Electronic Supplementary Information

Quantifying the barrier for the movement of cyclobis(paraquat-*p*-phenylene) over the dication of monopyrrolotetrathiafulvalene

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Assigned ¹H NMR spectrum of 2•4PF₆

Fig. S1 Partial ¹H NMR spectrum (400 MHz, 298 K, CD₃CN) of the [2]rotaxane **2**•4PF₆ (c = 2 mM). The (CH₃)₃C, CD₂HCN and H₂O signals are not shown in their full height.

Synthesis of compound 9



NaH / DMF / RT / 4.5 h



Scheme S1 Synthesis of the MPTTF derivative 9.

Compounds 11^{S1} and 4^{S2} were synthesised according to literature procedures and compound **9** was synthesised as illustrated in Scheme S1. Initial deprotonation of the pyrrole *N*-H proton by NaH was followed by *N*-alkylation with the iodide **11** affording the MPTTF derivative **9** in 69% yield.

Binding study between the model compound 9 and CBPQT⁴⁺



Scheme S2 Complexation between CBPQT⁴⁺ and the model compound 9 forming the [2]pseudorotaxane 9 CBPQT⁴⁺.

The binding constant K_a between the model compound **9** and CBPQT⁴⁺ and the related free energy ΔG^o for the complexation process were determined using the UV/Vis/NIR dilution method described by Nygaard *et al.*^{S3} Mixing equimolar amounts of the MPTTF derivative **9** (yellow) with a solution of CBPQT•4PF₆ (colourless) in MeCN immediately resulted in a green-coloured solution as a result of the formation of a charge-transfer complex with an absorption band centered at 815 nm in the UV-Vis-NIR spectrum. Two independent starting solutions (2.4 mL) with an absolute concentration of 8.0 × 10⁻⁴ M were prepared and allowed to equilibrate at 298 K in the thermostated cell compartment of the spectrophotometer, before the absorbance A_m was measured. Subsequently, the solutions were repeatedly diluted to 5/6 of its initial concentration with MeCN, resulting in two series of 13 data points [$A^{-1/2}$, cA^{-1}] (Table S1).

Optical path length:	/= 1 cm	
CBPQT•4PF6 in MeCN:	$\epsilon = 3.9 \text{ M}^{-1} \text{ cm}^{-1}$	at λ = 815 nm
9 in MeCN:	$\epsilon = 2.5 \text{ M}^{-1} \text{ cm}^{-1}$	at λ = 815 nm
Total background absorbance Ab:	$\epsilon = 6.4 \text{ M}^{-1} \text{ cm}^{-1}$	at λ = 815 nm

Plotting cA^{-1} as a function of $A^{-1/2}$ afforded a straight line with a slope α of $(1/K_a \epsilon))^{1/2}$ and a y-intercept y_0 of $1/\epsilon l$, where ϵ is the molar extinction coefficient and *l* is the optical path length. This linear plot gave a correlation coefficient of 0.998. The values for K_a and ϵ were obtained from the relationship $K_a = y_0/\alpha^2$ and the already shown relationship for y_0 , respectively. The free energies of complexation ΔG° were obtained from the relationship $\Delta G^\circ = -RT \ln K_a$, where *R* is the gas constant and *T* is the absolute temperature. The error of the binding constant (ΔK_a) is determined using the method described by Nygaard *et al.*^{S3}



Fig. S2 a) UV-Vis-NIR spectra recorded of a 1:1 mixture of **9** and CBPQT⁴⁺ at 298 K in MeCN. b) Plot of 1000 cA^{-1} as a function of $A^{-1/2}$.

