 3.960(3) Å. The pyrazole in the noncoordinating bridge does not deprotonate and lies to one side of the macrocyclic cavity. One of the aliphatic nitrogens of this bridge is protonated and hydrogen bonded to a water molecule, which is further connected to the sp² nitrogen of the pyrazole moiety through a hydrogen bond. The solution studies reveal a ready deprotonation of the pyrazole units induced by coordination to Cu²⁺. In the case of L₂ (L₂ = 3,6,9,12,13,16,19,22,25,26-decaazatricyclo[22.2.1.1^{11,14}]octacos-1(27),11,14(28),24-tetraene), deprotonation of both pyrazole subunits is already observed at pH ca. 4 for 2:1 Cu²⁺:L₂ molar ratios. All three free receptors interact with dopamine in aqueous solution. L₃ is a receptor particularly interesting with respect to the values of the interaction constants over five logarithmic units at neutral pH, which might suggest an encapsulation of dopamine in the macrocyclic cage. All three receptors form mixed complexes Cu²⁺–L–dopamine. The affinity for the formation of ternary dopamine complexes is particularly high in the case of the binuclear Cu²⁺ complexes of the 1-benzyl derivative L₁.

Introduction

Supramolecular chemistry bases its progress in the design of highly organized molecular receptors able to discriminate and/or induce characteristic properties in given substrates.¹ Among the wealth of molecules that can be the target of recognition, neurotransmitter catecholamines are particularly interesting due to their many biological implications. Concretely, dopamine is involved in the normal emotional and autonomic control of humans. The physiological level of dopamine is altered in

neurodegenerative and mental illnesses as well as in toxic syndromes induced by cocaine and psychotropic drugs.² Therefore receptors able to modulate the dopamine level by selective complexation or transport mechanisms represents a research goal of great interest.

In this respect, azacyclophanes containing pyrazole spacers can be adequate candidates since, apart from disposing of polyamine sites able to coordinate a variety of guests, their own pyrazole spacer may cooperate in the recognition event. Polyamine receptors can behave as ambivalent receptors since

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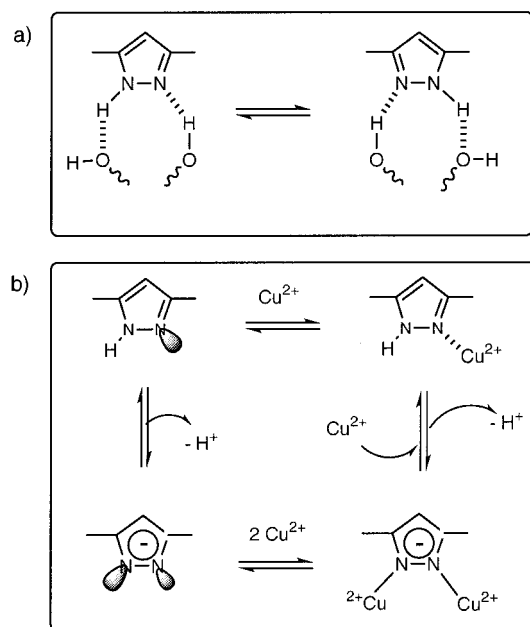
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Scheme 1



their partly or fully protonated forms can interact electrostatically or/and through hydrogen bonds with anionic or polar substrates such as catechol while, when deprotonated and, thus, disposing of free lone pairs, they can interact with metal ions acting as Lewis bases.^{3,4}

With respect to dopamine coordination, the pyrazole unit may participate in the binding donating or accepting hydrogen bonds through its NH or its sp^2 N groups, respectively (Scheme 1a). On the other hand, the neutral pyrazole unit can provide coordinative bonds to metal ions as an electron donor through the sp^2 nitrogen atom. When pyrazole deprotonates to give the pyrazolate anion, it can also act as a bridging ligand between metal centers (Scheme 1b).^{5,6}

Another possible coordination mode results from the use of either coordinatively unsaturated metal complexes or metal complexes with labile ligands in their first coordination sphere. These metal complexes behave as Lewis acids and may coordinate different substrates as exogenous ligands, giving rise to the formation of mixed or ternary complexes. Dopamine is a good candidate for this type of recognition due to the high strength of the catechol–copper(II) bonds and to the relevance of such complexes in biology.^{7,8}

Previously, we reported on the synthesis of 13,26-dibenzyl-3,6,9,12,13,16,19,22,25,26-decaazatricyclo[22.2.1.1^{11,14}]octacosal-1(27),11,14(28),24-tetraene (L_1), 3,6,9,12,13,16,19,22,25,26-decaazatricyclo[22.2.1.1^{11,14}]octacosal-1(27),11,14(28),24-tetraene

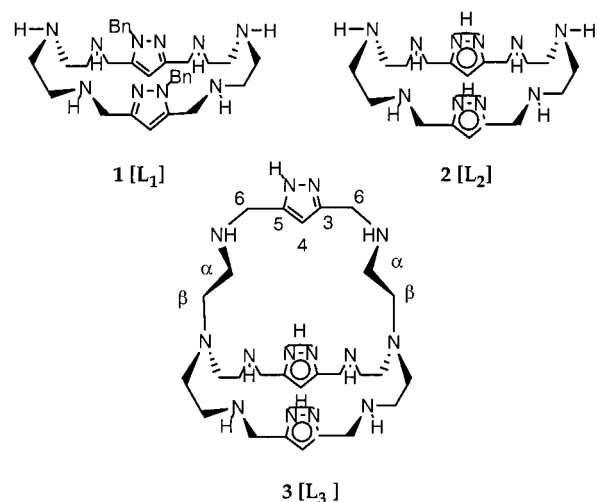
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(L_2), and the cryptand 1,4,7,8,11,14,17,20,21,24,29,32,33,36-tetradecaazapentacyclo[12.12.12.1^{6,9}.1^{19,22}.1^{31,34}]hentetraconta-6,9(41),19(40),21,31,34(39)-hexaene (L_3), as well as on the protonation behavior of L_1 and L_2 .^{9–11}



It was demonstrated that L_1 and L_2 present six protonation steps in aqueous solution that correspond to the number of nitrogen atoms in the polyamine bridges and that in the pH range 2–11 the pyrazole rings were not involved in protonation or deprotonation processes. On the other hand, a careful ^1H and ^{13}C NMR study of the 1-benzyl substituted receptor L_1 proved that its structure corresponds to a unique conformational isomer with the benzyl substituents located each other in opposite position.¹¹ Additionally, some of us have also presented the crystal structures of a binuclear Cu^{2+} complex of L_1 in which the pyrazole moieties were deprotonated acting as exobidentate ligands.¹²

In the present work we explore the capabilities of receptors L_1 , L_2 , and L_3 to coordinate Cu^{2+} and to interact with dopamine in aqueous solution, both in the absence and in the presence of Cu^{2+} ions. We show as Cu^{2+} coordination induces the ready deprotonation of the 1H-pyrazole rings at remarkably low pH values and how L_3 seems to be a particularly well-suited receptor for the recognition of dopamine in water. Additionally, we present the crystal structures of the complexes $[\text{Cu}_2(\text{L}_1)(\text{H}_2\text{O})_2](\text{ClO}_4)_4 \cdot 2\text{H}_2\text{O}$ and $[\text{Cu}_2(\text{H-L}_3)(\text{ClO}_4)_3] \cdot 2\text{H}_2\text{O}$. (Some of these results have been advanced in the following: Lamarque, L.; Miranda, C.; Navarro, P.; Escartí, F.; García-España, E.; Latorre, J.; Ramírez, J. A. *Chem. Commun.* **2000**, 1337.)

