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European Journal of Organic Chemistry



Accepted Article

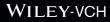
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To be cited as: Eur. J. Org. Chem. 10.1002/ejoc.202000997

Link to VoR: https://doi.org/10.1002/ejoc.202000997



Regulation of Macrocycle Shuttling Rates in [2]Rotaxanes by Amino-Acid Speed Bumps in Organic–Aqueous Solvent Mixtures

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Abstract: A homologous series of two-station [2]rotaxanes incorporating amino-acid units in the molecular thread has been developed. The degenerate [2]rotaxanes exhibit amino-acid specific shuttling rates between two fumaric stations related to the steric factor associated to the amino-acid side chain, as demonstrated by variable-temperature ¹H NMR spectroscopy and exchange spectroscopy (EXSY). This allows tuning of the macrocycle shuttling rate over 4 orders of magnitude, which further has a relatively small solvent dependency.

Introduction

Rotaxanes are the most commonly used systems for constructing artificial molecular machines.^[1] Indeed a variety of effective methods for the synthesis of rotaxanes has been established through numerous synthetic strategies.^[2-5] Among a wide diversity of reported interlocked structures, molecular shuttles comprising a mobile macrocycle threaded on a twostation axle, is one of the most emblematic types of architecture, which serves as an excellent platform for bottom-up studies of dynamic properties.^[6-8] With sub-molecular movement being integral to potential function in molecular machines, regulation of dynamics, both spontaneous and stimulus-induced, is essential and intimately linked with molecular design.[9-12] Developing methods to control the shuttling rate could unlock pioneering applications in the field of catalysis, as exemplified by rotating catalysts where the rotation speed had a direct influence in their performance.[13]

Despite the central importance of the ring shuttling rate in rotaxanes, only a relatively small number of studies have focused on detailed kinetic analyses providing key structureproperty information.^[14–16] Relevant works include the discovery of the key role of water lubricating hydrogen-bonded molecular machines by Leigh,^[17] Stoddart,^[18] and Harada.^[19] The development of systems with steric barriers for unidirectional movement by Stoddart,^[6] and the null impact of the rotaxane axle length on the ring shuttling reported by Hirose.^[14] These contributions are central to understanding the effect of the different macrocycle-thread interactions on the shuttling rate and incorporate such structural parameters in the design of rotaxanes with specific shuttling dynamics.

A family of interest are Leigh's rotaxanes, assembled in one synthetic step and facilitated by hydrogen bond preorganisation evoking a tetraamide macrocycle and fumaramide or succinamide station on the thread.^[20-23] In these systems, polar solvents have a significant effect on the shuttling rate, especially water,^[17,24] producing an increase of the shuttling rate as a consequence of disrupting hydrogen bonding between the macrocycle and the station, on providing competitive H-bonding interactions with the solvent. Despite the easy synthetic accessibility, the control of shuttling rate of the macrocycle has not been well-established and is highly dependent on the polarity solvent. Achieving this control in aqueous solvent mixtures, and ideally in water, is needed for developing operational rotaxanes for potential biological applications. Yet, few examples of rotaxane shuttles in water have been reported with limited ring shuttling kinetic information.^[25-28] While water is shown to accelerate ring movement in hydrogen-bonding rotaxanes, full control implies the ability to slow ring movement and even render it solvent independent to a large degree. For this purpose, introducing a custom steric barrier on the thread that the macrocyclic can only pass over slowly, can produce the desired shuttling rate reduction.

The selection of the steric barrier size is critical, as just one methyl difference may completely block the ring shuttling, as highlighted by macrocycle dethreading experiments in rotaxanes with stoppers of different sizes.^[29] In this work we developed designer rotaxane systems building on a benchmark system, namely Leigh's fumaramide rotaxane (Figure 1 top), by incorporating steric barriers to control the macrocycle shuttling rate in both non-polar organic solvents and organic aqueous mixtures.^[30,31] For this propose, we synthesized degenerate molecular shuttles containing specific amino-acid units in the thread which are able to act as speed bumps to control the macrocycle shuttling speed (Figure 1, bottom) and sought to generalize this approach for a range of amino-acid units. In these systems, which have two identical stations, the macrocycle would have an equal probability of residing on each station and stochastic macrocycle shuttling would be operational. These degenerate [2]rotaxanes systems provide an appropriate platform to measure the shuttling process by dynamic NMR spectroscopy using variable temperature ¹H NMR and EXSY NMR.

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FULL PAPER

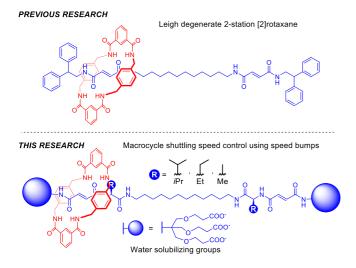


Figure 1. Representation of Leigh's degenerate 2-station [2]rotaxane (from Ref ^[21]) and the analogues developed in this project incorporating amino-acid speed bumps and water solubilizing groups on the stopper.

Results and Discussion

The prototype 2-station molecular machine design developed by Leigh and coworkers (Figure 1 top),^[20] which comprises a tetraamide macrocycle and two-fumaramide stations, has a typical energy barrier of 16.2 kcal/mol to ring displacement corresponding to a shuttling speed of 8 times per second in CDCl₃ at 25 °C.^[21] This barrier is greatly affected by polar solvents such as water that weaken the hydrogen-bond strength between the macrocycle and the station. This results in a reduction of the energy barrier and consequently an enhancement of the shuttling rate.^[17,24] In the current design (Figure 1 bottom), we integrated a multi-ester stopper that can be hydrolyzed to obtain six negatively charged carboxylate groups offering enhanced solubility in polar solvents, including water. Additionally, amino-acid units were integrated into the

molecular thread to act as speed bumps in our prototype rotaxane model (Figure 1, bottom). In particular, amino-acid speed bumps with R = Me, Et and *i*Pr (using the amino acids Ala, Abu and Val, respectively) have an ideal size to act as steric speed bumps for the macrocycle employed. The wide variety of commercially-available natural and non-natural amino acids affords a selection of side chains with the appropriate size and simplifies the synthesis of the corresponding axles. Among different possible stations, the fumaramide station was selected as it has the larger hydrogen bonding affinity towards the macrocycle it favours the positioning of the macrocycle over the station in polar solvents. Whereas the strongly hydrogen bond disrupting solvent DMSO displaces the macrocycle from the low affinity dipeptide stations,[32] the fumaramide station is still occupied by the macrocycle for at least 85% of the time in DMSO.^[33] Stopper 6 (Figure 2) was designed to impart water solubility of the rotaxanes in water upon hydrolysis of the tertbutyl groups. Furthermore, the tert-butyl groups enhanced the solubility of thread 2 and rotaxane 1 in organic solvents.