<i>c</i> [× 10 ^{−3} M]	Am	A	<i>cA</i> ⁻¹ [× 10 ⁻³ M]	A ^{-1/2}
0.668	0.878	0.874	0.764	1.070
0.557	0.717	0.714	0.780	1.184
0.464	0.584	0.581	0.799	1.312
0.387	0.474	0.471	0.821	1.457
0.322	0.383	0.381	0.846	1.621
0.268	0.307	0.305	0.878	1.810
0.224	0.246	0.245	0.916	2.022
0.186	0.196	0.195	0.953	2.263
0.155	0.156	0.155	0.997	2.536
0.129	0.124	0.123	1.046	2.847
0.108	0.099	0.099	1.096	3.185
0.090	0.078	0.078	1.157	3.585
0.075	0.062	0.061	1.227	4.044
0.668	0.865	0.860	0.776	1.078
0.557	0.705	0.702	0.794	1.194
0.464	0.573	0.570	0.814	1.324
0.387	0.465	0.463	0.836	1.470
0.322	0.376	0.374	0.861	1.635
0.268	0.301	0.299	0.896	1.828
0.224	0.243	0.241	0.928	2.035
0.186	0.196	0.195	0.956	2.267
0.155	0.157	0.156	0.993	2.531
0.129	0.126	0.125	1.033	2.830
0.108	0.100	0.100	1.083	3.167
0.090	0.079	0.079	1.145	3.567
0.075	0.062	0.062	1.212	4.020

Table S1 The measured absorbance A_m for a 1:1 mixture of CBPQT-4PF₆ and **9** (MeCN, 298 K) at $\lambda_{max} = 815$ nm at different concentrations *c*, and the actual absorbance *A* obtained by subtracting the background absorbance A_b at 815 nm from the measured absorbance A_m (*i.e.* $A = A_m - A_b = A_m - 6.4 \text{ M}^{-1}\text{cm}^{-1} \times c \times I$)

Binding study between the semidumbbell 1 and CBPQT⁴⁺



Scheme S3 Complexation between CBPQT⁴⁺ and the semidumbbell 1 forming the [2]pseudorotaxane 1 CBPQT⁴⁺.

The binding constant K_a between the semidumbbell **1** and CBPQT⁴⁺ and the related free energy ΔG^o for the complexation process were determined using the UV-Vis-NIR dilution method described by Nygaard *et al.*^{S3} As the complexation process is slow exchange, the stock solutions were made one day in advance by mixing equimolar amounts of the semidumbbell **1** (yellow) with a solution of CBPQT•4PF₆ (colorless) in MeCN. Overnight the stock solutions turned green as a result of the formation of a charge-transfer complex with an absorption band centered at 816 nm in the UV-Vis-NIR spectrum. Appropriate dilutions of the two independent stock solutions produced samples with absolute concentrations in the range of 6.7×10^{-5} to 8.0×10^{-4} M which again was left overnight to equilibrate. The samples were allowed to equilibrate at 298 K in the thermostated cell compartment of the spectrophotometer, before the absorbance A_m was measured which resulted in two series of 15 data points [$1A^{-1/2}$, cA^{-1}] (Table S2).

Optical path length:	<i>l</i> = 1 cm	
CBPQT•4PF ₆ in MeCN:	$\epsilon = 4.1 \text{ M}^{-1} \text{ cm}^{-1}$	at λ = 816 nm
1 in MeCN:	$\epsilon = 21 \text{ M}^{-1} \text{ cm}^{-1}$	at λ = 816 nm
Total background absorbance Ab:	$\epsilon = 25 \text{ M}^{-1} \text{ cm}^{-1}$	at λ = 816 nm

Plotting cA^{-1} as a function of $A^{-1/2}$ afforded a straight line with a slope α of $(1/K_a \epsilon I)^{1/2}$ and a y-intercept y_0 of $1/\epsilon I$, where ϵ is the molar extinction coefficient and I is the optical path length. This linear plot gave a correlation coefficient of 0.980. The values for K_a and ϵ were obtained from the relationship $K_a = y_0/\alpha^2$ and the already shown relationship for y_0 , respectively. The free energies of complexation ΔG° were obtained from the relationship $\Delta G^\circ = -RT \ln K_a$, where R is the gas constant and T is the absolute temperature. The error of the binding constant (ΔK_a) is determined using the method described by Nygaard *et al.*^{S3}



Fig. S3 a) UV-Vis-NIR spectra recorded of a 1:1 mixture of 1 and CBPQT⁴⁺ at 298 K in MeCN. b) Plot of 1000 cA^{-1} as a function of $A^{-1/2}$.

