



# Controlling molecular shuttling in a rotaxane with weak ring recognition sites†

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**We describe a rotaxane molecular shuttle encompassing triazolium and tertiary ammonium units as weak recognition sites for the ring. Such a design, which differs from that of typical controllable rotaxanes, allows the precise tuning of the ring distribution among the two sites – i.e., the coconformational equilibrium – by changing the solvent polarity or the nature of the counteranions. Shuttling of the ring between the two stations can also be toggled by acid–base stimuli. Such an approach is paradigmatic to obtain rotaxanes that can sense environmental changes and transduce them into a coconformational response and opens avenues for novel applications in sensing and stimuli-responsive materials.**

Controllable molecular shuttles are a class of bistable rotaxanes whose axle component is endowed with two different recognition sites (stations) for the ring component. These rotaxanes can exist in two isomers, called coconformers, which display different stabilities and are in equilibrium. Upon application of a suitable chemical, electrical or optical stimulus, the relative affinity of the macrocycle for the two stations can be modified, causing the former to move from one station to the other.<sup>1,2</sup> More precisely, the transformation triggered by the stimulus determines a new coconformational equilibrium, in which the distribution of the ring among the stations is different from the original one. The reversibility of the switching process, usually by application of an opposite stimulus, enables the reset to the original distribution.

Because of these unique characteristics, arising by the presence of the mechanical bond, molecular shuttles and related species have become attractive for utilization in fields such as functional materials, sensing, and information storage, as well as molecular machines and motors.<sup>1,3</sup> To ensure a clear-cut

coconformational switching, it is important that, in either state, the macrocycle interacts with the two stations with a markedly different strength, and that the pattern is inverted upon changing the state. In this manner, it can be ensured that the ring resides exclusively on station A in the original state, while it moves completely to station B in the switched state.<sup>1,4</sup>

A popular and successful design for chemically controllable reversible molecular shuttles is based on a secondary ammonium station encircled by a crown ether which, upon deprotonation of the ammonium center, shuttles to a pH-insensitive weaker station.<sup>5</sup> Among others, 4,4'-bipyridinium or triazolium units are good candidates for the role of the weaker station in combination with *sec*-ammonium.<sup>6</sup> As the strength of interaction between a crown ether such as dibenzo-24-crown-8 (DB24C8) and *sec*-ammonium centers is remarkably high,<sup>7</sup> in rotaxanes displaying this motif the dominant coconformer is the one in which the macrocycle sits onto the *sec*-ammonium station, regardless of the nature of the secondary station.<sup>6</sup>

The deprotonation of the ammonium center with a base switches off the recognition of the ring, thereby reversing the unbalanced distribution. Overall, it can be said that complete ring shuttling from one station to the other is achieved. This result requires (i) a large free energy difference between the two coconformations in either state, and (ii) the ability of the stimulus to change significantly such difference (Fig. 1a).

In 2008, Takata and coworkers showed that the *sec*-ammonium station of a rotaxane can be converted into a *tert*-ammonium moiety that exhibits a much lower affinity for DB24C8.<sup>8</sup> This approach could be utilized to make rotaxanes with coconformational equilibria characterized by much lower  $\Delta G^\circ$  and a markedly different switching behaviour. We envisioned that the combination of low affinity *tert*-ammonium and triazolium stations should result in a bistable rotaxane where subtle variations in the energy of the ground state suffice to bias the coconformational equilibrium (Fig. 1b). To date, however, only one rotaxane comprising a *tert*-ammonium and a urea station that responds to chemical stimuli ( $\text{Na}^+$  ions or protonation/deprotonation) has been described.<sup>9</sup>

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a) previous works: large  $\Delta G^\circ$ , strong switching, fixed distribution