The synthesis of these structures was carried out as described in Figure 2. Amide bond formation on reacting 1,12diaminododecane with the activated Boc-Aaa-OH followed by deprotection yielded compound 3 in 80-90% yield over two steps. Stopper 6 was prepared according to a literature procedure by condensation of tris(hydroxymethyl)aminomethane (TRIS) with tert-butyl acrylate [34] Amide bond formation with monomethyl fumarate followed by deprotection in basic conditions yielded 4 in 61% yield over two steps. Peptidic coupling of 3 with 4 yielded thread 2 in 40-50% yields. The final rotaxane formation reaction was performed by simultaneously separate addition of chloroform solutions of p-xylylenediamine and of isophthaloyl dichloride to that of thread 2 and triethylamine in chloroform during 2 h via a syringe pump. The target rotaxanes 1 were successfully isolated in 10-16% yield after column chromatography purification. Hydrolysis of rotaxane 1a in neat formic acid yielded rotaxane 1a-WS quantitatively. The rotaxanes were characterized by ¹H, ¹³C, 2D NMR and high resolution mass spectrometry.

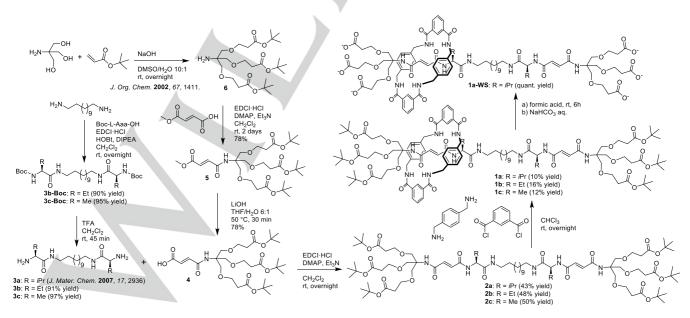


Figure 2. Synthesis of the [2]rotaxanes. Stopper 6 and valine amide 3a were prepared using literature procedures.^[34-36]

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Initial analysis of the ¹H NMR of rotaxane **1a** in CDCl₃ at room temperature showed two sets of signals for the fumaramide protons, which could be assigned to the free and occupied stations. Whereas the free station fumaramide proton resonances (S2) had the same chemical shift observed in the thread, the presence of the macrocycle produced a $\Delta \delta > 1$ ppm upfield shift in the fumaramide signals S2 attributed to the deshielding produced by the macrocycle phenyl rings (Figure 3). The dynamic nature of ring movement within the rotaxane was determined by EXSY ¹H NMR. EXSY allows direct measurement of dynamic process in equilibrated systems as cross peaks between the two signals that undergo exchange during the mixing time of the experiment.^[37]

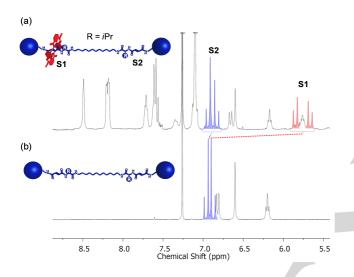


Figure 3. ¹H NMR in CDCl₃ at rt, 300 MHz: (a) Rotaxane 1a, (b) thread 2a. Fumaramide protons labelled as S1 for occupied station and S2 for free station.

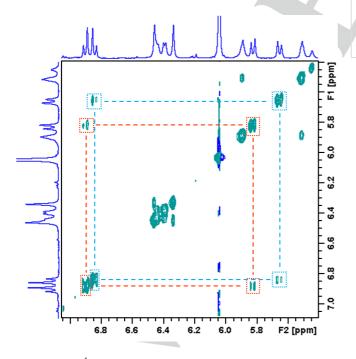


Figure 4. EXSY ¹H NMR of rotaxane **1a** in TCE-*d*₂ at 85 °C, 600 MHz, 250 ms mixing time.

A similar spectrum was obtained for rotaxane **1b**, comprising an ethyl appendage rather than an isopropyl group, two sets of signals at room temperature for the fumaramide protons being observed. In contrast, further downsizing to a methyl appendage in rotaxane **1c** resulted in a single set of broad signals for the fumaramide protons, indicating a relatively fast exchange between the signals of the two stations on the NMR timescale (Figure 5). This observation qualitatively confirms that the macrocycle undergoes a rapid shutting in this case.

To evaluate the effect of the amino-acid speed bump on the macrocycle shuttling, variable temperature (VT) ¹H NMR experiments in TCE- d_2 were performed (600 MHz spectrometer). Whereas no signal coalescence was observed for rotaxane 1a in the 25-105 °C temperature range, we could observe coalescence for rotaxanes 1b and 1c. This showed spontaneous shuttling on the ms-us timescale at higher temperature. The tBu signals ($\Delta \delta \approx 0.045$ ppm) of the stopper has a coalescence temperature > 105 °C for 1a, 75 °C for 1b and -10 °C for 1c, showing a direct effect of the steric bulk on the shuttling rate, the larger the substituent (*i*Pr > Et > Me) the lower the rate. A similar trend was observed for the fumaramide signals ($\Delta \delta \approx 1$ ppm). In this case we only observed a coalescence of the fumaramide signals for rotaxane 1c (Figure 5). Although no coalescence was reached for the fumaramide signals in rotaxanes 1a and 1b, signal broadening indicated the slow dynamics of the macrocycle shuttling in these systems. Overall, the differences observed in coalescence temperatures reflect a reduction of the ring shuttling rate by the increase in amino-acid unit steric size from 1c to 1a (Table 1).