Results and Discussion

Dopamine Recognition. Our first objective was the study of the interaction of receptors L_1 – L_3 with dopamine in aqueous solution. In Table 1 are presented the stability constants for the

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Table 1. Equilibrium Constants for the Interaction of Dopamine with the Receptors L_1 , L_2 , and L_3 Determined in 0.15 mol dm⁻³ NaCl at 298.1 K

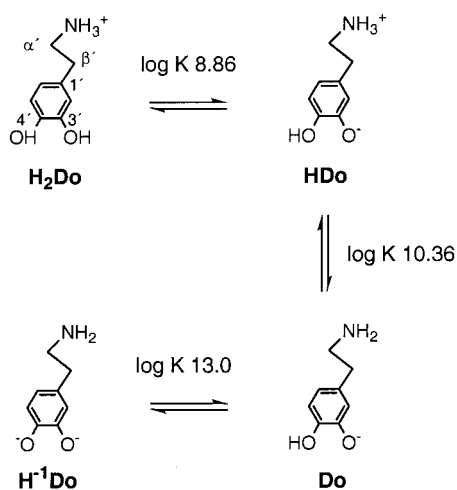
entry	reaction ^a	L_1	L_2	L_3
1	Do + L + H \rightleftharpoons H(Do)L			16.19(2)
2	Do + 2H + L \rightleftharpoons H ₂ (Do)L			25.72(2)
3	Do + 3H + L \rightleftharpoons H ₃ (Do)L	31.16(8)	32.08(4)	34.57(2)
4	Do + 4H + L \rightleftharpoons H ₄ (Do)L	39.42(6)	40.71(3)	43.04(2)
5	Do + 5H + L \rightleftharpoons H ₅ (Do)L	46.49(6)	48.13(4)	50.39(2)
6	Do + 6H + L \rightleftharpoons H ₆ (Do)L	52.52(6)	55.51(4)	57.08(2)
7	Do + 7H + L \rightleftharpoons H ₇ (Do)L	56.38(7)	59.96(6)	63.40(2)
8	Do + 8H + L \rightleftharpoons H ₈ (Do)L	59.72(6)	63.04(7)	
9	HDo + L \rightleftharpoons H(Do)L			5.8
10	HDo + HL \rightleftharpoons H ₂ (Do)L			5.9
11	HDo + H ₂ L \rightleftharpoons H ₃ (Do)L	3.6	3.1	6.0
12	H ₂ Do + HL \rightleftharpoons H ₃ (Do)L	3.1	3.1	5.8
13	H ₂ Do + H ₂ L \rightleftharpoons H ₄ (Do)L	3.0	2.9	5.6
14	H ₂ Do + H ₃ L \rightleftharpoons H ₅ (Do)L	3.4	2.9	5.3
15	H ₂ Do + H ₄ L \rightleftharpoons H ₆ (Do)L	3.6	2.9	5.2
16	H ₂ Do + H ₅ L \rightleftharpoons H ₇ (Do)L	3.8	2.8	5.4
17	H ₂ Do + H ₆ L \rightleftharpoons H ₈ (Do)L	5.0	2.7	

^a Charges omitted for clarity. ^b Values in parentheses are standard deviations in the last significant figure.

Table 2. Protonation Constants of Receptors L_1 – L_3 Determined in 0.15 mol dm⁻³ NaCl at 298.1 K

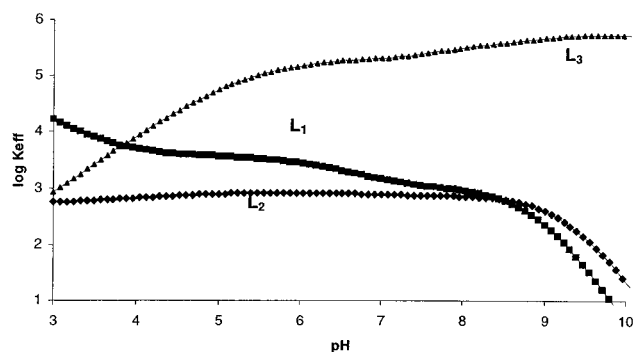
reaction	L_{1a}	L_{2a}	L_3
H + L \rightleftharpoons HL ^b	8.90	9.74	9.54(2) ^c
H + HL \rightleftharpoons H ₂ L	8.27	8.86	8.71(1)
H + H ₂ L \rightleftharpoons H ₃ L	6.62	7.96	7.72(2)
H + H ₃ L \rightleftharpoons H ₄ L	5.85	6.83	6.55(3)
H + H ₄ L \rightleftharpoons H ₅ L	3.37	4.57	6.51(3)
H + H ₅ L \rightleftharpoons H ₆ L	2.27	3.19	5.22(3)

^a Taken from ref 11. ^b Charges omitted for clarity. ^c Values in parentheses are standard deviations in the last significant figure.

Scheme 2

formation of the different adducts dopamine–receptor determined by pH-metric titration in the pH range 2.5–9.0. A step required for obtaining the adduct stability constants is the previous determination of the protonation constants of receptors and substrate under the same experimental conditions. The values for such constants are included in Table 2 and Scheme 2, respectively.

The stoichiometries derived from the analysis of the potentiometric data are always 1:1 receptor:dopamine. L_1 and L_2 form dopamine adducts with protonation degrees that vary between 3 ($[H_3LD_2]^{2+}$ species) and 8 ($[H_8LD_2]^{7+}$ species), while for L_3 the protonation degrees change from 1 to 8 ($[HL_3Do]$ and $[H_8L_3Do]^{7+}$ species) throughout the pH range explored (pH

**Figure 1.** Plot of the logarithms of the conditional constants (K_{eff}) vs pH for the systems L_1 –Do, L_2 –Do, and L_3 –Do.

2.5–9.5). At physiological pH the average protonation degree of the adducts formed is around 5 in all three systems. However, the protonation degrees achieved by the different systems deserve some comment. First of all, to understand the actual distribution of protons between receptor and substrate, the protonation constants of the receptors and of dopamine have to be evaluated. We define “Do” as the mononegatively charged species in which the amino group and one of the hydroxyl groups are deprotonated. The protonation of the amino group in our experimental conditions to give HDo presents $\log K_{\text{HDo}/\text{HDo}} = 10.36$, and the protonation of the phenolate group to yield the positively charged H_2Do^+ species has $\log K_{\text{H}_2\text{Do}/\text{HDo}} = 8.86$ (see Scheme 2). Therefore, in view of these values H_2Do^+ is the prevalent species at pH 7.4, and this also has to be the situation in the binary system receptor–dopamine, unless the interaction with the receptors dramatically changes the acidity of dopamine, which is not the case of the systems here reported.

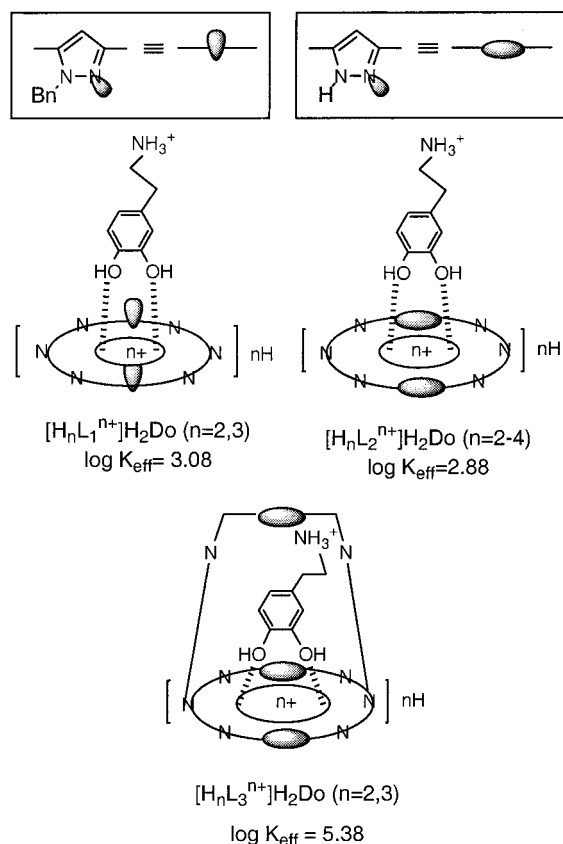
Therefore, taking into account this consideration and the protonation constants of the receptors (Table 2), the species $[H_5LD_2]^{4+}$ should be the result of the reaction between H_2Do^+ and H_3L^{3+} , and the $[H_6LD_2]^{5+}$ product, of the reaction of H_2Do^+ and H_4L^{4+} . The same reasoning shows that HDo is the intervening species in the formation of the mono- and diprotonated adducts. With all these considerations in mind, the overall association constants can be simplified to the step-wise constants shown at the bottom of Table 1. However, this analysis is not so straightforward for the species with intermediate protonation degrees (3 and 4), and several equilibria can simultaneously contribute to the formation of a given adduct. Therefore, to avoid any misinterpretation of the selectivity patterns due to an erroneous consideration of the basicity of receptor and substrate, it is convenient to define for each pH value an effective stability constant as the quotient between the overall amounts of complexed dopamine and those of non-complexed dopamine and ligand.¹³

$$\log K_{\text{eff}} = \sum[(H_i\text{Do})(H_j\text{L})] / \{ \sum[H_j\text{L}] \sum[H_i\text{Do}] \}$$

A plot of the logarithms of the stability constants vs pH confirms that the L_3 adducts prevail over those of L_1 and L_2 and permits one to establish selectivity ratios at any pH value by dividing the values of the effective constants (Figure 1). It is interesting to note that the profile of the $\log K_{\text{eff}}$ vs pH curve for L_3 showing an increase of the constant with the pH, is opposite those of L_1 and L_2 .