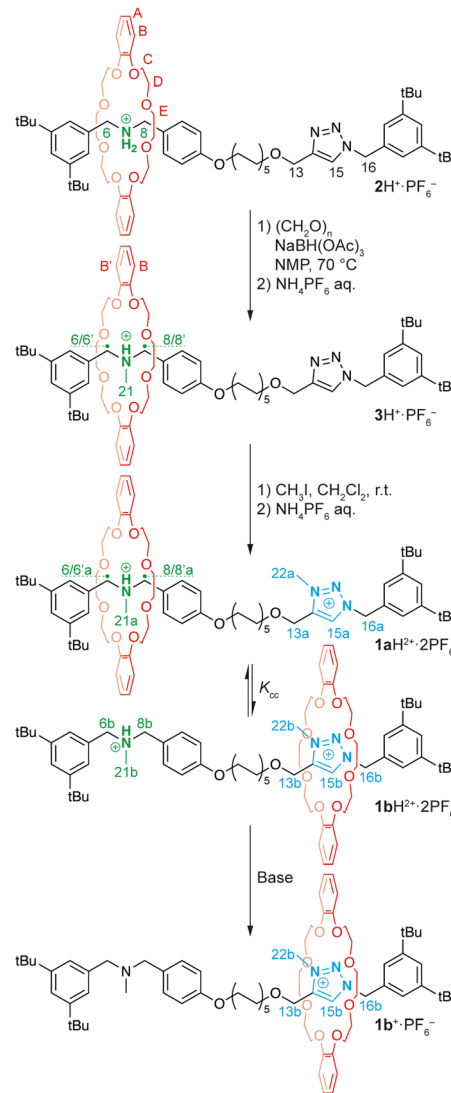
b) this work: small  $\Delta G^\circ$ , weak switching, tunable distribution



**Fig. 1** Schematic energy-level diagram illustrating the switching between two coconformational states in a molecular shuttle. In (a), the large  $\Delta G$  between the two coconformations determines a clear positioning of the rings in both the original and the switched state; to achieve this goal, the stimulus must cause a very significant change of  $\Delta G$ . In (b), the modest free energy difference between the coconformations can result in a small, potentially tunable, population bias. The red circles denote the rotaxane coconformations in a population; “+S” indicates the stimulus inducing the free energy change and “−S” is the opposite stimulus.

Herein, we report on the preparation and operation of a bistable molecular shuttle endowed with a *tert*-ammonium and a triazolium station (**1aH<sup>2+</sup>**, Scheme 1) in which the coconformational equilibrium is highly sensitive to the polarity of the medium. We also demonstrate that the relative distribution of the rings between the stations can be adjusted to different values by modifying the ion-pairing interactions between the positively charged stations and their counteranions. The position of the ring between the two stations can also be switched by classical acid/base treatment.

The synthesis of target rotaxane **1aH<sup>2+</sup>** is depicted in Scheme 1. The *sec*-ammonium [2]rotaxane **2H<sup>+</sup>**·PF<sub>6</sub><sup>−</sup> was synthesized by stoppering the pseudorotaxane complex formed between the DB24C8 and the axle equipped with a *sec*-ammonium recognition motif by copper-catalyzed azide–alkyne cycloaddition (CuAAC) (see the ESI†). In order to avoid simultaneous interaction with the two weak stations—*tert*-ammonium and triazolium—in the target rotaxane, a C10 spacer alkyl chain was introduced. The <sup>1</sup>H NMR spectral features of **2H<sup>+</sup>**·PF<sub>6</sub><sup>−</sup> in CD<sub>2</sub>Cl<sub>2</sub> are consistent with a [2]rotaxane structure where the DB24C8 encircles the dibenzyl ammonium moiety. The *tert*-ammonium station was installed by methylation of the *sec*-ammonium using a variation of the Eschweiler–Clarke reaction.<sup>8</sup> Reaction of **2H<sup>+</sup>**·PF<sub>6</sub><sup>−</sup> with paraformaldehyde and triacetoxyborohydride (20 equivalents each) in *N*-methyl-2-pyrrolidone at 70 °C, followed by anion exchange, allowed the isolation of **3H<sup>+</sup>**·PF<sub>6</sub><sup>−</sup> in good yield. Methylation of the ammonium moiety was confirmed by the appearance, in the <sup>1</sup>H NMR spectrum, of a doublet at 2.93 ppm corresponding to the methyl group (H<sub>21</sub>) bound to the nitrogen. Due to the loss of the symmetry plane containing the axle, combined with the slow nitrogen inversion on the NMR timescale,<sup>10</sup> the methylene protons become diastereotopic. Specifically, H<sub>6</sub> and H<sub>6'</sub>, as well as H<sub>8</sub> and H<sub>8'</sub>, resonate at different frequencies and display scalar coupling. Additionally, due to the constraints of the mechanical bond, H<sub>B</sub> splits into two sets of



**Scheme 1** Synthesis and coconformational equilibrium of rotaxane **1aH<sup>2+</sup>**·2PF<sub>6</sub><sup>−</sup> and **1bH<sup>2+</sup>**·2PF<sub>6</sub><sup>−</sup>.

resonances (H<sub>B</sub> and H<sub>B'</sub>), *syn* and *anti* with respect to the NMe group (see the ESI†). The signals associated to the triazole portion of the molecule remain mostly unchanged, indicating that, even upon methylation, the majority of rings encircle the ammonium station. ROESY data also confirm this conclusion. It is worth noting that this rotaxane equipped with a *tert*-ammonium and a neutral triazole station can also function as a pH-operated molecular switch (see the ESI†).