Table 1. Coalescence temperatures in TCE- d_2 , 600 MHz. Signal width at half-
height in brackets at the temperature displayed in the table for protons not
reaching coalescence. Fumaramide signals $\Delta \delta \approx 1$ ppm (600 Hz). *t*Bu signals
 $\Delta \delta \approx 0.045$ ppm (27 Hz).

	P			
Rotaxane	Fumaramide	Stopper <i>t</i> Bu		
1a	> 105 °C (7 Hz)	> 105 °C (8 Hz)		
1b	> 95 °C (20 Hz)	75 °C		
1c	25 °C	–10 °C		

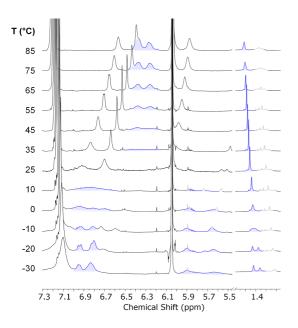


Figure 5. VT ¹H-NMR of alanine-containing rotaxane **1c** with the signals displaying coalescence marked in blue (600 MHz in TCE- d_2). Solvent impurities are marked in grey.

From the VT ¹H NMR experiments it is possible to obtain further information on the dynamics of the process. Increasing the temperature produces an increment of the exchange rate, producing a broadening of the signals until coalescence is achieved when the exchange rate approaches the frequency difference between the two signals. Further heating produces a fast exchange and a spectrum with individual signals at the average frequency of the signals in this symmetrical molecule. This phenomenon can be used to extract quantitative information of the dynamic process by performing a dynamic NMR lineshape fitting.^[14,38,39] The results show that the amino-acid speed bump has a direct influence in the shuttling rate, spanning the measured k_{ex} over 4 orders of magnitude (Figure 6 and Table 2). For the three rotaxanes 1a, 1b and 1c, EXSY ¹H NMR experiments showed cross peaks connecting the different fumaramide and the tBu protons allowing determination of the associated kex (see Figure 6 and supporting information for further details).

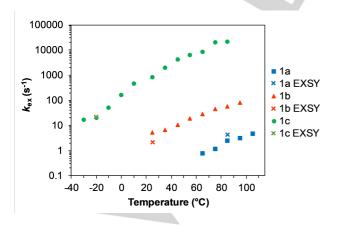


Figure 6. Exchange rate constants determined for rotaxanes **1a**, **1b** and **1c** in TCE- d_2 at different temperatures by lineshape analysis (squares, triangles and circles) and EXSY (crosses) ¹H NMR, 600 MHz.

In order to rationalize the significant changes in the shuttling rates introduced by the amino-acid speed bumps, we evaluated the precise steric hindrance introduced in each case. To this end, we used the steric parameter S⁰ of the Me, Et and *i*Pr groups to determine if there is a direct relation between sterics vs energy barrier.^[30,40] The steric parameter S⁰, derived empirically from the quaternization rate constant of methylation reactions of ortho-substituted pyridines, has been proposed as a quantitative scale in the evaluation the steric bulk of substituents in the shuttling behavior of [2]rotaxanes.^[30] A plot of ΔG^{\ddagger} vs S⁰ shows a linear relation showing the direct relation and preponderant nature of the steric parameter on the process. This data suggests that the speed bump does not significantly affect the complementary hydrogen bonding interactions between the tetraamide macrocycle and the fumaramide stations. Therefore, the energy levels of S1 and S2 would remain unchanged and the speed bumps only affects the energy of the transition state that connects stations S1 and S2 as schematized in Figure 7b.

We further considered the CPK substituent volume as an alternative parameter that can be used to evaluate the steric bulk of amino-acid side chain.^[41] The CPK volumes offer a good linearity with the observed experimental shuttling energy barrier (see Figure 7d and Supporting Information). We envisage that CPK volumes could be a useful tool to select appropriate speed bumps in rotaxane systems. As cited above, the parameter S⁰ provides a good description of the overall steric bulk in systems for small groups such as Me, Et and *i*Pr groups. However, S⁰ does not describe accurately the overall steric bulk of larger groups, while the use of the relative CPK volume of the amino-acid side chain with regard to the macrocycle cavity size provides a better description of the speed bump steric size (see Supporting information). Analysis of the macrocycle cavity showed a diameter of 4.88 Å, corresponding to a spherical void volume of 61 Å³. The overall size of the speed bump was obtained as the CPK volume giving a value of 49 Å³, 67 Å³, and 85 Å³ for Ala, Abu, and Val, respectively (see Figure 7d and Supporting Information). A plot of ΔG^{\ddagger} vs CPK volume shows a linear relation (Figure 7d), therefore providing a good representation of the overall steric bulk. This analysis also provides a physical meaning by comparing the relative size of the cavity of the macrocycle and the CPK volume of the speed bump, which can be readily estimated in analogy to Rebek's 55% optimal occupancy rule.^[42] The Ala CPK volume is smaller than the macrocycle cavity (speed bump factor = V_{Ala}/V_{Cavity} = 0.80), and therefore the steric speed bump effect will be relatively small or negligible. The Abu CPK volume is similar to that of the macrocycle cavity (speed bump factor = V_{Abu}/V_{Cavity} = 1.10) showing a medium steric speed bump effect, and the Val CPK volume is significantly larger than the macrocycle cavity (speed bump factor = V_{Val}/V_{Cavity} = 1.39) showing larger speed bump effect (Figure 7d). Overall, the relative speed bump CPK volume with regard to the macrocycle cavity, defined as speed *bump factor* = V_{Val}/V_{Cavity} , analysis described here represents the observed experimental speed bump effect. We envisage that the use of speed bump factors can be a useful tool in estimating custom macrocycle shuttling rates in designer systems. (Note: variants with significantly faster or slower ring shuttling rates than the examples measured herein are incompatible with NMR analysis).