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Chart 1



It is also remarkable that, particularly for L_3 , there is not a direct correlation between the charge of the receptor and the strength of the interaction. This might be an indication of the importance that forces other than charge-charge or charge-dipole interaction may have in these systems. To achieve a maximum interaction, a certain degree of conformational freedom is needed in order to rightly arrange the functionalities of receptor and substrate, and thereby a high protonation degree of the receptor might afford too much rigidity. This would suggest the participation of the additional arm of cryptand L_3 in the recognition of the neurotransmitter. Chart 1 depicts possible coordination modes for the prevailing species at physiological pH.

To confirm the high stability obtained for the L_3 -dopamine adducts, we have checked the pH-metric stability constants by an alternative UV-vis method. UV spectra of solutions containing a constant amount of L_3 and variable amounts of dopamine at pH 7 show an increase in absorptivity and eventual saturation of an absorption band centered at 214 nm. Fitting of the absorbance vs $[\text{Do}]/[\text{L}_3]$ plots confirmed the 1:1 Do: L_3 stoichiometry and allowed for the estimation of an effective stability constant of ca. 5 logarithmic units in agreement with the pH-metric studies.

With the purpose of getting additional insight into dopamine complexation, we have performed a NMR study on the formation of dopamine adducts of receptors L_2 and L_3 . ^{13}C NMR spectra of dopamine samples with a 3-fold excess of receptors L_2 and L_3 at pH values 6.2 and 7.5 (Table 3; for the labeling of the atoms see drawing of ligands 1-3 and Scheme 2) show, as the most noticeable features, downfield shifts for the signals of carbon atoms $\text{C}3'$ and $\text{C}4'$ of the catechol group upon interaction with L_2 and L_3 . The shifts are particularly significant at pH 7.5 and are greater for $\text{C}3$ than for $\text{C}4$, which

Table 3. Chemical Shifts ($\Delta\delta$, ppm) Induced in Dopamine (Do) ^{13}C NMR Spectra (D_2O) at pH 6.2 and pH 7.5 by Complexation with $2[\text{L}_2]$ and $3[\text{L}_3]$ [Do:L Molar Ratio, 1:3]

dopamine	Do- L_2 pH 6.2	Do- L_2 pH 7.5	Do- L_3 pH 6.2	Do- L_3 pH 7.5
$\text{C}\alpha'$	+0.25	+0.10	+0.08	+0.28
$\text{C}\beta'$	+0.12	+0.08	+0.06	+0.26
$\text{C}1'$	0.0	-0.01	-0.10	-0.02
$\text{C}2'$	+0.06	+0.03	-0.09	-0.09
$\text{C}3'$	+0.20	+0.43	+0.14	+0.65
$\text{C}4'$	+0.19	+0.29	+0.13	+0.58
$\text{C}5'$	+0.07	+0.06	-0.07	+0.13
$\text{C}6'$	-0.05	-0.08	-0.12	-0.14

Table 4. Chemical Shifts ($\Delta\delta$, ppm) Induced in $2[\text{L}_2]$ and $3[\text{L}_3]$ ^{13}C NMR Spectra (D_2O) at pH 6.2 and pH 7.5 by Complexation with Dopamine (L:Do Molar Ratio, 1:3)

ligand	L_2 -Do pH 6.2	L_2 -Do pH 7.5	L_3 -Do pH 6.2	L_3 -Do pH 7.5
$\text{C}3,5$	+0.23	1.35	+0.30	-1.01
$\text{C}4$	-0.15	+0.56	-0.20	+0.83
$\text{C}6$	+0.08	-0.21	-0.01	-0.01
$\text{C}\alpha$	-0.07	+0.09	+0.17	-0.11
$\text{C}\beta$	+0.11	-0.55	+0.04	-0.38

may suggest a partial participation of the HDo form in the complexation process.

^{13}C NMR spectra recorded in a 3-fold excess of dopamine with respect to L_2 and L_3 allowed observation of the changes in the receptor signals brought about by dopamine complexation (Table 4). At pH 6.2, the first aspect to be noted is that while the resonance corresponding to carbon atoms $\text{C}3,5$ appears for the free ligands as a broad signal almost embedded in the baseline of the spectra, it sharpens considerably upon addition of dopamine, yielding a broad singlet centered at 141.2 ppm. Moreover, in the case of L_3 , $\text{C}6$ presents in the ^{13}C NMR spectrum a signal of anomalous low intensity that becomes notably more intense with the addition of dopamine. All these spectral features provide evidence for the formation of the adducts. In the free receptors, at this pH there would be an interaction between the sp^2 pyrazole nitrogens and the protonated neighboring amine groups that slow the prototropic equilibria of the ring. The formation of the dopamine adducts alters the prototropic exchange rate that becomes remarkably faster. Also, the slight deshielding of the ^{13}C NMR signals of carbons $\text{C}3,5$ and the shielding of $\text{C}4$ denote an increase of the negative charge in the pyrazole ring and agree with the preceding observations (Table 4).

At pH 7.5, the NMR behavior of the pyrazole rings is completely different. Both in the free coronand L_2 and in the free cryptand L_3 , the signal of two magnetically equivalent carbons $\text{C}3,5$ are more deshielded and appear as a sharp singlet indicating the absence of interaction with protonated neighboring amine groups. Interaction with dopamine yields shielding and deshielding effects of opposite sign than at pH 6.6, suggesting an increment of positive charge in the pyrazole ring. A plausible explanation for this observation would be a direct interaction of the OH groups of the catechol moiety and the pyrazolic nitrogens. Such an interaction would also be the responsible for the largest shifts of $\text{C}3'$ and $\text{C}4'$ of dopamine observed at this pH (Table 3). The significant shielding of the signals of carbons $\text{C}\beta$ of both receptors (L_2 , -0.55 ppm; L_3 , -0.38 ppm (Table 4)) that occurs at this pH can be ascribed to the simultaneous interaction of dopamine with protonated amine groups close to pyrazole.

As a last aspect in this study, we have modeled the interaction of dopamine with the three receptors with the program MACRO-

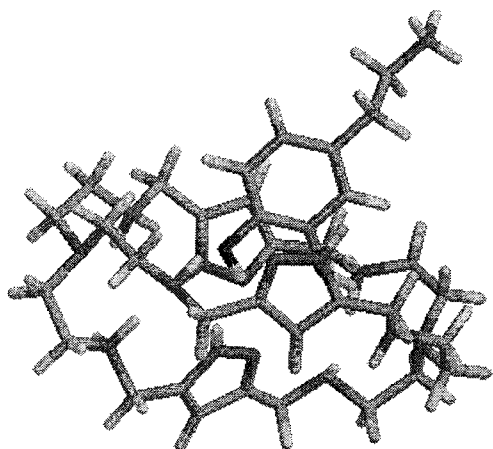
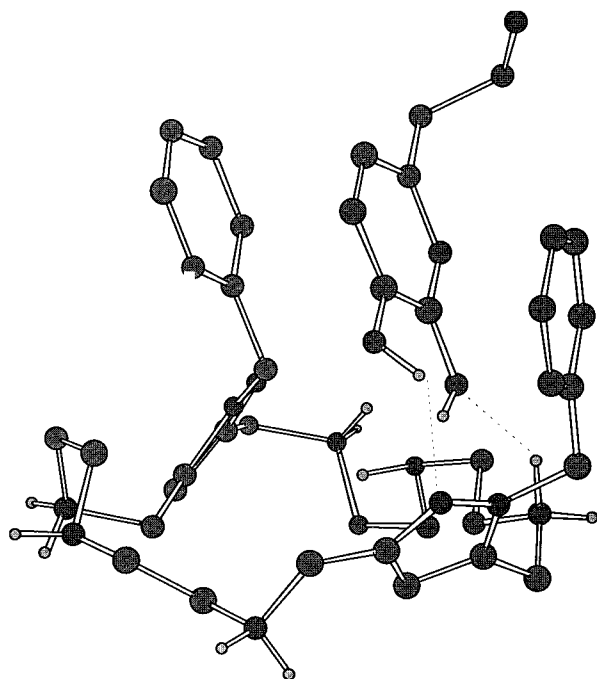


Figure 2. (A, top) Alternative minimum energy conformers for the adduct $H_6L_1(Do)^{5+}$ ($H_4L_1^{4+} \cdot H_2Do^+$) showing stacking contributions. (B, bottom) Minimum energy conformers for the adduct $H_5L_3(Do)^{4+}$ ($H_3L_2 \cdot H_2Do^{4+}$).

