

# Stereoselective self-assembly of atropoisomeric Pd(II) metallocycles induced by an aromatic guest†

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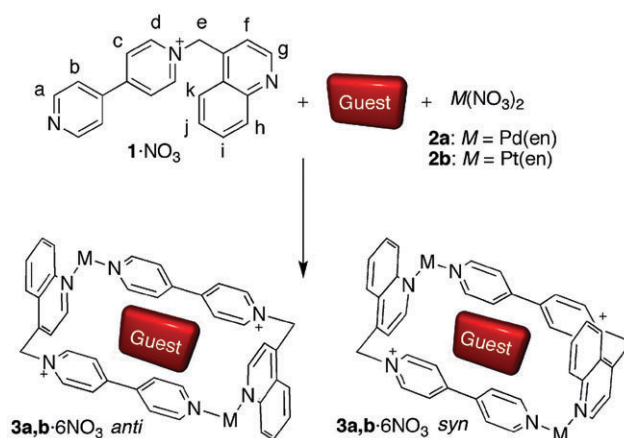
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A new dinuclear Pd(II) metallocycle consisting of a 4,4'-bipyridin-1-ium ligand with a quinoline moiety was self-assembled in aqueous solution with the aid of template molecules; the situation found in solution in which both *syn* and *anti* or only the *anti* atropoisomers are observed strongly relies on the intermolecular host-guest interactions.

The facile obtention of macrocycles with cavities large enough to accommodate substrate molecules in complexation processes is of enormous importance in host-guest chemistry.<sup>1</sup> The structural characteristics of their cavities, including shape, size and flexibility, determinate the usefulness of the receptors towards target substrates. Metal-directed self-assembly has emerged as a powerful tool to obtain molecular hosts of defined shapes and sizes.<sup>2</sup> The design of receptors capable of adapting their structure to bind the guest in a manner reminiscent of Koshland's induced model<sup>3</sup> for substrate binding to enzymes has attracted much attention because such induced fit is the basis of many allosteric systems.<sup>4</sup>

In our previous reports, we have described different strategies and procedures towards the self-assembly of square and rectangular Pd(II) and Pt(II) metallocycles and related structures such as inclusion complexes and catenanes.<sup>5</sup> Here, we use 4-substituted quinoline moieties in the short sides of a molecular rectangle, which can rotate around the CH<sub>2</sub>-N axis giving rise to the existence of two atropoisomers with different cavity characteristics. The advantage of this "flexible" receptor is that it can change its shape in order to maximize the intermolecular interactions with different guests (induced-fit).

Ligand **1**·NO<sub>3</sub> was easily obtained from 4,4'-bipyridine and 4-chloromethylquinoline<sup>6</sup> in moderate yield. The addition of one equivalent of hydroquinone (HQ) to an equimolar 10 mM D<sub>2</sub>O solution of ligand **1**·NO<sub>3</sub> and Pd(en)(NO<sub>3</sub>)<sub>2</sub> results in the appearance of a reddish colour solution. The <sup>1</sup>H and <sup>13</sup>C NMR spectra showed an approximate 1 : 1 mixture of the *syn* and *anti* HQ·**3a**·6NO<sub>3</sub> atropoisomers (Scheme 1), as can be concluded from the two sets of signals and 2D NMR data (Fig. 1). The chemical shifts are compatible with the formation of the coordinative bonds ( $\Delta\delta = 1.82$  for H<sub>h</sub> and  $\Delta\delta = 0.58$  ppm for H<sub>g</sub> from those of **1**·NO<sub>3</sub>). The



Scheme 1

upfield shift of the signal of HQ ( $\Delta\delta = -1.10$  ppm from those of the free HQ, for the aromatic protons of HQ) along with the upfield shift of the  $\beta$  protons of the bipyridine units (H<sub>b,c</sub>) suggests that HQ is inserted into the cavity of metallocycle. Similar results have been obtained with resorcinol, catechol, and phloroglucinol. It is noteworthy that the interconversion of the stereoisomers is slow in the NMR timescale, probably as a result of the energy barrier for the rotation of the quinoline units, in addition to the steric and energetic impediments caused by the hydroquinone guest inside the cavity.