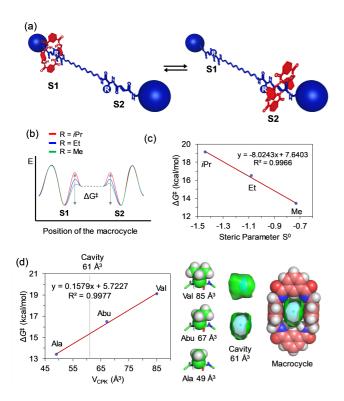


Figure 7. (a) Cartoon representation of the degenerate 2-station [2]rotaxanes developed in this project. (b) Schematic representation of the amino-acid speed bump effect on the ring-on-thread energy profile. (c) Experimental shuttling energy barrier vs steric parameter S^0 (see refs^[30,40] for definition of S^0). (d) Experimental shuttling energy barrier vs amino-acid side chain CPK volume. Atoms used to determine the amino-acid speed bump volume and representation of the macrocycle cavity. Ala = alanine, Abu = 2-aminobutanoic acid, Val = Valine. Experimental data in TCE- d_2 at 25 °C.

Further evidence of the small/negligible alanine steric speed bump effect can be obtained by comparing the 13.4 kcal/mol free energy of activation in rotaxane **1c** with more weakly binding stations - typically 11–12 kcal/mol,^[12,20–22,32,43] and 16.2 kcal/mol for the high affinity fumaramide station (Figure 1).^[21] The activation energy found for rotaxane **1c** is already limited by bond breaking between macrocycle and initial station and chain folding. Therefore, the size of the alanine speed bump is effectively not significantly hindering the shuttling rate.

Having measured processes in a non-competitive solvent, we further evaluated the effect of polar and protic solvents in the macrocycle shuttling rate in rotaxane **1a**. While addition of 5% MeOD to TCE- d_2 did not produce any measurable acceleration, the more polar mixture CDCl₃/MeOD/D₂O 45:45:10 v/v produced an increase of the shuttling rate from ca. 0.05 s⁻¹ to 0.14 s⁻¹. Then we hydrolysed the rotaxane **1a** to obtain the hexa-anionic rotaxane **1a-WS**. Unfortunately, attempts to dissolve **1a-WS** in pure D₂O at the concentrations required for ¹H NMR spectroscopy were unsuccessful. Full solubilisation was achieved in MeOD/D₂O solutions when high proportions of MeOD were employed, i.e. MeOD/D₂O 9:1, v/v. The determined shuttling rate in this solvent mixture was 0.97 s⁻¹, highlighting the lesser effect of polar solvents on the shuttling rate in these systems, compared with steric influences (Table 2).

Table 2. Shuttling constants (k_{ex}) and the corresponding ΔG^{\ddagger} values for the rotaxanes **1a**, **1b**, **1c** and **1a-WS** in different polar solvent mixtures at 25 °C, 600 MHz

Solvent	rotaxane	<i>k</i> _{ex} (s ⁻¹)	∆G [‡] (kcal/mol)
TCE-d ₂	1a ^[a]	0.07	19.1
TCE-d ₂	1b ^[a]	4.6	16.5
TCE-d ₂	1c ^[a]	980	13.4
TCE-d ₂ /MeOD 95:5	1a ^[b]	<0.05	>19.3
CDCl ₃ /MeOD/D ₂ O 45:45:10	1a ^[b]	0.14	18.6
MeOD/D ₂ O 9:1	1a-WS ^[b]	0.97	17.5

[a] Determined by lineshape fitting VT ¹H NMR and extrapolated at 25 °C. [b] Determined by EXSY ¹H NMR.

Conclusion

A family of degenerate 2-station [2]rotaxanes incorporating amino-acid units in conjunction with fumaramide stations is reported. In these systems the appendages on the amino-acid units act as speed bumps, which serve to slow down (up to 4 orders of magnitude) ring shuttling in accordance with a linear dependence on the appendage steric factor or CPK volume. In this regard a degree of predictability in shuttling rates is possible, profiting from a range of commercially-available amino acids with appendages presenting different steric factors. Equally noteworthy, the effect of aqueous solvents on ring shuttling is relatively small (<10) for the range of different solvent mixtures assayed, compared to the predominant steric influence.

Experimental Section

Materials and methods

All chemicals and solvents were obtained from commercial sources and used without further purification unless specified. Mass spectra and NMR were performed in the CESAMO analytical facilities (Bordeaux, France). **NMR**: ¹H and ¹³C NMR spectra were recorded in a Bruker Avance 300 or Bruker Avance III 600 spectrometer. Chemical shifts are reported in ppm and referenced to the solvent residual peaks. **Mass spectrometry**: Field desorption spectra (FD) were recorded on a AccuTOF (JEOL) mass spectrometer using an FD emitter with an emitter voltage of 10 kV. The sample (1–2 μ L) was deposited on a 13 μ m emitter wire. Electrospray spectra (ESI) were recorded on a Qexactive (Thermo) mass spectrometer. The instrument is equipped with an ESI source and spectra were recorded in positive mode. The spray voltage was maintained at 3200 V and capillary temperature set to 320 °C. Samples were introduced by injection through a 20 μ L loop into a 300 μ L/min flow of methanol from the LC pump.

Synthetic procedures

Rotaxanes were prepared using the synthetic scheme described in Figure 2. The TRIS based stopper **6** was prepared using a literature procedure.^[34] Compounds **3a**, **3b**, and **3c** were prepared using literature procedures.^[35,36] Signal labels used in the assignment and ¹H and ¹³C NMR spectra are provided in the Supporting Information.