MODEL 7.0 using AMBER* as a force field.^{14a} The hydration energy has been considered with the GB/SA^{14b} model incorporated in the MACROMODEL package.

First, a point has to be raised concerning the importance that entropy has in these kind of interactions. Thereby, although the results obtained by models may be corroborative, they can hardly be conclusive.

The family of minimum energy conformers of the adduct $[H_6L_1Do]^{5+}$ ($H_4L_1^{4+} \cdot H_2Do^+$) implies structures in which the hydroxyl groups of the catechol fragment are hydrogen bonded to all four protonated amine groups of the receptor close to the pyrazole moiety. Also in several minima of close energy, contributions from stacking interactions between the benzyl groups of the receptor and the dopamine ring can be found (Figure 2A). The minimum energy conformers for $[H_6L_2Do]^{5+}$

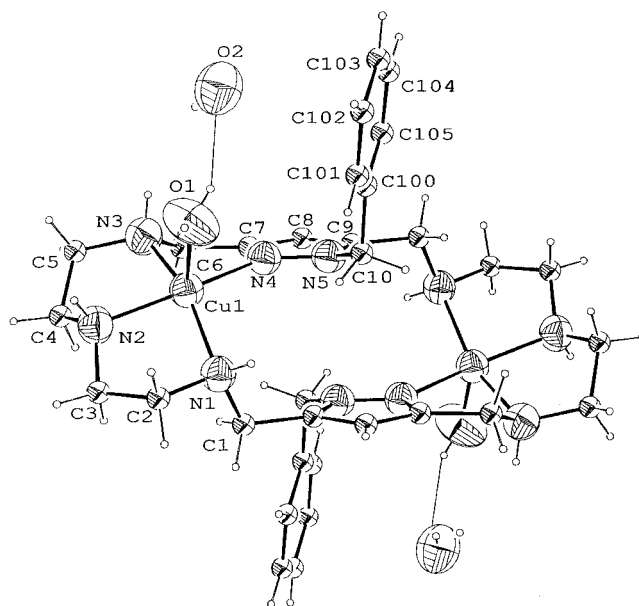


Figure 3. ORTEP drawing for $[Cu_2(L_1)(H_2O)_2]^{4+}$. Thermal ellipsoids are plotted at the 50% probability level.

show also that formation of hydrogen bonds between the tetraprotonated receptor and the phenolic groups of the catechol fragment. These data afford some indications that the larger interaction observed for dopamine- L_1 in relation to dopamine- L_2 may be due to additional stacking interactions.

The conformational analysis of the cage L_3 is particularly difficult for the many prototropic tautomers that can be present. $[H_5L_3Do]^{4+}$ ($H_3L_3^{3+} \cdot H_2Do^+$) presents a structure similar to those found for L_1 and L_2 in which different possibilities of formation of hydrogen networks between receptor and substrate are open (Figure 2B). Finally, for $H_6L_3(Do)^{5+}$ ($H_4L_3^{4+} \cdot H_2Do^+$) the minimum energy structure also reveals the important contribution of hydrogen bonds in the stabilization of the adduct. In some of these structures the participation of the 1H-pyrazole fragment in the hydrogen bond network with the substrate is observed (Figure 2B). None of these structures reveals inclusion of dopamine in the cavity; nevertheless the importance that the entropic term has in these interactions again has to be emphasized, and this parameter could favor an inclusive coordination.

Cu^{2+} Coordination. (a) **Crystal Structure of $[Cu_2(L_1)-(H_2O)_2](ClO_4)_4 \cdot 2H_2O$ (4).** Crystals of the compound $[Cu_2(L_1)-(H_2O)_2](ClO_4)_4 \cdot 2H_2O$ (4) consists of $[Cu_2(L_1)(H_2O)_2]^{4+}$ cations, ClO_4^- anions, and lattice water molecules (Figure 3). Table 5 collects details of the crystal structure refinement and Table 6a selected distances and angles. The coordination geometry around each copper atom is square pyramidal, being the base of the pyramid formed by the three nitrogen atoms of the polyamine bridge [$Cu(1)-N(1) = 2.060(3)$ Å, $Cu(1)-N(2) = 2.003(3)$ Å, $Cu(1)-N(3) = 2.027(4)$ Å, $Cu(1)-N(4) = 2.016(3)$ Å] and the nonbenzylated sp^2 nitrogen of the *N*-benzylpyrazole spacer. The axial position is occupied by a water molecule ($Cu(1)-O(1) = 2.170(4)$ Å). The axial water molecule in each coordination site is hydrogen bonded to another water molecule ($O(1)-O(2) = 2.84(1)$ Å). The copper atom almost resides in the equatorial plane of the pyramid with a mean elevation over the plane defined by the coordinated nitrogen atoms of 0.325(2) Å. The $Cu \cdots Cu$ distance is 6.788(2) Å. The molecule presents an inversion center, and therefore the water molecules are pointing toward opposite sides of the mean plane defined by the macrocyclic cavity. According to those previously observed using

(14) (a) Mohamadi, F.; Richards, N. G. J.; Guida, W. C.; Liskamp, R.; Lipton, M.; Canfield, C.; Chang, G.; Hendrickson, T.; Still, W. C. *J. Comput. Chem.* **1990**, *11*, 440. (b) Qiu, D.; Shenkin, P. S.; Hollinger, F. P.; Still, W. C. *J. Phys. Chem.* **1997**, *101*, 3005. (c) Stewart, J. J. P. *J. Comput. Chem.* **1989**, *10*, 209.

Table 5. Crystallographic Data for **4** and **6**

	4	6
empirical formula	C ₁₆ H ₂₇ C ₁₂ Cu N ₅ O ₁₀	C ₅₄ H ₁₀₂ C ₁₆ Cu ₄ N ₂₈ O ₂₈
fw	583.87	2058.50
space group	<i>P</i> 2 ₁ / <i>a</i>	<i>C</i> 2/ <i>c</i>
<i>V</i> , Å ³	2310(2)	8098(5)
<i>a</i> , Å	9.381(5)	44.667(5)
<i>b</i> , Å	19.418(5)	9.124(5)
<i>c</i> , Å	12.804(5)	20.050(5)
β , deg	97.931(5)	97.687(5)
<i>T</i> , K	293(2)	293(2)
<i>D</i> _{calcd.} , g/cm ³	1.679	1.688
μ (Mo K α), mm ⁻¹	1.24	1.33
2 θ range, deg	1–25	1–25
total no. of data	4142	6759
no. of unique data	3902	3411
no. of obsd data ^a	3132	1623
no. of params	326	325
<i>R</i> ^b	0.0663	0.0684
<i>R</i> _w ^c	0.1594	0.1776

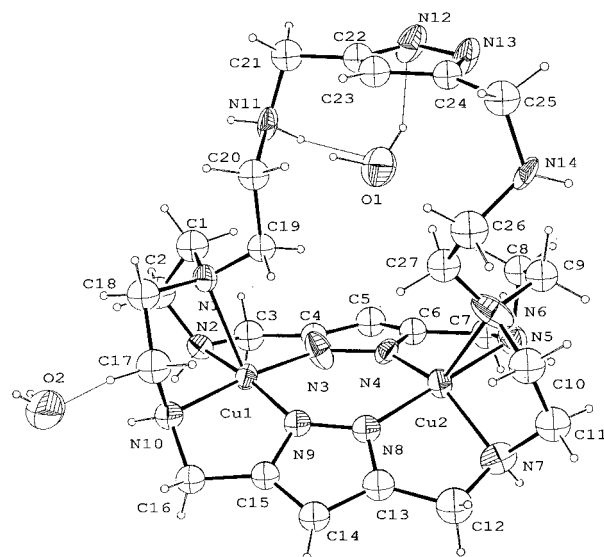
^a Observed criterion: $F_o > 4\sigma F_o$. ^b $R_1 = \sum ||F_o| - |F_c|| / \sum |F_o|$.
^c $R_w = [\sum w(F_o^2 - F_c^2)^2 / \sum w(F_o^2)]^{1/2}$. For **4**, $w = 1/[\sigma^2(F_o^2) + (0.10P)^2]$,
 $P_{max} = (\text{Max}(F_o)^2 + 2(F_c)^2)/3$; for **6**, $w = 1/[\sigma^2(F_o^2) + (0.14P)^2]$,
 $P_{max} = (\text{Max}(F_o)^2 + 2(F_c)^2)/3$.