<i>c</i> [× 10 ⁻³ M]	Am	А	<i>cA</i> ⁻¹ [× 10 ⁻³ M]	A ^{-1/2}
0.799	0.942	0.922	0.867	1.041
0.666	0.764	0.747	0.892	1.157
0.549	0.607	0.594	0.925	1.298
0.466	0.503	0.492	0.948	1.426
0.383	0.398	0.388	0.987	1.605
0.316	0.317	0.309	1.024	1.800
0.266	0.257	0.250	1.063	1.999
0.216	0.200	0.194	1.111	2.268
0.183	0.158	0.153	1.192	2.553
0.150	0.125	0.121	1.241	2.876
0.133	0.106	0.103	1.295	3.121
0.117	0.090	0.087	1.338	3.382
0.100	0.072	0.070	1.434	3.786
0.083	0.060	0.058	1.423	4.141
0.067	0.041	0.039	1.702	5.040
0.808	0.961	0.941	0.859	1.031
0.674	0.771	0.755	0.893	1.151
0.556	0.615	0.601	0.925	1.290
0.471	0.506	0.494	0.953	1.422
0.387	0.403	0.393	0.984	1.595
0.320	0.321	0.313	1.022	1.787
0.269	0.257	0.251	1.074	1.998
0.219	0.200	0.194	1.128	2.269
0.185	0.156	0.151	1.222	2.570
0.152	0.115	0.111	1.366	2.998
0.135	0.103	0.099	1.360	3.174
0.118	0.087	0.084	1.405	3.451
0.101	0.068	0.066	1.541	3.906
0.084	0.057	0.055	1.539	4.281
0.067	0.039	0.038	1.786	5.164

Table S2 The measured absorbance A_m for a 1:1 mixture of CBPQT•4PF₆ and **1** (MeCN, 298 K) at λ_{max} = 816 nm at different concentrations *c*, and the actual absorbance *A* obtained by subtracting the background absorbance A_b at 816 nM from the measured absorbance A_m (*i.e.* $A = A_m - A_b = A_m - 25 \text{ M}^{-1} \text{ cm}^{-1} \times c \times I$)

Illustration of the redox processes taking place upon electrochemical oxidation of 2^{4+} and $1 \subset CBPQT^{4+}$



Scheme S4 Cartoon representation of the redox processes to produce 2•OP⁶⁺ upon electrochemical oxidation of 2•MPTTF⁴⁺.



Scheme S5 Cartoon representation of the redox processes to produce $1 \subset CBPQT \cdot OP^{6+}$ upon electrochemical oxidation of $1 \subset CBPQT \cdot MPTTF^{4+}$.

Kinetic investigations of 1⊂CBPQT•MPTTF⁴⁺



Fig. S4 UV-Vis-NIR absorption spectra of the [2]pseudorotaxane 1⊂CBPQT•MPTTF⁴⁺ in MeCN at 298 K. The top spectrum is recorded as soon as possible after heating the solution to 298 K and the subsequent spectra are recorded every 10 s.



Assigned ¹H NMR spectrum of 1⊂CBPQT⁴⁺

Fig. S5 Partial ¹H NMR spectrum (400 MHz, 298 K, CD₃CN, 2 mM) recorded of an equilibrated 1:2 mixture of **1** and CBPQT⁴⁺, where assignments in green are associated with the [2]pseudorotaxane **1** \subset CBPQT⁴⁺ and assignments in blue are associated with the uncomplexed CBPQT⁴⁺. The region from 1.20 to 1.32 is scaled down.

Oxidation experiments on 1⊂CBPQT⁴⁺ and 2⁴⁺



Fig. S6 Partial ¹H NMR spectra (400 MHz, 298 K, CD₃CN, 2 mM) of **1** and CBPQT⁴⁺ (1:2 ratio), **1** and CBPQT⁴⁺ (1:2 ratio) + ten equiv. TBPASbCl₆ as soon as possible after mixing and **1** with ten equiv. Fe(ClO₄)₃. The areas omitted have no signals. The area from 1.2 to 1.4 ppm is scaled down to clarify the separation and height of the two (CH₃)₃C signals. The CD₂HCN signal as well as the H₂O signals are not shown in their full height. Blue assignments are from free CBPQT⁴⁺ and red assignments are from **1**⊂CBPQT•OP⁶⁺. Only distinct signals are assigned.