Synthesis of rotaxane 1a: A solution of p-xylylenediamine (35 mg, 0.255 mmol, 4 eq.) in chloroform (10 mL) and a solution of isophthaloyl dichloride (52 mg, 0.255 mmol, 4 eq.) in chloroform (10 mL) were simultaneously added at room temperature with a syringe pump during 2h to a solution of thread 2a (100 mg, 0.0637 mmol, 1 eq.) and dry triethylamine (71 µL, 0.510 mmol, 8 eq.) in chloroform (20 mL). The reaction mixture was stirred at room temperature overnight under N2 atmosphere. The reaction mixture was filtered through a Celite® pad. The filtrate was washed successively with a sodium bicarbonate solution (10%), water and brine. The organic layer was dried over magnesium sulfate, filtered and concentrated in vacuo. Purification by silica gel column chromatography (cyclohexane/i-PrOH, 90:10 to 84:16, v/v) afforded 1a as a colourless solid (13 mg, 10% yield). ¹H NMR (600 MHz, TCE-d₂) δ 8.55 (s, 2H, H_u), 8.20 (t, J = 7.6 Hz, 4H, H_s), 7.68 (m, 2H, H_v or H_v), 7.63 (t, 2H, J = 7.8 Hz, H_t), 7.61 (m, 2H, H_v or H_v), 7.10 (psq, J = 7.9 Hz, 8H, H_x), 6.94 (m, 1H, H_h or H_h), 6.88 (d, J = 15.0 Hz, 1H, H_g or H_f), 6.87 (s, 1H, $H_{e'}$), 6.80 (d, J = 15.1 Hz, 1H, H_a or H_f), 6.56 (d, J = 8.6 Hz, 1H, H_h or $H_{h'}$), 6.49 (s, 1H, H_e), 6.06 (m, 1H, H_l), 5.84 (d, J = 14.6 Hz, 1H, $H_{f'}$ or $H_{a'}$), 5.69 (brs, 1H, $H_{f'}$), 5.58 (d, J = 14.6 Hz, 1H, $H_{f'}$ or $H_{a'}$), 4.60 (m, 4H, H_w), 4.40 – 4.18 (m, 5H, H_w + H_i), 3.71 (m, 7H, H_i + H_d), 3.68 (m, 6H, $H_{d'}$), 3.64 (t, J = 6.3 Hz, 12H, H_c), 3.41 – 3.03 (m, 4H, H_m), 2.45 (m, 12H, H_b), 2.11 - 2.06 (m, 1H, H_j), 1.88 - 1.81 (m, 1H, H_{j'}), 1.50 - 1.44 (m, 4H, H_n), 1.43 (s, 27H, H_a), 1.39 (s, 27H, $H_{a'}$), 1.33 – 1.19 (m, 16H, H_{o-r}), 0.97 - 0.74 (m, 12H, H_k + H_k). ¹³C NMR (151 MHz, TCE-d₂) δ 170.95, 170.83, 170.17, 169.11, 166.03, 164.90, 164.69, 163.97, 163.71, 136.85, 136.75, 134.47, 133.54, 133.42, 131.77, 131.55, 131.30, 129.29, 129.25, 129.18, 124.33, 80.64, 80.46, 68.71, 68.60, 67.07, 66.95, 61.21, 60.09, 59.36, 58.57, 44.09, 39.71, 39.52, 36.03, 35.99, 31.11, 29.61, 29.33, 29.27, 29.03, 28.96, 27.99, 27.92, 19.14, 18.03 (1 signal overlapping/missing). **HRMS** (ESI): m/z calcd for $C_{112}H_{168}N_{10}O_{28}+Na^{+}$: 2124.19218 $[M+Na]^{+}$; found: 2124.19153.

Synthesis of rotaxane 1b: A solution of p-xylylenediamine (35 mg, 0.259 mmol, 4 eq.) in chloroform (10 mL) and a solution of isophthaloyl dichloride (53 mg, 0.259 mmol, 4 eq.) in chloroform (10 mL) were simultaneously added at room temperature with a syringe pump during 2h to a solution of thread 2b (100 mg, 0.0649 mmol, 1 eq.) and dry triethylamine (72 µL, 0.519 mmol, 8 eq.) in chloroform (20 mL). The reaction mixture was stirred at room temperature overnight under N2 atmosphere. The reaction mixture was filtered through a Celite® pad. The filtrate was washed successively with a sodium bicarbonate solution (10%), water and brine. The organic layer was dried over magnesium sulfate, filtered and concentrated in vacuo. Purification by silica gel column chromatography (cyclohexane/i-PrOH, 9:1 to 8:2, v/v) afforded 1b as a colourless solid (21 mg, 16% yield). ¹H NMR (600 MHz, TCE-d₂) δ 8.54 (s, 2H, H_u), 8.18 (t, J = 8.0 Hz, 4H, H_s), 7.72 (m, 2H, H_v or $H_{v'}$), 7.67 (m, 2H, H_v or $H_{v'}$), 7.62 (t, J = 7.8 Hz, 2H, H_t), 7.21 (m, 2H, $H_{h'}$), 7.11 (s, 8H, H_x), 6.88 (d, J = 14.7 Hz, 1H, H_f or H_g), 6.87 (s, 1H, $H_{e'}$), 6.79 (d, J= 14.9 Hz, 1H, H_f or H_g), 6.72 (d, J = 7.8 Hz, 1H, H_h), 6.52 (s, 1H, H_e), 6.25 (m, 1H, H_l or $H_{l'}$), 5.84 (d, J = 14.9 Hz, 1H, $H_{f'}$ or $H_{g'}$), 5.82 (m, 1H, H_l or $H_{i'}$), 5.61 (d, J = 14.6 Hz, 1H, $H_{f'}$ or $H_{g'}$), 4.61 (m, 4H, H_w), 4.35 (m, 1H, *H_i*), 4.30 (m, 4H, *H_w*), 3.87 (m, 1H, *H_i*), 3.71 (s, 6H, *H_d*), 3.69 (m, 6H, *H_d*), 3.64 (m, 12H, H_c), 3.34 - 3.03 (m, 4H, H_m), 2.45 (m, 12H, H_b), 1.92 -1.61 (dm, 4H, H_j), 1.50 – 1.45 (m, 4H, H_n), 1.43 (s, 27H, H_a or H_a), 1.39 (s, 27H, H_a or $H_{a'}$), 1.33 – 1.16 (m, 16H, H_{0-r}), 0.93 – 0.78 (m, 6H, H_k + H_k). ¹³C NMR (151 MHz, CDCl₃) δ 171.13, 171.10, 171.04, 170.22, 166.71, 166.62, 165.30, 165.11, 164.45, 164.18, 137.06, 134.81, 133.88, 133.80, 132.05, 131.74, 131.51, 129.53, 129.49, 129.28, 124.78, 80.92, 80.73, 69.10, 68.96, 67.34, 67.24, 61.44, 60.48, 55.55, 54.79, 44.53, 44.47, 39.91, 39.75, 36.27, 36.26, 31.58, 30.33, 29.84, 29.57, 29.43, 28.26, 28.20, 26.04, 25.71, 10.15, 10.00 (1 signal overlapping/missing). HRMS (ESI): *m*/z calcd for C₁₁₀H₁₆₄N₁₀O₂₈+Na⁺: 2096.160888 [*M*+Na]⁺; found: 2096.160868 .