Table 6. Selected Bond Distances (Å) and Bond Angles (deg)

(a) For Compound 4			
Cu(1)–N(2)	2.003(3)	N(2)–Cu(1)–N(3)	84.1(1)
Cu(1)–N(4)	2.016(3)	N(4)–Cu(1)–N(3)	80.0(1)
Cu(1)–N(3)	2.027(4)	N(2)–Cu(1)–N(1)	85.6(1)
Cu(1)–N(1)	2.060(3)	N(4)–Cu(1)–N(1)	104.5(1)
Cu(1)–O(1)	2.170(4)	N(2)–Cu(1)–O(1)	98.2(2)
Cu–Cu	6.788(2)	N(4)–Cu(1)–O(1)	98.2(2)
		N(3)–Cu(1)–O(1)	101.6(2)
		N(1)–Cu(1)–O(1)	99.2(2)
(b) For Compound 6			
Cu(1)–N(3)	1.90(1)	N(3)–Cu(1)–N(9)	94.7(6)
Cu(1)–N(9)	1.93(2)	N(9)–Cu(1)–N(10)	79.6(6)
Cu(1)–N(10)	2.05(1)	N(3)–Cu(1)–N(2)	81.4(6)
Cu(1)–N(2)	2.11(1)	N(10)–Cu(1)–N(2)	103.7(5)
Cu(1)–N(1)	2.29(2)	N(3)–Cu(1)–N(1)	102.9(6)
Cu(2)–N(8)	1.92(1)	N(9)–Cu(1)–N(1)	111.3(5)
Cu(2)–N(4)	1.92(1)	N(10)–Cu(1)–N(1)	81.4(5)
Cu(2)–N(5)	2.10(1)	N(2)–Cu(1)–N(1)	82.0(5)
Cu(2)–N(7)	2.12(2)	N(8)–Cu(2)–N(4)	94.6(5)
Cu(2)–N(6)	2.40(1)	N(4)–Cu(2)–N(5)	80.2(6)
Cu(1)–Cu(2)	3.960(3)	N(8)–Cu(2)–N(7)	81.4(6)
		N(5)–Cu(2)–N(7)	101.4(7)
		N(8)–Cu(2)–N(6)	105.6(5)
		N(4)–Cu(2)–N(6)	123.7(6)
		N(5)–Cu(2)–N(6)	80.1(4)
		N(7)–Cu(2)–N(6)	81.7(5)

NMR techniques,¹¹ the benzyl groups are disposed trans to each other, pointing also outward from the macrocyclic cavity.

(b) Crystal Structure of [Cu₂(H–L₃)](ClO₄)₃·2H₂O (6**).** The crystal structure of **6** consists of [Cu₂(H–L₃)]³⁺ cations, ClO₄⁻ anions, and water molecules. Table 5 presents details of the crystal structure and of the refinement procedure. The coordination geometry around each Cu²⁺ is square pyramidal, the base of the pyramid being formed by two secondary nitrogens of the bridge and two nitrogen atoms of two different pyrazolate moieties which act as exobidentate ligands (Figure 4). The axial positions are occupied by the bridgehead nitrogen atoms, being the distortion more pronounced in one of the two sites [Cu(1)–N(1) = 2.29(2) Å and Cu(2)–N(6) = 2.40(1) Å, Table 6b]. The Cu²⁺–L distances involving the sp² pyrazolate nitrogen atoms [Cu(1)–N(3) = 1.90(1) Å, Cu(1)–N(9) = 1.93(2) Å, Cu(2)–N(4) = 1.92(1) Å, and Cu(2)–N(8) = 1.92(1) Å] are much shorter than those of the secondary nitrogen

**Figure 4.** ORTEP drawing of the complex [Cu₂(HL₃)](ClO₄)₃·2H₂O. Thermal ellipsoids are plotted at the 50% probability level. ClO₄⁻ anions are removed for clarity.

atoms [Cu(1)–N(2) = 2.11(1) Å, Cu(1)–N(10) = 2.05(1) Å, Cu(2)–N(5) = 2.10(1) Å, and Cu(2)–N(7) = 2.12(2) Å]. The distance Cu(1)–Cu(2) is 3.960(3) Å. Similar coordination features were observed in the crystal structure of the binuclear complex [Cu₂(H–L₂)](ClO₄)₂,¹² and in several crystal structures of other pyrazole containing ligands.^{6,15} Interestingly enough, we have observed that Cu²⁺ produces the ready deprotonation of the two pyrazole fragments involved in the coordination, without requiring any addition of base (vide infra). However, the pyrazole in the noncoordinating bridge of L₃ (see Figure 4) does not deprotonate, pointing at one side of the macrocyclic cavity. One of the nitrogen atoms of the aliphatic chains is protonated and hydrogen bonded to a water molecule (N(11)–O(1) = 2.79(2) Å) that is placed at one side of the macrocyclic cavity and further connected to the sp² nitrogen of the pyrazole group through an additional hydrogen bond (N(12)–O(1) = 3.12(2) Å). The distances of the water molecule to the nonprotonated nitrogen atom of the bridge and to the other pyrazole nitrogen are much longer (5.38 and 3.79 Å, respectively). The other water molecule, placed completely outside of the cavity, is hydrogen bonded to N(10) (N(10)–O(2) = 2.98(2) Å).

The coordination arrangement keeps both Cu²⁺ metal ions close to one face of the cage (the elevation of the Cu atoms over the mean plane defined by the nitrogen donors of the base of the square pyramid are 0.142(7) and 0.272(7) Å for Cu(1) and Cu(2), respectively) leaving enough free room within the cavity for allowing the encapsulation of further substrates as exogenous ligands.

Electromotive Force Measurements. The stability constants for the interaction with Cu²⁺ of receptors L₁–L₃ determined at 298.1 K in 0.15 mol dm⁻³ are presented in Table 7. In all three systems formation of mono- and binuclear complexes is observed. In the system Cu²⁺–L₃ also a trinuclear species of [Cu₃(H–L₃)]³⁺ stoichiometry is even detected.

The thermodynamic data in Table 7 and the distribution diagrams in Figure 5 provide important clues about the coordination characteristics of these pyrazole containing polyaza macrocycles.

(15) (a) Behle, L.; Neuburger, M.; Zehnder, M.; Kaden, T. A. *Helv. Chim. Acta* **1995**, *78*, 693. (b) Weller, H.; Kaden, T. A.; Hopfgartner, G. *Polyhedron* **1998**, *17*, 4543.

Table 7. Stability Constants for the Formation of Cu^{2+} Complexes of Receptors L_1 , L_2 , and L_3 Determined in 0.15 mol dm^{-3} NaCl at 298.1 K

entry	reaction ^a	L_1	L_2	L_3
1	$\text{Cu} + \text{L} + 2\text{H} \rightleftharpoons \text{CuH}_2\text{L}$	27.9(1) ^b	35.05(2)	35.42(8)
2	$\text{Cu} + \text{L} + \text{H} \rightleftharpoons \text{CuHL}$	22.13(8)	30.81(2)	29.85(7)
3	$\text{Cu} + \text{L} \rightleftharpoons \text{CuL}$	13.68(6)	23.47(3)	21.76(6)
4	$\text{Cu} + \text{L} + \text{H}_2\text{O} \rightleftharpoons \text{CuH}_{-1}\text{L} + \text{H}$			13.05(3)
5	$2\text{Cu} + \text{L} + \text{H} \rightleftharpoons \text{Cu}_2\text{HL}$	29.37(5)		39.04(7)
6	$2\text{Cu} + \text{L} \rightleftharpoons \text{Cu}_2\text{L}$	26.27(3)	34.45(3)	36.72(3)
7	$2\text{Cu} + \text{L} \rightleftharpoons \text{Cu}_2\text{H}_{-1}\text{L} + \text{H}$	16.85(4)		31.31(5)
8	$2\text{Cu} + \text{L} \rightleftharpoons \text{Cu}_2\text{H}_{-2}\text{L} + 2\text{H}$	7.06(3)	26.56(3)	24.13(5)
9	$3\text{Cu} + \text{L} \rightleftharpoons \text{Cu}_3\text{H}_{-3}\text{L} + 3\text{H}$			25.78(4)
10	$\text{CuHL} + \text{H} \rightleftharpoons \text{CuH}_2\text{L}$	5.8	4.2	6.6
11	$\text{CuL} + \text{H} \rightleftharpoons \text{CuHL}$	8.5	7.3	7.1
12	$\text{CuL} \rightleftharpoons \text{CuH}_{-1}\text{L} + \text{H}$			-8.7
13	$\text{CuL} + \text{Cu} \rightleftharpoons \text{Cu}_2\text{L}$	12.6	11.0	15.0
14	$\text{Cu}_2\text{L} + \text{H} \rightleftharpoons \text{Cu}_2\text{HL}$	3.1		2.3
15	$\text{Cu}_2\text{L} \rightleftharpoons \text{Cu}_2\text{H}_{-1}\text{L} + \text{H}$	-9.4		-5.4
16	$\text{Cu}_2\text{H}_{-1}\text{L} \rightleftharpoons \text{Cu}_2\text{H}_{-2}\text{L} + \text{H}$	-9.8		-7.2
17	$\text{Cu}_2\text{L} \rightleftharpoons \text{Cu}_2\text{H}_{-2}\text{L} + 2\text{H}$	-19.2	-7.9	-12.6