Addition of NaNO<sub>3</sub> results in the slow crystallization of the inclusion complex. X-Ray analysis of these crystals shows the presence of the *anti* stereoisomer exclusively (Fig. 2).<sup>7</sup> Both components lie about a common inversion centre located at

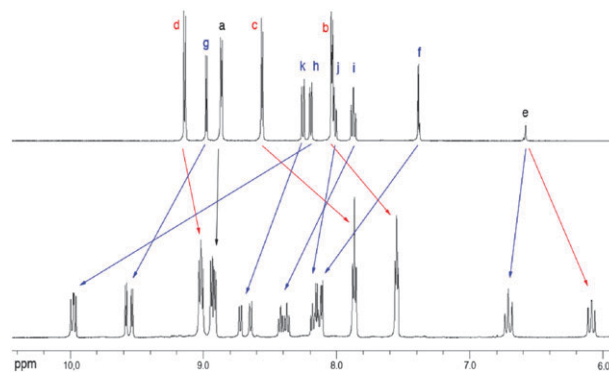
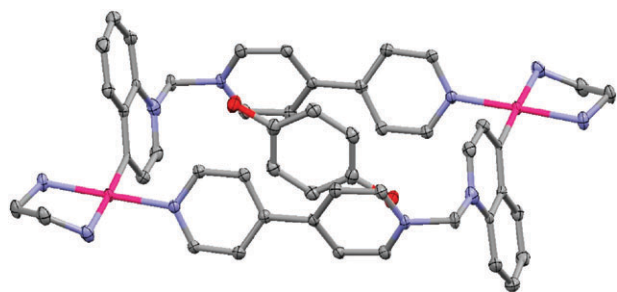


Fig. 1 Partial <sup>1</sup>H NMR spectra (500 MHz, D<sub>2</sub>O, 298 K). Top: free ligand **1**·NO<sub>3</sub> (10 mM). Bottom: **1**·NO<sub>3</sub> (10 mM), Pd(en)(NO<sub>3</sub>)<sub>2</sub> (10 mM), and HQ (5 mM). See Scheme 1 for proton assignments.

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**Fig. 2** Crystal structure of HQ-c3a-6NO<sub>3</sub> showing the 50% probability thermal ellipsoids. Solvent molecules, counterions, and hydrogen atoms have been omitted for clarity.

the centroid defined by the guest. Immediately upon redissolution of the crystals in D<sub>2</sub>O the <sup>1</sup>H NMR spectrum showed a 1 : 1 mixture of its *syn* and *anti* atropoisomers, indicating that the initial equilibrium is re-achieved in seconds. The fact that *anti* stereoisomer of HQ-c3b-6NO<sub>3</sub> crystallizes out preferentially over the *syn* one suggests that crystal packing forces must play a significant role in determining the crystal structure.

To obtain more detailed information about the mechanism of interconversion between the *syn* and *anti* atropoisomers we prepared the corresponding platinum metallocycle. The reaction between ligand 1-NO<sub>3</sub> and Pt(en)(NO<sub>3</sub>)<sub>2</sub> was carried out at 100 °C to exploit the “molecular lock” strategy introduced by Fujita.<sup>8</sup> The reaction allowed the isolation of a 1 : 1 mixture of the *syn* and *anti* atropoisomers of 3b as their hexafluorophosphate salts. The <sup>1</sup>H NMR spectrum of the mixture recorded in a CD<sub>3</sub>NO<sub>2</sub> solution at room temperature is consistent with the presence of a 1 : 1 mixture of the *syn* and *anti* atropoisomers. Upon increasing the temperature the signals gradually broaden, reflecting conformational exchange processes. The coalescence method was used to calculate the  $\Delta G^\ddagger_c$  (75.3 kJ mol<sup>-1</sup>) value for the interconversion process.<sup>9</sup> Similar activation barriers have been determined for the rotation of Pt-N(quinoline) bonds in square planar Pt(II) complexes.<sup>10</sup>

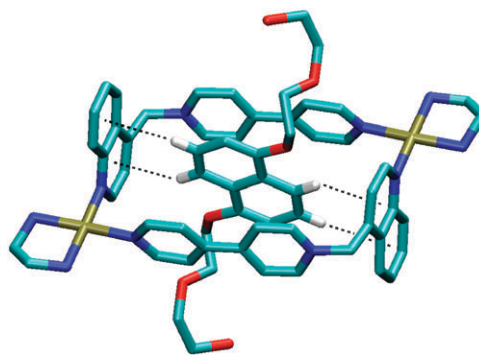
To support that the interconversion between the *syn* and *anti* atropoisomers occurs by rotation around the Pt-N(quinoline)-CH<sub>2</sub> axis we performed DFT calculations at the B3LYP level. In these calculations we used the standard 6-31G(d) basis set for C, H and N ligand atoms, while the LanL2DZ valence and effective core potential functions were used for Pt. Our calculations provide two minimum energy conformations for the metallocycle that correspond to the expected *syn* and *anti* atropoisomers. According to the calculations the *syn* isomer is slightly more stable than the *anti* one in the gas phase, the relative free energy  $\Delta G^\circ_{298} = G^\circ_{298(\text{syn})} - G^\circ_{298(\text{anti})}$  amounting only to -4.52 kJ mol<sup>-1</sup>.<sup>11</sup> In the *anti* isomer the two quinoline units are parallel to each other, while in the *syn* form the metallocycle adopts a cone-like conformation with the least-squares planes defined by the two quinoline units forming an angle of 31.6° (Fig. S24, ESI†). The *syn* → *anti* interconversion process was studied *in vacuo* by using the synchronous transit-guided quasi-Newton method.<sup>12</sup> Our results show that the interconversion between the two isomers is a single-step process involving the rotation of one quinoline unit. Both the *syn* and *anti* isomers are characterised by dihedral angles of *ca.* 75° between the best plane of the