The target rotaxane **1H<sup>2+</sup>**·2PF<sub>6</sub><sup>−</sup> was obtained by regioselective methylation of the triazole moiety of rotaxane **3H<sup>+</sup>**·PF<sub>6</sub><sup>−</sup> with iodomethane, followed by anion exchange.<sup>6,11</sup> The <sup>1</sup>H NMR spectrum of **1H<sup>2+</sup>**·2PF<sub>6</sub><sup>−</sup> in CD<sub>2</sub>Cl<sub>2</sub> shows that each signal exists as two distinct sets of resonances in a 95 : 5 ratio. In particular, the triazolium singlet (H<sub>15</sub>) displays two distinct resonances at 9.37 and 8.23 ppm associated with the encircled and free triazolium stations, respectively (Fig. 2a). The same ratio is observed for the two resonances associated with the methyl group of the *tert*-ammonium moiety (H<sub>21</sub>, Fig. 2a). The two sets of





Fig. 2 Partial  $^1\text{H}$  NMR spectra (500 MHz,  $\text{CD}_2\text{Cl}_2$ , 298 K) of a 95:5 mixture of rotaxane  $\mathbf{1aH}^{2+}\cdot 2\text{PF}_6^-$  and  $\mathbf{1bH}^{2+}\cdot 2\text{PF}_6^-$  (a) and the same sample after addition of 1 equivalent of  $\text{tBu}_4\text{NCl}$  displaying mainly  $\mathbf{1bH}^{2+}\cdot 2\text{Cl}^-$  (b). Signals are color-coded and numbered according to Scheme 1. The residual solvent and the tetrabutylammonium signals are color-coded in grey.

resonances can thus be attributed to the two distinct coconformers of rotaxane  $\mathbf{1H}^{2+}$ , in which the ring encircles either the tertiary ammonium ( $\mathbf{1aH}^{2+}\cdot 2\text{PF}_6^-$ ) or the triazolium ( $\mathbf{1bH}^{2+}\cdot 2\text{PF}_6^-$ ) station (Scheme 1).

The presence of both coconformers points to a similar affinity of the DB24C8 for the two stations. In the low polarity  $\text{CD}_2\text{Cl}_2$  solvent with the non-coordinating  $\text{PF}_6^-$  counterion, 95% of the rings encircle the ammonium station; this figure corresponds to a conformational equilibrium constant ( $K_{cc} = [\mathbf{1bH}^{2+}]/[\mathbf{1aH}^{2+}]$ ) of about 0.05 and a free energy difference between the two coconformers of less than 2 kcal mol $^{-1}$ . On the basis of this small  $\Delta G^\circ$ , we hypothesized that even minor perturbations of the medium (e.g. the addition of a different anion or a change in solvent polarity) could affect significantly the population of the two coconformers. We initially investigated the influence of solvent polarity on the conformational equilibrium.  $^1\text{H}$  NMR spectra of rotaxane  $\mathbf{1aH}^{2+}\cdot 2\text{PF}_6^-$  were recorded in acetone- $d_6$ , a solvent remarkably more polar than  $\text{CD}_2\text{Cl}_2$ . Spectra at 298 K revealed significant line broadening indicative of a fast dynamics of exchange between the two stations on the NMR timescale, most likely due to the lower association of the ring to the cationic stations in highly polar solvents. The chemical shift of protons  $\text{H}_{15}$  indicates that in such a polar environment the majority of the rings encircle the triazolium station (see the ESI $^\dagger$ ). To accurately determine the relative population of the two coconformers, the temperature was lowered to 223 K to obtain a slow exchange process on the NMR timescale. In these conditions, the  $[\mathbf{1bH}^{2+}\cdot 2\text{PF}_6^-]/[\mathbf{1aH}^{2+}\cdot 2\text{PF}_6^-]$  ratio is 70:30, corresponding to a  $K_{cc}$  of about 2.3 ( $\Delta G^\circ = -0.5$  kcal mol $^{-1}$ , see the ESI $^\dagger$ ).