^a Charges omitted for clarity. ^b Values in parentheses are standard deviations in the last significant figure.

The speciation shows for the three systems the formation of the mononuclear species $[\text{CuH}_2\text{L}]^{4+}$, $[\text{CuHL}]^{3+}$, and $[\text{CuL}]^{2+}$; for the system $\text{Cu}^{2+}-\text{L}_3$ also the species $[\text{CuH}_{-1}\text{L}_3]^+$ was observed. The most important difference between the three systems is the much lower constants obtained for the benzylated receptor L_1 with respect to L_2 and L_3 ; the formation constant for the $[\text{CuL}_1]^{2+}$ species is 13.68 logarithmic units, while those for the corresponding species of L_2 and L_3 are over 20 logarithmic units (entry 3 in Table 7). This situation can be ascribed to the fact that the benzylated receptor is unable to deprotonate and yield the stronger field pyrazolate ligand. UV-vis spectra are not very clear in identifying this situation since formation of the binuclear complexes is even detected for low $\text{Cu}^{2+}:\text{L}$ metal ratios (see Figure 5). On the other hand, the large protonation constants of the $[\text{CuL}]^{2+}$ complexes (entry 11) denotes that these processes are occurring in non-coordinated nitrogen atoms of the three receptors.

The thermodynamic data for the binuclear $\text{Cu}^{2+}-\text{L}_1$ complexes agree well with the fundamental coordination characteristics evidenced by the crystal structure (vide supra). Indeed in this case, both coordination sites seem to act almost independently. The stepwise addition of the second Cu^{2+} ion displays a constant almost as high as that of the addition of the first metal ion (entries 13 and 3 in the table, respectively). Furthermore, the hydrolysis of the water molecules in the binuclear complex $[\text{Cu}_2\text{L}_1]^{4+}$ to give the hydroxo species present pK_a values close between them (9.4 and 9.8, entries 15 and 16) and high enough to discard the formation of hydroxo bridges between the copper(II) metal ions.

In view of these thermodynamic parameters and of the X-ray study previously discussed, the structures in Chart 2 can be proposed for the three main binuclear species in solution for a molar $\text{Cu}:\text{L}_1$ ratio 2:1. $[\text{Cu}_2(\text{L}_1)(\text{H}_2\text{O})_2]^{4+}$ would be coincident with the X-ray structure, while all the data strongly suggest that neither in $[\text{Cu}_2(\text{L}_1)(\text{H}_2\text{O})(\text{OH})]^{3+}$ nor in $[\text{Cu}_2(\text{L}_1)(\text{OH})_2]^{2+}$ is there formation of hydroxo bridges between the metal centers.

The distribution diagram of the system $\text{Cu}^{2+}-\text{L}_2$ in molar ratio 2:1 shows the formation of $[\text{Cu}_2\text{H}_{-2}\text{L}_2]^{2+}$ as the unique species in solution above pH 4.5. The formation of this species implies the deprotonation of the two pyrazole spacers that according to the X-ray structure¹² will be acting as exobidentate bridging ligands between both metal centers. As shown by the constants (entries 8 and 17 in Table 7), these deprotonation

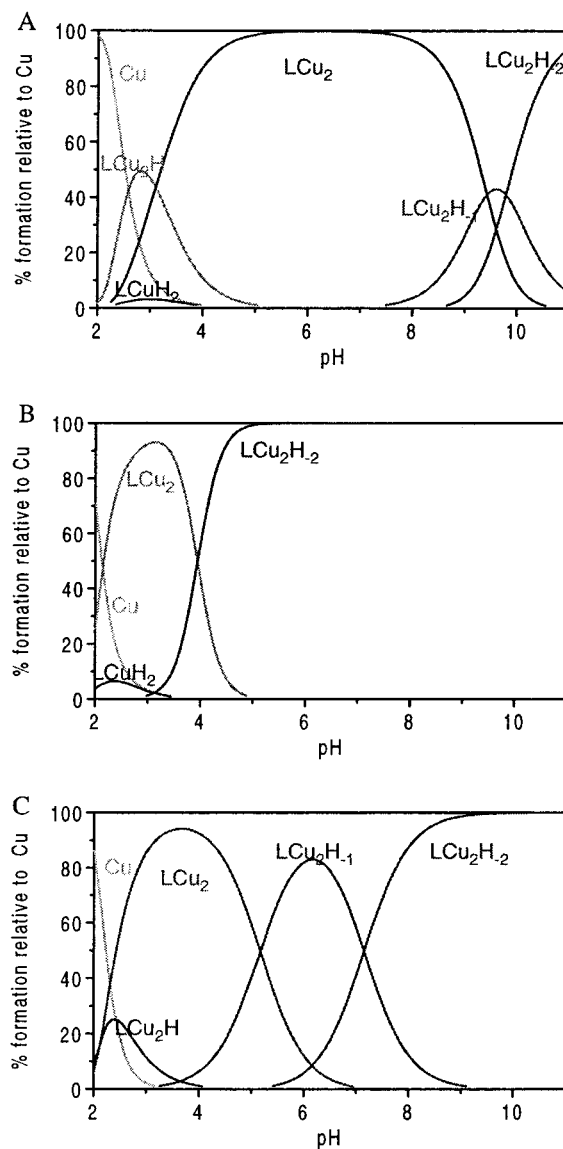
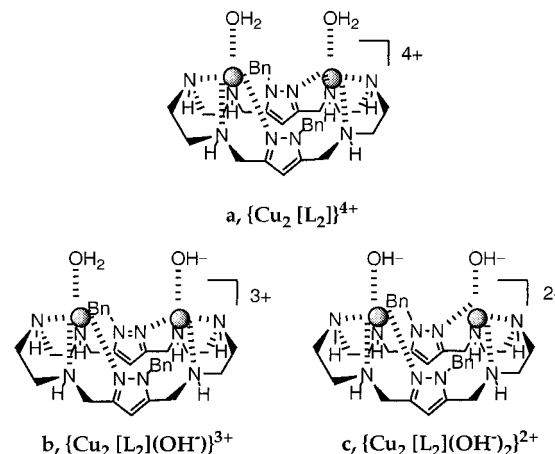


Figure 5. Distribution diagram for the systems (A) $\text{Cu}^{2+}-\text{L}_1$, (B) $\text{Cu}^{2+}-\text{L}_2$, and (C) $\text{Cu}^{2+}-\text{L}_3$, calculated for molar ratios 2:1 $\text{Cu}^{2+}:\text{L}$ ($[\text{Cu}^{2+}] = 2 \times 10^{-3} \text{ mol dm}^{-3}$).

Chart 2



processes occur cooperatively, and as a consequence we could not calculate a constant for the first one. The binuclear $[\text{Cu}_2\text{H}_{-2}\text{L}_2]^{2+}$ species are the main ones in solution even for low metal:ligand molar ratios (see Figure 5). Therefore, Cu^{2+}

coordination promotes a dramatic increase of the acidity of the pyrazole fragment than when free does not deprotonate up to pH 12.¹¹ Also the cooperative behavior of both coordination sites in L_2 should be noted.