metallocycle and the two planes described by the quinoline units. However, the structure of the transition state that connects the two minimum energy conformations presents a dihedral angle of *ca.* 75°, while the second one defined by the quinoline unit undergoing the rotation process amounts to *ca.* 6° (See ESI†). The computed Gibbs free energy barrier for the *syn* → *anti* interconversion process amounts to 69.9 kJ mol<sup>-1</sup>,<sup>11</sup> which is in reasonably good agreement with the experimental value obtained in CD<sub>3</sub>NO<sub>2</sub> solution (75.3 kJ mol<sup>-1</sup>). This relatively high energy barrier for the interconversion between the *syn* and *anti* isomers is related, at least in part, to the distortion of the Pt(II) coordination environment experienced by the molecule to reach the transition state. Indeed, the calculated bond distances between Pt and the nitrogen atoms of the bipyridine (Pt-N<sub>b</sub>) and quinoline (Pt-N<sub>q</sub>) moieties are 2.11–2.12 Å in both isomers, while in the transition state the Pt-N<sub>b</sub> and Pt-N<sub>q</sub> distances amount to 2.12 and 2.27 Å, respectively. The lengthening of the Pt-N<sub>q</sub> distance in the transition state is a consequence of the steric hindrance caused by the ethylenediamine ligand on the quinoline unit undergoing the rotation process.

The inclusion complex with 1,5-bis[2-(2-hydroxyethoxy)ethoxy]naphthalene (1,5-NPH) has also been studied. In this case, the *anti* stereoisomer is the only one present in solution, as shown by NMR analysis (Fig. S16†). The *anti* atropoisomer is the only one compatible with the presence of the inversion centre evidenced by the NMR spectrum.

As expected, the X-ray structure of 1,5-NPH-c3a-6NO<sub>3</sub> (Fig. 3) shows the exclusive presence of the *anti* stereoisomer in the solid state.<sup>13</sup> Again, both components lie about a common inversion centre located at the centroid defined by the guest. The shorter distance between the centroids of the central C–C bonds of the two bipyridine moieties (0.14 Å shorter than in the hydroquinone inclusion complex) together with the almost coplanar arrangement of the pyridine rings (15° dihedral angle in comparison to the 28° dihedral angle found in HQ-c3a-6NO<sub>3</sub>) demonstrate enhanced host–guest interactions, undoubtedly due to the larger  $\pi$ -surface and the stronger  $\pi$ -donor character of the naphthalene guest.

Four C–H··· $\pi$  interactions between the guest (at positions C-3, C-4, C-7, and C-8) and the quinoline rings, resulting from the perfect host–guest fit, further contribute to the structure stabilization in solution and in solid phase.<sup>14</sup> In the *syn* isomer,



**Fig. 3** Crystal structure of 1,5-NPH-c3a-6NO<sub>3</sub> showing the four C–H··· $\pi$  interactions.<sup>14</sup> Solvent molecules and hydrogen atoms have been omitted for clarity.

only three of these interactions would be possible. Steric hindrance between the polyether chains and the quinoline could also contribute to the destabilization of the *syn* system. A third experiment was carried out trying to force the formation of the *syn* stereoisomer. This time, the template of choice, 1,3-bis[2-(2-hydroxyethoxy)ethoxy]naphthalene, is the result of shifting the unfavourable polyether chain at the C-5 position, that creates steric hindrance with the quinoline moiety in the *syn* stereoisomer, to the C-3 position, in which a similar impediment is introduced with the quinoline in the undesired *anti* isomer. The  $^1\text{H}$  NMR spectrum of the resulting solution shows the presence of two isomers in an approximate 2 : 1 ratio. Although a specific assignment of the two sets of signals to a particular isomer (*syn* or *anti*) was not possible, these results clearly indicate that the nature of the aromatic guest plays an important role in the proportion of the two atropoisomers present in solution. The presence of two stereoisomers can probably be ascribed to the fact that, assuming a similar insertion mode for 1,3-NPH, only three C–H $\cdots\pi$  bonds (at positions C-4, C-7 and C-8) can be formed between 1,3-NPH and the quinoline systems in a *syn* conformation.

In conclusion, we have demonstrated that a judicious selection of the aromatic guest allows the stereoselective self-assembly of atropoisomeric Pd(II) metallocycles containing quinoline moieties.

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