

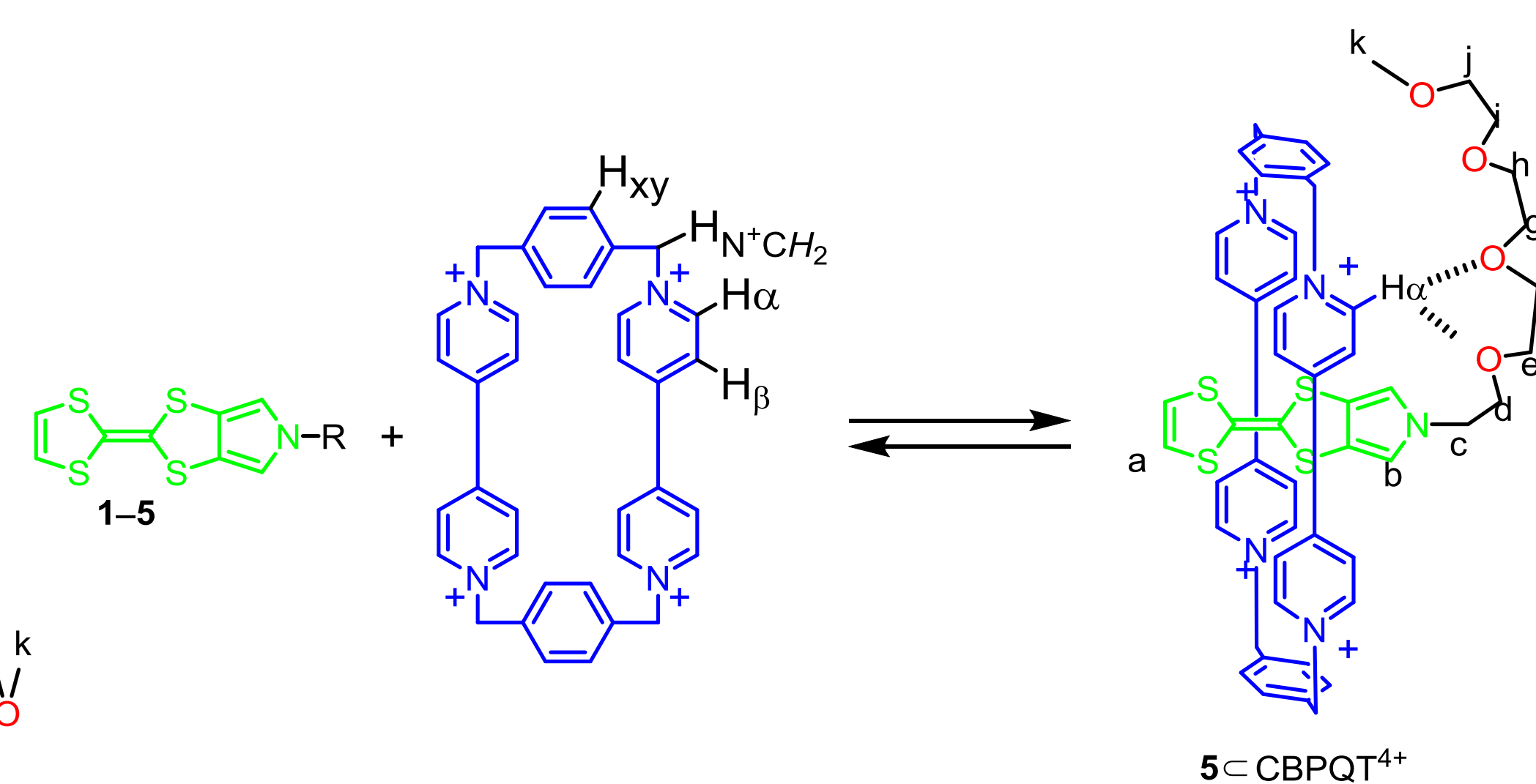
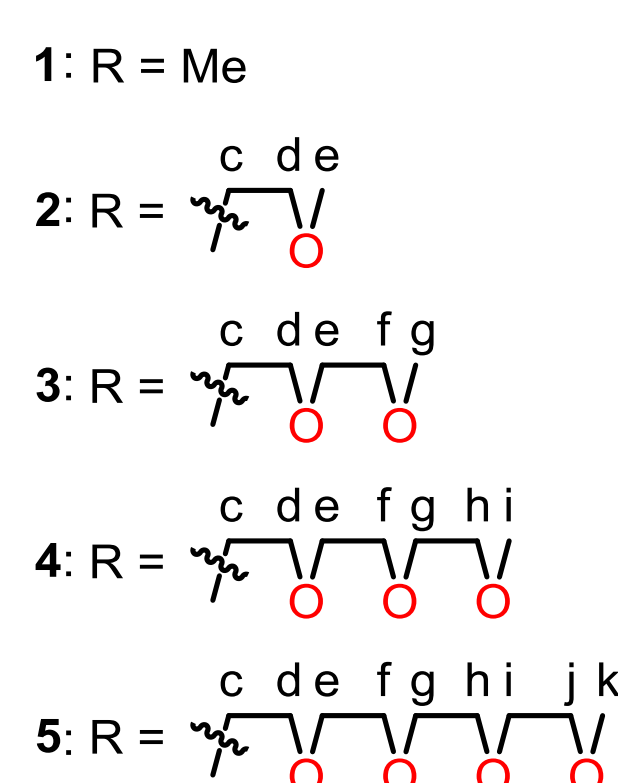
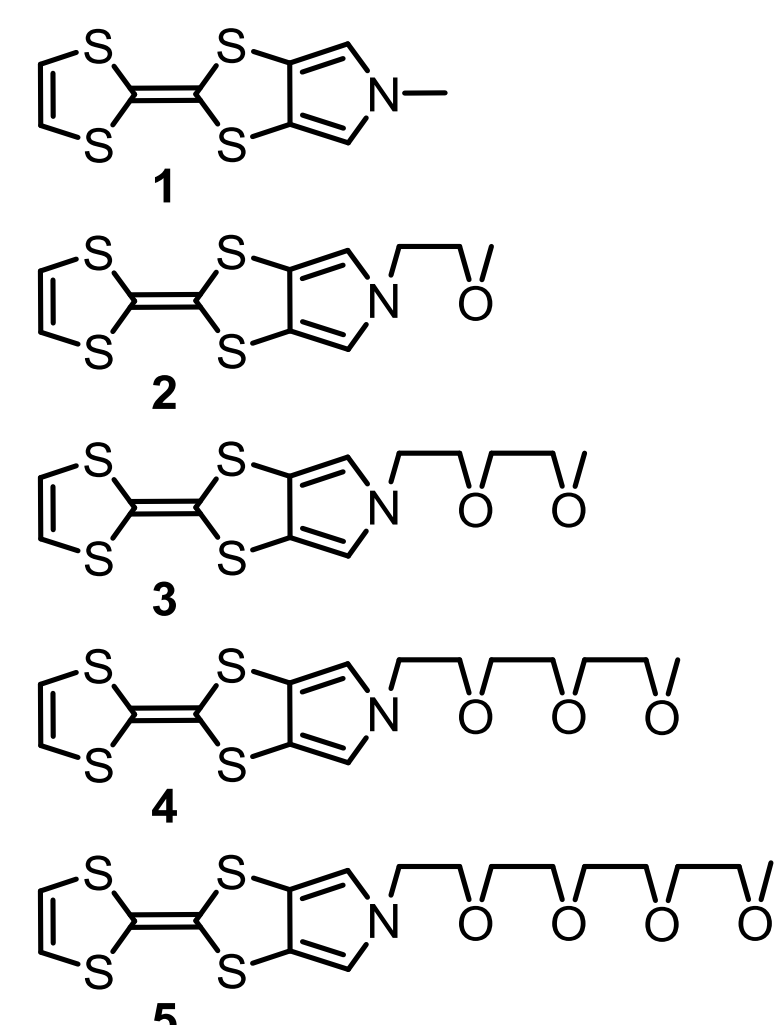
Probing the Role of Glycol Chain Lengths in π -Donor-Acceptor [2]Pseudorotaxanes Based on Monopyrrolo-Tetrathiafulvalene and Cyclobis(paraquat-*p*-phenylene)