Fig. S7 COSY spectrum (400 MHz, 298) recorded as fast as possible after addition of ten equiv. TBPASbCl₆ to a solution of $1 \subset CBPQT^{4+}$ in CD₃CN (*c* = 2 mM) resulting in $1 \subset CBPQT \cdot OP^{6+}$. The structure shown is the part of the stopper which is encircled by CBPQT⁴⁺.



Fig. S8 COSY spectrum (400 MHz, 298 K) as fast as possible after addition of ten equiv. TBPASbCl₆ to a solution in CD₃CN of 2^{4+} (c = 2 mM) resulting in 2•OP ⁶⁺. The structure shown is the part of the stopper encircled by CBPQT⁴⁺.



Fig. S9 Partial ¹H NMR spectra (400 MHz, 298 K, CD₃CN) of 2^{4+} (2 mM) + tenequiv. TBPASbCl₆ recorded 25, 45 and 97 h after mixing and of $6 + CBPQT^{4+}$ (1:1, 2 mM) + ten equiv. TBPASbCl₆. The (CH₃)₃C, CD₂HCN, ⁺NCH₂ and xylyl-H are not shown in full height. The blue assignments are from free CBPQT⁴⁺ and the black assignments are from 2^{4+} .



Fig. S10 Partial ¹H NMR spectra (400 MHz, 298 K, CD₃CN) of 2^{4+} (c = 2 mM with 0.2 M NH₄PF₆) + ten equiv. TBPASbCl₆ recorded 1, 6, 23 and 48 h after mixing. The (CH₃)₃C, CD₂HCN, N⁺CH₂, NH₄⁺ and xylyl-H from free CBPQT⁴⁺ signals are not shown in their full height. The assigned resonances are all from 2^{6+} or residual solvent peaks. The complete assignment can be found in Fig. S11.

It is evident that all four spectra in Fig. S10 are almost identical. By integrating the signals from the phenylene group in the bulky stopper unit and comparing the integrals to those of the tetramethylsilane(TMS) signal, it becomes clear that they are not changing over time. This observation indicates that 2^{6+} is stable for more than 48 h in the presence of 100 equiv. NH₄PF₆ at 298 K and that CBPQT⁴⁺ does not deslip (Scheme S6).



Scheme S6 Movement mechanism in the [2]rotaxane 2•MPTTF⁴⁺ after addition of ten equiv. TBPASbCl₆.

Determining the size of the MPTTF²⁺/SMe/TDEG barrier in 1⊂CBPQT⁶⁺



Fig. S11 Partial ¹H NMR spectra (400 MHz, 298 K) of $1 \subset CBPQT^{4+}$ produced by mixing **1** (2 mM) and CBPQT⁴⁺ (4 mM) in CD₃CN and then allowed to equilibrate for 10 h in the presence of NH₄PF₆ (0.2 M) followed by oxidation with ten equiv. TBPASbCl₆. The spectra are recorded at different delay times after addition of TBPASbCl₆. The areas omitted have no signals or only contain the residual solvent peak. The area from 1.2 to 1.4 ppm is scaled down to clarify the separation and height of the two (CH₃)₃C signals. The NH₄ signal as well as the H₂O signal are not shown in their full height. Assignments in blue are from 1_CCBPQT⁴⁺, assignments in red are from 1_CCBPQT•OP⁶⁺ and assignments in black are from 1²⁺.

The stability of $1 \subset CBPQT^{6+}$ over time is poor. To keep it stable all experiments are performed with the presence of 0.2 M NH₄PF₆, which is found to increase the stability of oxidised MPTTF.

Because it is possible to follow the movement of CBPQT⁴⁺ from the position around the OP station in $1 \subset CBPQT \cdot OP^{6+}$ and over the MPTTF²⁺/SMe/TDEG barrier, it is possible to qualitatively determine the size of the combined MPTTF²⁺/SMe/TDEG barrier. The experiments shown in Fig. S11 were repeated, but spectra were recorded more frequently (every hour for three days) to get enough data points to calculate the rate of CBPQT⁴⁺ deslipping from $1 \subset CBPQT^{6+}$. The experiments were repeated at 294, 303, 308, 313 and 323 K, and the first ¹H NMR spectrum was recorded as soon as possible after addition of ten equiv. TBPASbCl₆ to an equilibrated 1:2 mixture of the semidumbbell **1** and CBPQT⁴⁺ in CD₃CN with NH₄PF₆ (0.2 M). By treating the deslipping as a first order process, it is possible to use first order kinetics (ln *I* as a function of time, *t*, Fig. S12) to determine the rate constant k_d for the movement of CBPQT⁴⁺.