Synthesis of rotaxane **1c**: A solution of *p*-xylylenediamine (36 mg, 0.264 mmol, 4 eq.) in chloroform (10 mL) and a solution of isophthaloyl dichloride (54 mg, 0.264 mmol, 4 eq.) in chloroform (10 mL) were simultaneously added at room temperature with a syringe pump during 2h to a solution of thread **2c** (100 mg, 0.0661 mmol, 1 eq.) and dry

triethylamine (74 µL, 0.528 mmol, 8 eq.) in chloroform (20 mL). The reaction mixture was stirred at room temperature overnight under N2 atmosphere. The reaction mixture was filtered through a Celite® pad. The filtrate was washed successively with a sodium bicarbonate solution (10%), water and brine. The organic layer was dried over magnesium sulfate, filtered and concentrated in vacuo. Purification by silica gel column chromatography (cyclohexane/i-PrOH, 9:1 to 8:2, v/v) afforded 1c as a colourless solid (16 mg, 12% yield). ¹H NMR (600 MHz, 85 °C, TCEd₂) δ 8.51 (s, 2H, H_t), 8.24 (m, 4H, H_r), 7.64 (m, 2H, H_s), 7.57 (m, 4H, H_u), 7.18, (s, 8H, H_w), 6.53 (m, 2H, H_h), 6.35 (m, 2H, H_e), 6.28 (m, 4H, H_f + Hg), 5.84 (m, 2H, Hk), 4.53 (m, 8H, Hv), 4.23 (m, 2H, Hi), 3.74 (m, 12H, H_d), 3.70 (m, 12H, H_c), 3.32 - 3.11 (m, 4H, H_l), 2.48 (m, 12H, H_b), 1.71 -1.49 (m, 10H, $H_j + H_m$), 1.47 (s, 54H, H_a), 1.41 – 1.20 (m, 16H, H_{n-q}). ¹³C NMR (151 MHz, TCE-d₂) ō 170.91, 166.16, 164.07, 136.88, 133.58, 131.35, 129.19, 124.34, 80.55, 68.64, 67.00, 60.57, 49.09, 44.08, 39.62, 36.00, 31.32, 30.07, 29.61, 29.21, 29.17, 28.93, 27.95 (5 signals overlapping/missing). HRMS (ESI): m/z calcd for C₁₀₈H₁₆₀N₁₀O₂₈+Na⁺: 2068.12957 [*M*+Na]⁺; found: 2068.12963.

Synthesis of water soluble rotaxane 1a-WS: The valine-containing rotaxane 1a (14 mg, 6.66 µmol) was dissolved in formic acid (1 mL) and stirred at room temperature for 6h. Volatiles were removed in vacuo. Methanol (0.9 mL) and a solution of sodium bicarbonate (3.4 mg, 40.0 µmol, 6 eq.) in water (0.1 mL) were added successively to the residue. Rotaxane 1a-WS was obtained upon removal of the volatiles in vacuo as a colourless solid (quantitative yield). ¹H NMR (600 MHz, MeOD/D₂O 9:1, v/v) δ 8.75 (s, 2H, H_q), 8.14 (dm, 4H, H_o), 7.70 (t, J = 7.8 Hz, 2H, H_p), 7.16 (s, 8H, H_s), 6.97 (d, J = 15.4 Hz, 1H, H_d or H_e), 6.91 (d, J = 15.4 Hz, 1H, H_d or H_e), 5.97 (d, J = 15.0 Hz, 1H, $H_{d'}$ or $H_{e'}$), 5.83 (d, J = 15.0 Hz, 1H, $H_{d'}$ or $H_{e'}$, 4.53 – 4.36 (m, 8H, H_{r}), 4.19 (d, J = 7.3 Hz, 1H, H_{f}), 3.86 $(d, J = 8.7 Hz, 1H, H_f)$, 3.77 (s, 6H, H_c), 3.69 (t, $J = 6.8 Hz, 6H, H_b$), 3.67 - 3.60 (m, 12H, $H_{c'}$ + $H_{b'}$), 3.25 - 3.00 (m, 4H, H_i), 2.42 (t, J = 6.9 Hz, 12H, Ha), 2.13 (m, 1H, Hg), 1.95 (m, 1H, Hg'), 1.49 (m, 4H, Hj), 1.44 (m, 16H, H_{k-n}), 1.04 – 0.90 (m, 12H, H_h + H_h). ¹³C NMR (151 MHz, MeOD/D2O 9:1, v/v) & 180.37, 180.04, 173.40, 173.06, 169.14, 169.10, 167.25, 167.18, 167.06, 166.91, 161.35, 137.63, 137.56, 135.42, 134.94, 134.90, 132.80, 132.56, 132.46, 132.43, 130.59, 130.56, 130.36, 129.86, 126.95, 70.25, 70.15, 70.12, 70.07, 62.69, 62.02, 60.98, 45.16, 40.43, 39.38, 39.29, 30.50, 30.15, 19.68, 19.63 (3 signals overlapping/missing). HRMS (ESI): m/z calcd for $(C_{88}H_{114}N_{10}O_{28})^{6-}+4(H^{+})+2(Na^{+})$: 905.40290 [M+4H+2Na]²⁺; found: 905.40192.