Deprotonation of the pyrazole units in aqueous solution induced by coordination to metal is a characteristic feature of this heterocycle. Kaden et al.¹⁵ have evidenced this behavior in the Cu^{2+} complexes of a bis(macrocylic) receptor containing two 1,4,7-triazacyclononane ([9]aneN₃) unit connected to the 3,5 positions of a pyrazole spacer through methylenic groups (L_5).¹⁵ Deprotonation of the 1H-pyrazole group in the $[\text{Cu}_2\text{L}_5]^{4+}$ complex presented a pK_a value of 3.67 that falls in the range of the values here reported (Table 7). Similar conclusions were derived for these authors for a related bis(macrocylic) ligand with 1,4,7-azadithianonane ([9]aneNS₂) units.^{15b}

In a previous paper some of us¹² reported on the crystal structure of complex 4. In such a structure both pyrazole units were deprotonated, behaving as bis(bidentate) anions. The crystals were grown from an ethanolic solution of the ligand to which 2 mol of NaOH/(mol of ligand) were added. Here, we have arrived at the same complex without the addition of base following a re-crystallization in water (see Experimental Section). This experiment gives further support to the ready deprotonation that the pyrazole groups experience in aqueous solution induced by the presence of metal ions.

The stability data for the mono- and binuclear Cu^{2+} complexes of the cryptand L_3 shows similarities with those discussed for the system $\text{Cu}^{2+}-\text{L}_2$. The greatest differences between the two systems are the larger stability of the $[\text{Cu}_2\text{L}_3]^{4+}$ complex with respect to $[\text{Cu}_2\text{L}_2]^{4+}$ and the reduced acidity of the pyrazole moieties in the binuclear Cu^{2+} complexes of L_3 ($\text{pK}_{a1} = 5.4$, $\text{pK}_{a2} = 7.2$, entries 15 and 16 in Table 7; see Figure 5). While, for L_2 , the two pyrazole moieties are already fully deprotonated at pH 5, for L_3 such a situation seems to be only achieved at pH 8.5. Probably, the presence of the additional bridge in the cryptand structure can contribute to make more rigid the molecule, disrupting somewhat the cooperativity observed in L_2 for these processes.

A final point that deserves some comment is the formation of a trinuclear $[\text{Cu}_3\text{L}_3\text{H}_{-3}]^{3+}$ complex (Table 7, entry 9) which is the species majoritary in solution for molar ratios 3:1 $\text{Cu}^{2+}:\text{L}_3$ over pH 5. The formation of this complex should imply the deprotonation of all three pyrazole fragments of the cage at a relatively low pH value. Currently we are performing experiments to try to unravel the molecular structure of this complex.

Mixed Complexes $\text{Cu}^{2+}-\text{L}$ -Dopamine. The formation of mixed complexes $\text{Cu}^{2+}-\text{L}$ -dopamine has been followed by potentiometric measurements at 298.1 K in 0.15 mol dm^{-3} NaCl using molar ratios $\text{Cu}^{2+}-\text{L}$ -dopamine ranging from 2:1:1 to 2:1:3. The model species and stability constants that best fit the experimental data correspond to the formation of ternary complexes with stoichiometries $\text{Cu}^{2+}-\text{L}$ -dopamine 2:1:1. The stability constants for such systems are included in Table 8. The distribution diagrams for the three systems calculated for molar ratios 1:2:1 $\text{Cu}^{2+}:\text{L}:\text{dopamine}$ are reported in Figure 6.

These ternary complexes are for L_2 only the main species in solution above pH 8.5 in correspondence with the formation of the species $[\text{Cu}_2(\text{H}_{-2}\text{L}_2)(\text{DoH})]^{2+}$ and $[\text{Cu}_2(\text{H}_{-2}\text{L}_2)(\text{Do})]^+$. However, in the case of L_1 the mixed complexes are always the main ones in solution prevailing throughout the entire pH range studied (pH 3–11). This behavior can find its explanation in the different nature of the binuclear complexes formed. As discussed previously, for L_2 , the coordination sites of the Cu^{2+}

Table 8. Stability Constants for the Formation of $\text{Cu}^{2+}-\text{L}-\text{Do}$ Complexes of Receptors L_1 , L_2 , and L_3 (Do = Dopamine) Determined in 0.15 mol dm^{-3} NaCl at 298.1 K

entry	reaction ^a	L_1	L_2	L_3
1	$2\text{Cu} + \text{L} + \text{Do} + 2\text{H} \rightleftharpoons \text{Cu}_2\text{H}_2\text{LDo}$	49.01(4) ^b		59.6(1)
2	$2\text{Cu} + \text{L} + \text{Do} + \text{H} \rightleftharpoons \text{Cu}_2\text{HLDo}$	42.15(6)	51.7(2)	54.2(1)
3	$2\text{Cu} + \text{L} + \text{Do} \rightleftharpoons \text{Cu}_2\text{LDo}$			47.0(1)
4	$2\text{Cu} + \text{L} + \text{Do} \rightleftharpoons \text{Cu}_2\text{H}_{-1}\text{LDo} + \text{H}$	26.20(4)	40.56(3)	38.5(1)
5	$2\text{Cu} + \text{L} + \text{Do} \rightleftharpoons \text{Cu}_2\text{H}_{-2}\text{LDo} + 2\text{H}$	18.40(4)	30.36(4)	
6	$2\text{Cu} + \text{L} + \text{Do} \rightleftharpoons \text{Cu}_2\text{H}_{-3}\text{LDo} + 3\text{H}$	8.19(4)		
8	$\text{Cu}_2\text{L} + \text{H}_2\text{Do} \rightleftharpoons \text{Cu}_2\text{H}_2\text{LDo}$	3.5		3.6
9	$\text{Cu}_2\text{H}_{-1}\text{L} + \text{H}_2\text{Do} \rightleftharpoons \text{Cu}_2\text{HLDo}$	6.2		3.9
10	$\text{Cu}_2\text{L} + \text{HDo} \rightleftharpoons \text{CuHLDo}$	5.6		
11	$\text{Cu}_2\text{H}_{-2}\text{L} + \text{HDo} \rightleftharpoons \text{CuH}_{-1}\text{LDo}$	8.8	3.7	4.0
12	$\text{Cu}_2\text{H}_{-3}\text{L} + \text{Do} \rightleftharpoons \text{CuH}_{-3}\text{LDo}$	11.3	3.8	

^a Charges omitted for clarity. ^b Values in parentheses are standard deviations in the last significant figure.

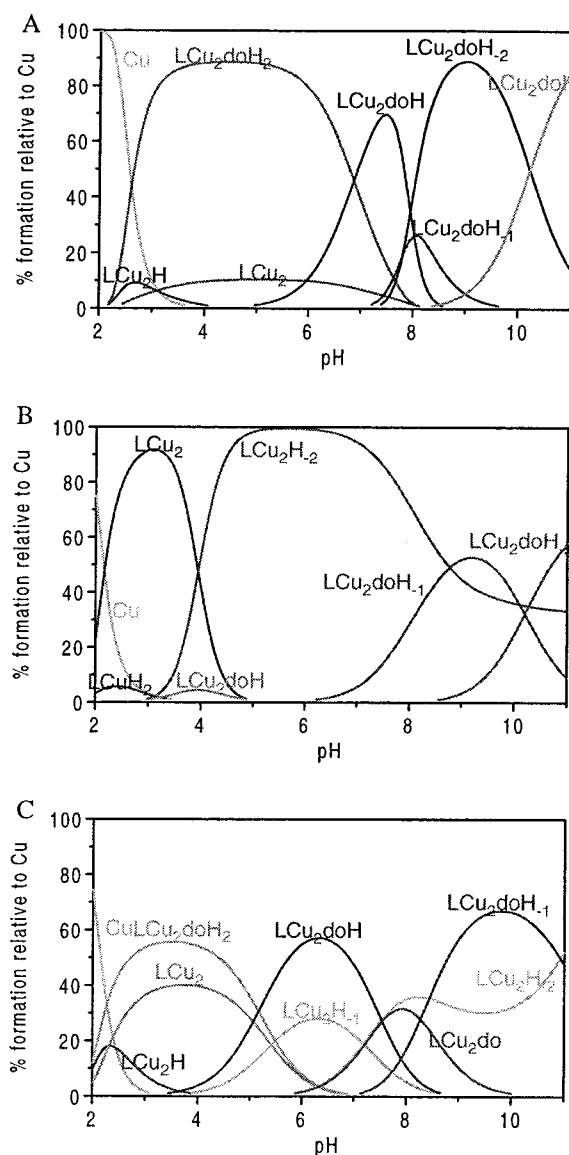


Figure 6. Distribution diagram for the systems (A) $\text{Cu}^{2+}-\text{L}_1-\text{Do}$, (B) $\text{Cu}^{2+}-\text{L}_2-\text{Do}$, and (C) $\text{Cu}^{2+}-\text{L}_3-\text{Do}$, calculated for molar ratios 2:1:1 $\text{Cu}^{2+}:\text{L}:\text{Do}$ ($[\text{Cu}^{2+}] = 2 \times 10^{-3}$ mol dm^{-3}).