Fig. S12 The natural logarithm of the integral of the singlets (294 K) resonating at 1.29 and 1.37 ppm, respectively, as a function of time. The red data points are used to make a linear regression, from which the slope and thereby the rate constant of CBPQT⁴⁺ deslipping over the MPTTF²⁺/SMe/TDEG barrier can be found.

Table S3 Rate constants, k_d , energies of activation, ΔG^{\ddagger}_d and correlation coefficients, R^2 , from CBPQT⁴⁺ deslipping over the MPTTF²⁺/SMe/TDEG barrier measured at 294 K in CD₃CN (0.2 M NH₄PF₆) by following the process with ¹H NMR spectroscopy (500 MHz), using different signals as probes. Errors are calculated based on Koumura *et al.*^{S4} with $\Delta T = 0.2$ K and $\Delta I = 0.05\%$

Assignment	δ/ppm	<i>k</i> _d / s ⁻¹	ΔG^{\ddagger}_{d} / kcal mol ⁻¹	R^2
2 ²⁺	1.29	1.24 × 10 ⁻⁶ ± 2.78 × 10 ⁻¹⁰	25.17 ± 0.02	0.995
2⊂CBPQT•OP ⁶⁺	1.37	$4.73 \times 10^{-7} \pm 7.12 \times 10^{-11}$	25.73 ± 0.02	0.989
2 ²⁺	3.21	$1.62 \times 10^{-6} \pm 3.67 \times 10^{-9}$	25.02 ± 0.02	0.973
2⊂CBPQT•OP ⁶⁺	3.24	$3.98 \times 10^{-7} \pm 7.59 \times 10^{-10}$	25.83 ± 0.02	0.977
CBPQT ⁴⁺	7.54	$3.71 \times 10^{-7} \pm 1.87 \times 10^{-10}$	25.88 ± 0.02	0.979

Table S4 Rate constants, k_d , energies of activation, ΔG^{\ddagger}_d and correlation coefficients, R^2 , from CBPQT⁴⁺ deslipping over the MPTTF²⁺/SMe/TDEG barrier measured at 303 K in CD₃CN (0.2 M NH₄PF₆) by following the process with ¹H NMR spectroscopy (500 MHz), using different signals as probes. Errors are calculated based on Koumura *et al.*^{S4} with $\Delta T = 0.2$ K and $\Delta I = 0.05\%$

Assignment	δ/ppm	<i>k</i> _d / s ⁻¹	ΔG^{\ddagger}_{d} / kcal mol ⁻¹	R^2
2 ²⁺	1.29	$3.12 \times 10^{-6} \pm 4.14 \times 10^{-10}$	25.40 ± 0.02	0.982
2⊂CBPQT•OP ⁶⁺	1.37	$9.40 \times 10^{-7} \pm 1.22 \times 10^{-10}$	26.13 ± 0.02	0.969
2 ²⁺	3.21	5.56 × 10 ⁻⁶ ± 7.67 × 10 ⁻⁹	25.06 ± 0.02	0.978
2⊂CBPQT•OP ⁶⁺	3.24	8.20 × 10 ⁻⁷ ± 1.29 × 10 ⁻⁹	26.21 ± 0.02	0.942
CBPQT ⁴⁺	7.54	$8.31 \times 10^{-7} \pm 2.62 \times 10^{-10}$	26.20 ± 0.02	0.943

Table S5 Rate constants, k_d , energies of activation, ΔG^{\ddagger}_d and correlation coefficients, R^2 , from CBPQT⁴⁺ deslipping over the MPTTF²⁺/SMe/TDEG barrier measured at 308 K in CD₃CN (0.2 M NH₄PF₆) by following the process with ¹H NMR spectroscopy (500 MHz), using different signals as probes. Errors are calculated based on Koumura *et al.*^{S4} with $\Delta T = 0.2$ K and $\Delta I = 0.05\%$