Synthesis of thread 2a: To a solution of valine 3a (199 mg, 0.5 mmol, 1 eq.), fumaric acid stopper 4 (604 mg, 1 mmol, 2 eq.), DMAP (147 mg, 1.2 mmol, 2.4 eq.) and triethylamine (0.17 mL, 1.2 mmol, 2.4 eq.) in dichloromethane (16 mL), EDCI·HCI (230 mg, 1.2 mmol, 2.4 eq.) was added in one portion at room temperature. The reaction mixture was stirred at room temperature for 16h under N2 atmosphere. The reaction mixture was diluted with dichloromethane and washed successively with a citric acid solution (10%), a sodium bicarbonate solution (10%), water and brine. The organic layer was dried over magnesium sulfate, filtered and concentrated in vacuo. Purification by silica gel column chromatography (DCM/MeOH, 100:0 to 96:4, v/v) afforded 2a as a colourless sticky solid (336 mg, 43% yield). ¹H NMR (300 MHz, CDCl₃) δ 6.96 (d, J = 15.0 Hz, 2H, H_f or H_g), 6.88 (d, J = 15.0 Hz, 2H, H_f or H_g), 6.82 (d, J = 8.8 Hz, 2H, H_h), 6.60 (s, 2H, H_e), 6.20 (t, J = 5.7 Hz, 2H, H_l), 4.25 (pst, J = 8.9 Hz, 2H, H_i), 3.74 (s, 12H, H_d), 3.64 (t, J = 6.3 Hz, 12H, H_c), 3.23 (m, 4H, H_m), 2.44 (t, J = 6.1 Hz, 12H, H_b), 2.13 (m, 2H, H_j), 1.53 - 1.46 (m, 4H, H_n), 1.44 (s, 54H, H_a), 1.34 - 1.19 (m, 16H, H_{o-r}), 0.95 (d, ¹³C NMR J = 6.8 Hz, 6H, H_k or $H_{k'}$), 0.93 (d, J = 6.8 Hz, 6H, H_k or $H_{k'}$). (151 MHz, CDCl₃) δ 171.04, 170.82, 164.62, 164.26, 134.74, 132.33, 80.65, 69.03, 67.21, 60.51, 59.30, 39.63, 36.25, 31.00, 29.53, 29.34, 29.13, 28.24, 26.87, 19.42, 18.51. HRMS (ESI): m/z calcd for C₈₀H₁₄₀N₆O₂₄+Na⁺: 1591.98112 [*M*+Na]⁺; found: 1591.98344.

Synthesis of thread **2b**: To a solution of valine **3b** (185 mg, 0.5 mmol, 1 eq.), fumaric acid stopper **4** (604 mg, 1 mmol, 2 eq.), DMAP (147 mg, 1.2 mmol, 2.4 eq.) and triethylamine (0.17 mL, 1.2 mmol, 2.4 eq.) in

dichloromethane (16 mL), EDCI·HCI (230 mg, 1.2 mmol, 2.4 eq.) was added in one portion at room temperature. The reaction mixture was stirred at room temperature for 24h under N2 atmosphere. The mixture was diluted with dichloromethane and washed successively with a citric acid solution (10%), a sodium bicarbonate solution (10%), water and brine. The organic layer was dried over magnesium sulfate, filtered and concentrated in vacuo. Purification by silica gel column chromatography (DCM/MeOH, 100:0 to 96:4, v/v) afforded 2b as a colourless sticky solid (372 mg, 48% yield). ¹H NMR (300 MHz, CDCl₃) δ 6.95 (d, J = 15.0 Hz, 2H, H_f or H_q), 6.84 (d, J = 15.0 Hz, 2H, H_f or H_q), 6.76 (d, J = 8.0 Hz, 2H, H_h), 6.61 (s, 2H, H_e), 6.23 (t, J = 5.7 Hz, 2H, H_l), 4.39 (q, J = 7.1 Hz, 2H, H_i), 3.74 (s, 12H, H_d), 3.65 (t, J = 6.3 Hz, 12H, H_c), 3.23 (m, 4H, H_m), 2.44 (t, J = 6.3 Hz, 12H, H_b), 1.99 – 1.61 (dm, J = 71.3 Hz, 4H, H_j), 1.54 – 1.47 (m, 4H, H_n), 1.44 (s, 54H, H_a), 1.34 - 1.20 (m, 16H, H_{o-r}), 0.92 (t, J = 7.4 Hz, 6H, H_k). ¹³C NMR (101 MHz, CDCl₃) δ 171.16, 171.04, 164.55, 164.23, 134.75, 132.20, 80.66, 69.09, 67.23, 60.52, 54.91, 39.67, 36.27, 29.56, 29.38, 29.17, 28.25, 26.85, 25.62, 10.09 (1 signal overlapping/missing). HRMS (ESI): m/z calcd for C₇₈H₁₃₆N₆O₂₄+Na⁺: 1563.94982 [*M*+Na]⁺; found: 1563.95145.

Synthesis of thread 2c: To a solution of valine 3c (171 mg, 0.5 mmol, 1 eq.), fumaric acid stopper 4 (604 mg, 1 mmol, 2 eq.), DMAP (147 mg, 1.2 mmol, 2.4 eq.) and triethylamine (0.17 mL, 1.2 mmol, 2.4 eq.) in dichloromethane (16 mL), EDCI·HCI (230 mg, 1.2 mmol, 2.4 eq.) was added in one portion at room temperature. The reaction mixture was stirred at room temperature for 24h under N_2 atmosphere. The mixture was diluted with dichloromethane and washed successively with a citric acid solution (10%), a sodium bicarbonate solution (10%), water and brine. The organic layer was dried over magnesium sulfate, filtered and concentrated in vacuo. Purification by silica gel column chromatography (DCM/MeOH, 100:0 to 96:4, v/v) afforded 2c as a colourless sticky solid (375 mg, 50% yield). ¹H NMR (300 MHz, CDCl₃) δ 6.96 (d, J = 15.0 Hz, 2H, H_f or H_a), 6.87 (m, 2H, H_h), 6.82 (d, J = 15.2 Hz, 2H, H_f or H_a), 6.64 (s, 2H, H_e), 6.37 (t, J = 5.7 Hz, 2H, H_k), 4.53 (m, 2H, H_i), 3.74 (s, 12H, H_d), 3.64 (t, J = 6.3 Hz, 12H, H_c), 3.23 (q, J = 6.7 Hz, 4H, H_i), 2.44 (t, J = 6.2 Hz, 12H, H_b), 1.56 – 1.46 (m, 4H, H_m), 1.44 (s, 54H, H_a), 1.39 (d, J = 6.9Hz, 6H, H_i) 1.33 – 1.17 (m, 16H, H_{n-q}). ¹³C NMR (101 MHz, CDCl₃) δ 171.85, 171.07, 164.41, 164.20, 134.85, 132.05, 80.68, 69.10, 67.23, 60.51, 49.21, 39.71, 36.26, 29.52, 29.39, 29.18, 28.25, 26.83, 18.20 (1 signal overlapping/missing). HRMS (ESI): m/z calcd for C₇₆H₁₃₂N₆O₂₄+Na⁺: 1535.91852 [*M*+Na]⁺; found: 1535.92002.