ions were made up by five nitrogen donors, occupying the central nitrogens of the diethylenetriamine subunits in the axial positions of the square pyramid. Formation of mixed complexes requires the detachment of at least one of the nitrogen donors (most likely the one in the axial position), and this would be

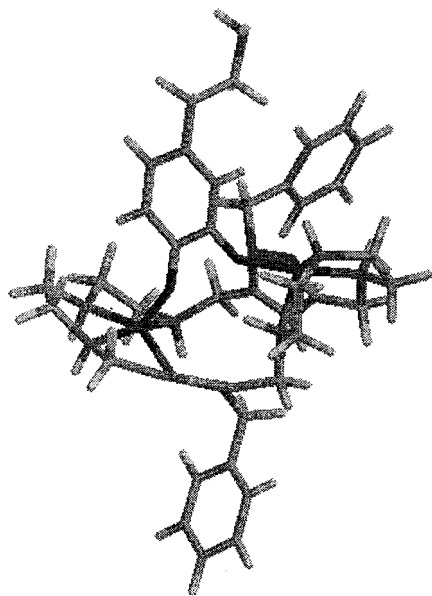


Figure 7. Model for the formation of ternary complexes $\text{Cu}^{2+}\text{-L}_1\text{-Do}$.

only efficient when the hydroxo groups of dopamine deprotonate to bind the Cu^{2+} ions at around pH 8.

For L_1 , however, the axial positions in the first coordination sphere are occupied by labile water molecules, which are more easily replaced by catecholamine favoring the formation of the ternary species. Although it is difficult to advance an hypothesis on structural aspects of the mixed complexes, the stoichiometries found and initial molecular modeling studies performed with the program SPARTAN^{14c} suggest the possibility that dopamine could act as a bridging exogenous ligand between both coordination sites. Cu^{2+} coordination would be facilitating the partial deprotonation of the phenolic groups (Figure 7).

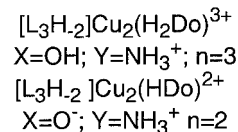
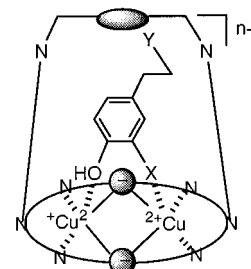
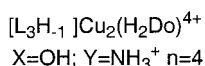
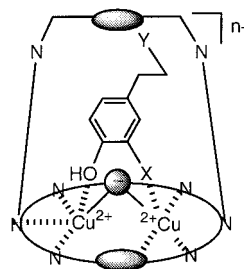
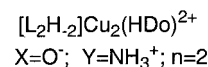
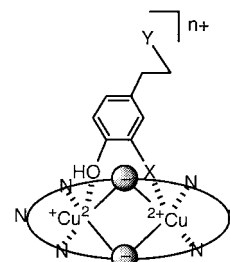
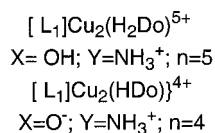
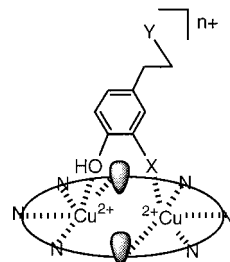
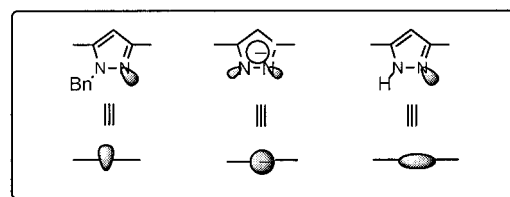
For the system $\text{Cu}^{2+}\text{-L}_3\text{-dopamine}$ formation of mixed complexes is observed throughout all of the pH range investigated. These species display intermediate stabilities between those of the ternary complexes of L_1 and L_2 . Although the coordination arrangement around the Cu^{2+} ions in the binary complexes of L_2 and L_3 is similar, inclusion of dopamine in the void upper part of the cavity might contribute to the larger stability of the L_3 ternary complexes.

Chart 3 shows possible coordination modes for the ternary complexes prevalent around physiological pH.

Conclusions

The results analyzed show that cryptand L_3 is a particularly well-suited receptor for dopamine recognition in aqueous solution. All three receptor form very stable binuclear Cu^{2+} complexes in aqueous solution. Formation of the Cu^{2+} complexes induces the ready deprotonation of the pyrazole units in L_2 and L_3 . Crystal structures of the complex $[\text{Cu}_2(\text{L}_1)(\text{H}_2\text{O})_2](\text{ClO}_4)_4 \cdot 2\text{H}_2\text{O}$ (**4**) denote the participation of the sp^2 nitrogen donors of the nonbenzylated *N*-benzylpyrazole nitrogen donors in the coordination of the Cu^{2+} metal ions which present square pyramidal geometry with a water molecule in axial position (Figure 3). The crystal structure of $[\text{Cu}_2(\text{H-L}_3)](\text{ClO}_4)_3 \cdot 2\text{H}_2\text{O}$ (**6**) shows also square pyramidal coordination geometry around each Cu^{2+} . The axial positions are occupied in this case by the bridgehead nitrogen of the cage. The two deprotonated pyrazole subunits behave as exobidentate ligands. In this structure there is a water molecule strongly retained by hydrogen bonds with

Chart 3



a protonated amine group and the nondeprotonated pyrazole of the bridge that does not participate in the coordination to the metal. Solution studies agree with the fundamental characteristic denoted by the crystal structures and also suggest the possible formation of trinuclear species for L_3 .

Formation of ternary complexes $\text{Cu}^{2+}\text{-dopamine-L}$ is particularly relevant in the case of L_1 where most likely the hydroxyl groups of the catechol would be replacing the axially coordinated water molecules.

The results here presented give interesting perspectives for the design of receptors for neurotransmitters. Currently we are exploring these insights.

Experimental Section

The starting materials were purchased from commercial sources and used without further purification. The solvents were dried using standard techniques. All reactions were monitored by thin-layer chromatography using DC-Alufolien silica gel 60PF₂₅₄ (Merck; layer thickness, 0.2 mm). Compounds were detected with iodine or with phosphomolybdic acid reagent. Melting points were determined in a Reichert-Jung hot-stage microscope and are uncorrected. ¹H and ¹³C NMR spectra were recorded on Varian Unity Inova-400 and Varian Unity-500 spectrometers. The chemical shifts are reported in parts per million from tetramethylsilane but were measured against the solvent signals; dioxane ($\delta = 67.4$ ppm) was used as reference for ¹³C NMR spectra in D₂O. All assignments

During the final stages of the refinement the positional parameters and the anisotropic thermal parameters of the non-hydrogen atoms were refined. The hydrogen atoms were refined with a common thermal parameter. The final conventional agreement factors for **4** and **6** were respectively $R_1 = 0.0663$, $R_{w2} = 0.1594$ and $R_1 = 0.0684$, $R_{w2} = 0.1776$. Atomic scattering factors were taken from the ref 28. Molecular plots were produced by the program ORTEP.²⁹

Molecular Mechanics Calculations. Molecular mechanics calculations were performed with MACROMODEL 7.0 package using AMBER*^{14a} as force-field and GB/SA^{14b} solvation model for water.

Semiempirical calculations of the metal complex structures were performed with PC SPARTAN PRO 1.0.5 using the PM3 parametrization.^{14c}

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(29) Johnson, C. K. *ORTEP*; Report ORNL-3794; Oak Ridge National Laboratory: Oak Ridge, TN, 1971.

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Supporting Information Available: Crystal data and structure refinement details (Tables S1 and S1'), atomic coordinates (Table S2 and S2'), bond lengths and angles (Table S3 and S3'), anisotropic displacement parameters (Tables S4 and S4'), and hydrogen coordinates for **4** and **6** (Tables S5 and S5') (PDF). This material is available free of charge via the Internet at <http://pubs.acs.org>.

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