Assignment	δ/ppm	<i>k</i> _d / s ⁻¹	ΔG^{\ddagger}_{d} / kcal mol ⁻¹	R^2
2 ²⁺	1.29	$3.47 \times 10^{-6} \pm 1.64 \times 10^{-9}$	25.77 ± 0.02	0.998
2⊂CBPQT•OP ⁶⁺	1.37	$1.08 \times 10^{-6} \pm 6.94 \times 10^{-10}$	26.48 ± 0.02	0.957
2 ²⁺	3.21	5.51 × 10 ⁻⁶ ± 2.23 × 10 ⁻⁸	25.48 ± 0.02	0.978
2⊂CBPQT•OP ⁶⁺	3.24	1.24 × 10 ⁻⁶ ± 7.01 × 10 ⁻⁹	26.40 ± 0.02	0.956
CBPQT ⁴⁺	7.54	1.67 × 10 ⁻⁶ ± 1.21 × 10 ⁻⁹	26.22 ± 0.02	0.985

Table S6 Rate constants, k_d , energies of activation, ΔG^{\ddagger}_d and correlation coefficients, R^2 , from CBPQT⁴⁺ deslipping over the MPTTF²⁺/SMe/TDEG barrier measured at 313 K in CD₃CN (0.2 M NH₄PF₆) by following the process with ¹H NMR spectroscopy (500 MHz), using different signals as probes. Errors are calculated based on Koumura *et al.*^{S4} with $\Delta T = 0.2$ K and $\Delta I = 0.05\%$

Assignment	δ/ppm	<i>k</i> _d / s ⁻¹	ΔG^{\ddagger}_{d} / kcal mol ⁻¹	R^2
2 ²⁺	1.29	6.23 × 10 ⁻⁶ ± 7.67 × 10 ⁻¹⁰	25.83 ± 0.02	0.995
2⊂CBPQT•OP ⁶⁺	1.37	$2.14 \times 10^{-6} \pm 2.89 \times 10^{-10}$	26.50 ± 0.02	0.961
2 ²⁺	3.21	1.02 × 10 ⁻⁵ ± 1.20 × 10 ⁻⁸	25.53 ± 0.02	0.990
2⊂CBPQT•OP ⁶⁺	3.24	$2.36 \times 10^{-6} \pm 3.18 \times 10^{-9}$	26.43 ± 0.02	0.972
CBPQT ⁴⁺	7.54	$2.53 \times 10^{-6} \pm 5.55 \times 10^{-10}$	26.39 ± 0.02	0.986

Table S7 Rate constants, k_d , energies of activation, ΔG^{\ddagger}_d and correlation coefficients, R^2 , from CBPQT⁴⁺ deslipping over the MPTTF²⁺/SMe/TDEG barrier measured at 323 K in CD₃CN (0.2 M NH₄PF₆) by following the process with ¹H NMR spectroscopy (500 MHz), using different signals as probes. Errors are calculated based on Koumura *et al.*^{S4} with $\Delta T = 0.2$ K and $\Delta I = 0.05\%$

Assignment	δ/ppm	<i>k</i> _d / s ⁻¹	ΔG^{\ddagger}_{d} / kcal mol ⁻¹	R^2
2 ²⁺	1.29	1.19 × 10 ⁻⁵ ± 1.88 × 10 ⁻⁹	26.26 ± 0.02	0.901
2⊂CBPQT•OP ⁶⁺	1.37	$9.20 \times 10^{-6} \pm 1.47 \times 10^{-9}$	26.43 ± 0.02	0.931
2 ²⁺	3.21	1.19 × 10 ⁻⁵ ± 3.10 × 10 ⁻⁸	26.26 ± 0.02	0.879
2⊂CBPQT•OP ⁶⁺	3.24	$8.90 \times 10^{-6} \pm 1.60 \times 10^{-8}$	26.45 ± 0.02	0.872
CBPQT ⁴⁺	7.54	$9.09 \times 10^{-6} \pm 2.04 \times 10^{-9}$	26.43 ± 0.02	0.960

An average value of all five results (Table S3–S7) was used to calculate the final rate constants for CBPQT⁴⁺ deslipping over the MPTTF²⁺/SMe/TDEG barrier The results obtained at the five different temperatures are listed in Table S8.