Synthesis of 4: To a solution of methyl ester 5 (920 mg, 1.49 mmol, 1 eq.) in a mixture of THF (100 mL) and water (12 mL), a solution of LiOH·H₂O (118 mg, 4.47 mmol, 3 eq.) in water (5 mL) was added. The reaction mixture was stirred under reflux for 30 min and the reaction progression was monitored by TLC (cyclohexane/EtOAc 1:1). THF was evaporated in vacuo. The aqueous solution was acidified with a 1M HCI solution to pH ≈ 1 and extracted with dichloromethane. The organic layer was washed with brine, dried over magnesium sulfate, filtered and concentrated in vacuo. Purification by silica gel column chromatography (cyclohexane/EtOAc, 9:1 + 0.05% AcOH, v/v) afforded fumaric acid stopper 4 as a thick colourless oil (699 mg, 78% yield). ¹H NMR (300 MHz, CDCl₃) δ 7.21 (d, J = 15.4 Hz, 1H, H_f or H_g), 7.03 (s, 1H, H_e), 6.73 (d, J = 15.4 Hz, 1H, H_f or H_g), 3.76 (s, 6H, H_d), 3.65 (t, J = 6.1 Hz, 6H, H_c), 2.45 (t, J = 6.1 Hz, 6H, H_b), 1.44 (s, 27H, H_a) (1 signal overlapping/missing). ¹³C NMR (151 MHz, CDCl₃) δ 171.35, 169.94, 163.67, 139.66, 129.04, 80.88, 69.11, 67.25, 60.58, 36.13, 28.20. HRMS (ESI): *m/z* calcd for C₂₉H₄₉NO₁₂+Na⁺: 626.31470 [*M*+Na]⁺; found: 626.31508.

Synthesis of 5: To a solution of amine stopper 6 (1.00 g, 1.98 mmol, 1 eq.), monomethyl fumarate (283 mg, 2.18 mmol, 1.1 eq.) and DMAP (293 mg, 2.40 mmol, 1.2 eq.) in dichloromethane (45 mL), EDCI·HCI (460 mg, 2.40 mmol, 1.2 eq.) was added in one portion at room temperature. The reaction mixture was stirred at room temperature for 48h under N₂ atmosphere and monitored by TLC (EtOAc/cyclohexane 3:1). The mixture was washed successively with a citric acid solution (10%), a

sodium bicarbonate solution (10%), water and brine. The organic layer was dried over magnesium sulfate, filtered and concentrated *in vacuo*. Purification by silica gel column chromatography (cyclohexane/EtOAc, 7:3, v/v) afforded methyl ester **5** as a colourless oil (951 mg, 78% yield). ¹**H NMR** (300 MHz, CDCl₃) δ 7.10 (d, *J* = 15.4 Hz, 1H, *H*_f or *H*_g), 6.88 (s, 1H, *H*_e), 6.74 (d, *J* = 15.4 Hz, 1H, *H*_f or *H*_g), 3.76 (s, 3H, *H*_h), 3.75 (s, 6H, *H*_d), 3.65 (t, *J* = 6.2 Hz, 6H, *H*_c), 2.44 (t, *J* = 6.2 Hz, 6H, *H*_b), 1.45 (s, 27H, *H*_g). ¹³**C NMR** (76 MHz, CDCl₃) δ 171.26, 166.22, 163.73, 138.19, 129.24, 80.78, 69.15, 67.26, 60.54, 52.04, 36.18, 28.24. **HRMS** (ESI): *m/z* calcd for C₃₀H₅₁NO₁₂+H⁺: 618.34840 [*M*+H]⁺; found: 618.34733.

Dynamic NMR

Line-broadening dynamic NMR (DNMR) fittings were performed using the DNMR module in Bruker TopSpin 4.0.6. The fumaramide and *t*Bu signals were suitable to perform the fittings to obtain the experimental exchange rate constants (k_{ex}).^[44] The first-order exchange rate constants (k_{ex}) were obtained from the intensities of the EXSY diagonal peaks (I_{AA} and I_{BB}) and the EXSY cross peaks (I_{AB} and I_{BA}) at different mixing times (t_{mix}) as described in the literature to obtain a straight line with slope k_{ex} .^[45,46] Further details and sample spectra are provided in the Supporting Information.

Molecular Modelling

Basic modelling was carried out to evaluate the steric size of the amino-acid speed bumps. Molecular modelling was carried out with the Spartan '18 software using the MMFF force field.^[47] See Supporting information for further details.

Acknowledgements

This project has received funding from the European Union's Horizon 2020 research and innovation programme under the Marie Skłodowska-Curie grant agreement No 796612. Equally, financial support from the Agence Nationale de la Recherche (project ANR-16-CE29-0011) and CNRS is gratefully acknowledged.

Keywords: rotaxanes • molecular shuttles • dynamics • EXSY NMR • hydrogen bonding

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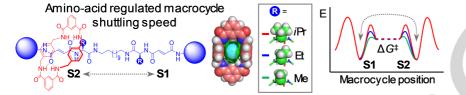
10.1002/ejoc.202000997

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Key Topic: Rotaxane dynamics

FULL PAPER



A family of degenerate 2-station [2]rotaxanes incorporating amino-acid units in conjunction with fumaramide stations is reported in aqueous and pure organic solvents. Appendages on the amino-acid units act as speed bumps, which serve to slow down (up to 4 orders of magnitude) ring shuttling in accordance with a linear dependence on the appendage steric factor and/or CPK volume.

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