In a second series of experiments, chloride anions, which are known to pair with the ammonium site, $^{7,12}$  were added to bias the equilibrium towards the triazolium station in  $\text{CD}_2\text{Cl}_2$ . We measured the relative population of the two stations at increasing concentration of tetrabutylammonium chloride. To our delight, all distributions between the two stations, ranging from 95:5 to less than 1:99 ammonium/triazolium (Fig. 2b), could be obtained, as shown by the relative integration of the signals for the free and complexed triazolium station at 8.23

and 9.37 ppm, respectively (Fig. 3a). Moreover, a complete shift of the equilibrium towards  $\mathbf{1bH}^{2+}$  was reached upon the addition of about one equivalent of  $\text{Cl}^-$ , suggesting that ring shuttling takes place as a consequence of the competing ion-pairing equilibrium (Fig. 3a). $^{12}$  Finally, the  $\text{NH}^+$  proton becomes strongly deshielded, shifting from 7.6 ppm to 12.7 in the presence of chloride anions, while the formation of a tight ion pair between the ammonium and the chloride is indicated by the upfield shift of  $\text{H}_{21b}$ . Interestingly, the absence of hydrogen bonding between the DB24C8 ring and the ammonium  $\text{NH}^+$  restores the fast nitrogen inversion on the NMR timescale. The diastereotopic  $\text{CH}_2$  signals become isochronous and protons  $\text{H}_B$  of the ring are magnetically equivalent. These observations confirm that the ring has moved from the ammonium to the triazolium station. $^{13}$  The identity of  $\mathbf{1bH}^{2+}\cdot 2\text{Cl}^-$  was unambiguously proven by deprotonating the ammonium station: the deprotonated rotaxane  $\mathbf{1b}^+\cdot \text{PF}_6^-$  is afforded, in which the rings are located around the triazolium station (see the ESI $^\dagger$ ). The presence of a single coconformation is confirmed by the presence of sharp resonances for all protons. Comparison of the  $^1\text{H}$  NMR spectra of  $\mathbf{1b}^+\cdot \text{PF}_6^-$  and  $\mathbf{1bH}^{2+}\cdot 2\text{Cl}^-$  showed remarkably similar spectral features. In particular, the



Fig. 3 (a) Titration of  $\mathbf{1aH}^{2+}\cdot 2\text{PF}_6^-$  with  $\text{tBu}_4\text{NCl}$ . Molar fraction of  $\mathbf{1aH}^{2+}$  (hollow circles) and  $\mathbf{1bH}^{2+}$  (full circles). (b) Partial EXSY spectrum (500 MHz,  $\text{CD}_2\text{Cl}_2$ , 298 K,  $t_{\text{mix}} = 200$  ms) showing the chemical exchange between the triazolium signals of  $\mathbf{1aH}^{2+}$  ( $\text{H}_{15a}$ ) and  $\mathbf{1bH}^{2+}$  ( $\text{H}_{15b}$ ).



signal at 9.37 ppm is associated with the complexed triazolium station, while deprotonation of the ammonium was indicated by the disappearance of the doublet at 2.9 ppm and the appearance of a sharp singlet at 2.2 ppm, consistent with the methyl group of a tertiary amine. This experiment also confirms the possibility to operate the system as a classical acid/base molecular switch.

From a kinetic point of view, the analysis of the  $^1\text{H}$  NMR spectra in  $\text{CD}_2\text{Cl}_2$  revealed significant line broadening regardless of the relative population of the stations. This observation is indicative of an exchange equilibrium between the two coconformers that is intermediate-to-slow on the NMR time-scale as confirmed also by exchange NMR (EXSY) experiments (see the ESI†). The in-phase cross-peaks between protons  $\text{H}_{15a}$  and  $\text{H}_{15b}$  are indicative of chemical exchange between  $1a\text{H}^{2+}$  and  $1b\text{H}^{2+}$  (Fig. 3b), which can occur by ring shuttling.

This combined set of data indicates that shuttling between the two chemically distinct stations occurs without the need to formally deactivate one of them and that fine tuning of the conformational equilibrium constant can be achieved.

In conclusion, we reported the synthesis and operation of a rotaxane molecular switch equipped with two weak stations for a DB24C8 macrocycle. Due to the tiny energy difference between the two coconformers, the corresponding equilibrium is highly influenced by the medium (for example, changes in the solvent polarity or the presence of ions). Specifically, the nature and concentration of anions, capable of pairing with the positively charged *tert*-ammonium station, can be exploited to fine tune the distribution of the rings between the two stations. These findings open interesting possibilities to make mechanically interlocked molecules that can sense anions and probe environmental changes in solution and, in perspective, in materials. Bistable rotaxanes with chemically distinct stations but with no strong bias in the ring distribution could be useful also from the viewpoint of molecular machinery.

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## Conflicts of interest

There are no conflicts to declare.

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