Table S8 Average rate constants and ΔG^{\ddagger} values from CBPQT⁴⁺ deslipping over the MPTTF²⁺/SMe/TDEG barrier measured at different temperatures in CD₃CN (0.2 M NH₄PF₆) by following the process with ¹H NMR spectroscopy (500 MHz). Errors are calculated based on Koumura et al^{S4} with ΔT = 0.2 K and ΔI = 0.05%

<i>T /</i> K	n	<i>k</i> _d / s ⁻¹	ΔG^{\ddagger}_{d} / kcal mol ⁻¹
294	51	$8.20 \times 10^{-7} \pm 9.93 \times 10^{-10}$	25.53 ± 0.02
303	45	2.25 × 10 ⁻⁶ ± 1.95 × 10 ⁻⁹	25.80 ± 0.02
308	24	$2.59 \times 10^{-6} \pm 6.57 \times 10^{-9}$	26.07 ± 0.02
313	22	$4.69 \times 10^{-6} \pm 3.36 \times 10^{-9}$	26.14 ± 0.02
323	8	$1.02 \times 10^{-5} \pm 1.05 \times 10^{-8}$	26.37 ± 0.02

By plotting the derived activation energies (ΔG^{\ddagger}_d) as a function of temperature (*T*), it is possible to determine the enthalpy (ΔH^{\ddagger}_d) and the entropy (ΔS^{\ddagger}_d) according to the equation $\Delta G^{\ddagger}_d = \Delta H^{\ddagger}_d - T \times \Delta S^{\ddagger}_d$. The plot is shown in Fig. S13 and the kinetic parameters are summarized in Table S9.



Fig. S13 The average activation energy (ΔG^{\ddagger}_d) plotted against the temperature (*T*) in order to find the enthalpy (ΔH^{\ddagger}_d) and the entropy (ΔS^{\ddagger}_d). The slope and intercept of the linear regression give the values – ΔS^{\ddagger}_d and ΔH^{\ddagger}_d , respectively from the equation $\Delta G^{\ddagger}_d = \Delta H^{\ddagger}_d - T \times \Delta S^{\ddagger}_d$.

Table S9 Kinetic parameters for the movement of CBPQT⁴⁺ from the OP station over the MPTTF²⁺/SMe/TDEG barrier in 2²⁺ obtained by ¹H NMR spectroscopy (500 MHz, CD₃CN, 0.2 M NH₄PF₆) at 298 K

R^2	ΔG^{\ddagger}_{d} / kcal mol ^{-1 a}	ΔH^{\ddagger}_{d} / kcal mol ^{-1 a}	$\Delta S^{\ddagger}_{ m d}$ / cal mol ⁻¹ K ⁻¹ a
0.968	25.7 ± 0.1	15.5 ± 5.2	-33.9 ± 16.8

^{*a*} Errors are calculated using Koumura *et al.*^{S4} with $\Delta K = 0.2$ K.



Fig. S14 ¹H NMR spectrum (400 MHz, 298K, CD₃CN) of the semidumbbell 1.



Fig. S15 ¹³C NMR spectrum (100 MHz, 298K, CDCl₃) of the semidumbbell 1.



Fig. S16 1 H NMR spectrum (400 MHz, 298K, CD₃CN) of the dumbbell 6.



Fig. S17 ¹³C NMR spectrum (100 MHz, 298K, CDCI₃) of the dumbbell 6.



Fig. S18 ¹H NMR spectrum (400 MHz, 298K, CD₃CN) of the [2]rotaxane 2•4PF₆.



Fig. S19 ¹³C NMR spectrum (100 MHz, 298K, CD₃CN) of the [2]rotaxane 2·4PF₆.



Fig. S20 ^1H NMR spectrum (400 MHz, 298K, CD_3CN) of the model compound 9.



Fig. S21 ^{13}C NMR spectrum (100 MHz, 298K, CD_3CN) of the model compound 9.

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