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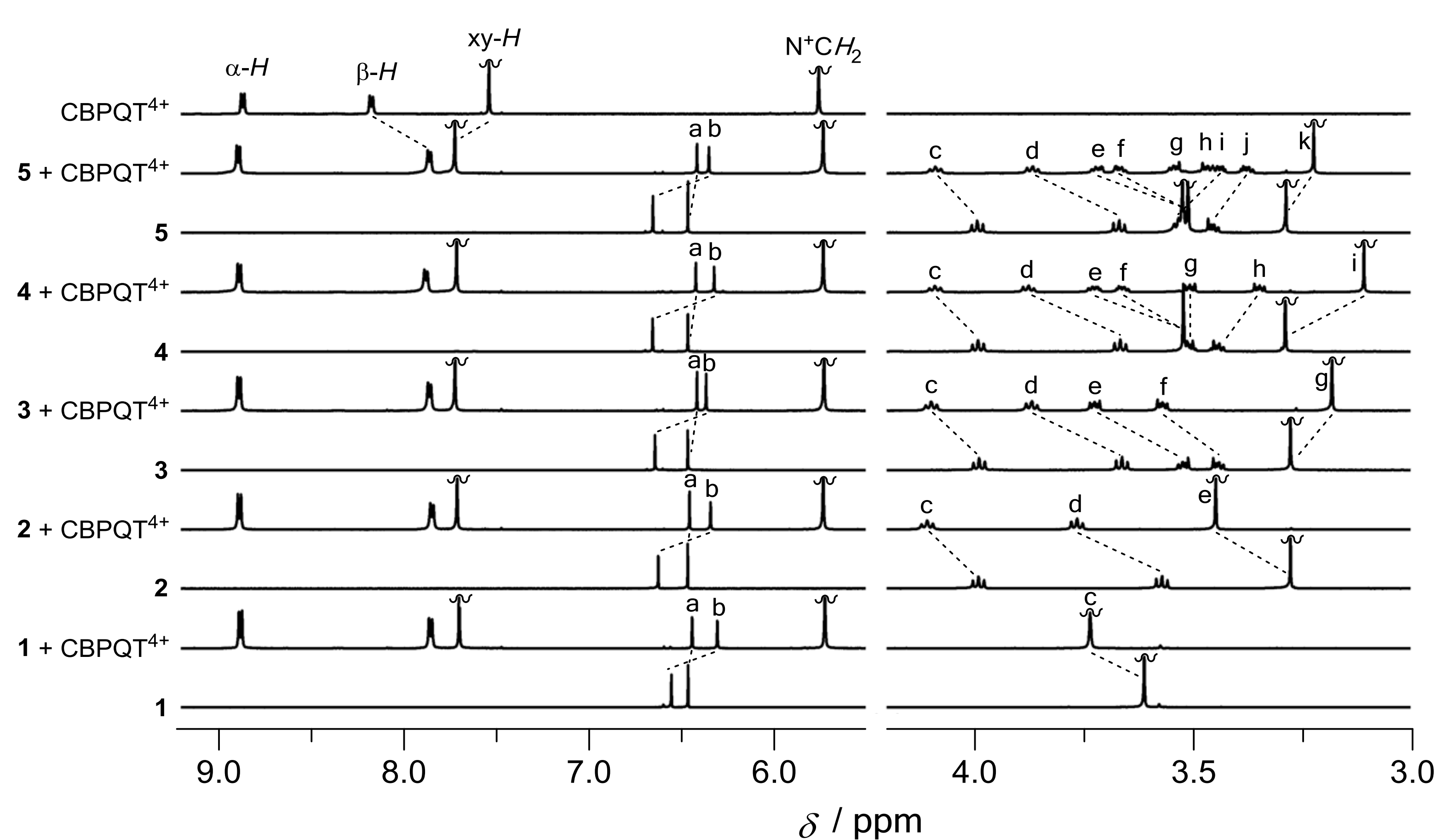


Aim of the Project

A homologous series of *N*-substituted monopyrrolo-tetrathiafulvalenes (MPTTF), only differing in the number of ethyleneoxy units were synthesized. These formed [2]pseudorotaxanes upon mixing with cyclobis(paraquat-*p*-phenylene) (CBPQT⁴⁺). The resulting [2]pseudorotaxanes were investigated in order to probe the role of the glycol chain lengths on the complexation between 1–5 and CBPQT⁴⁺. [1]

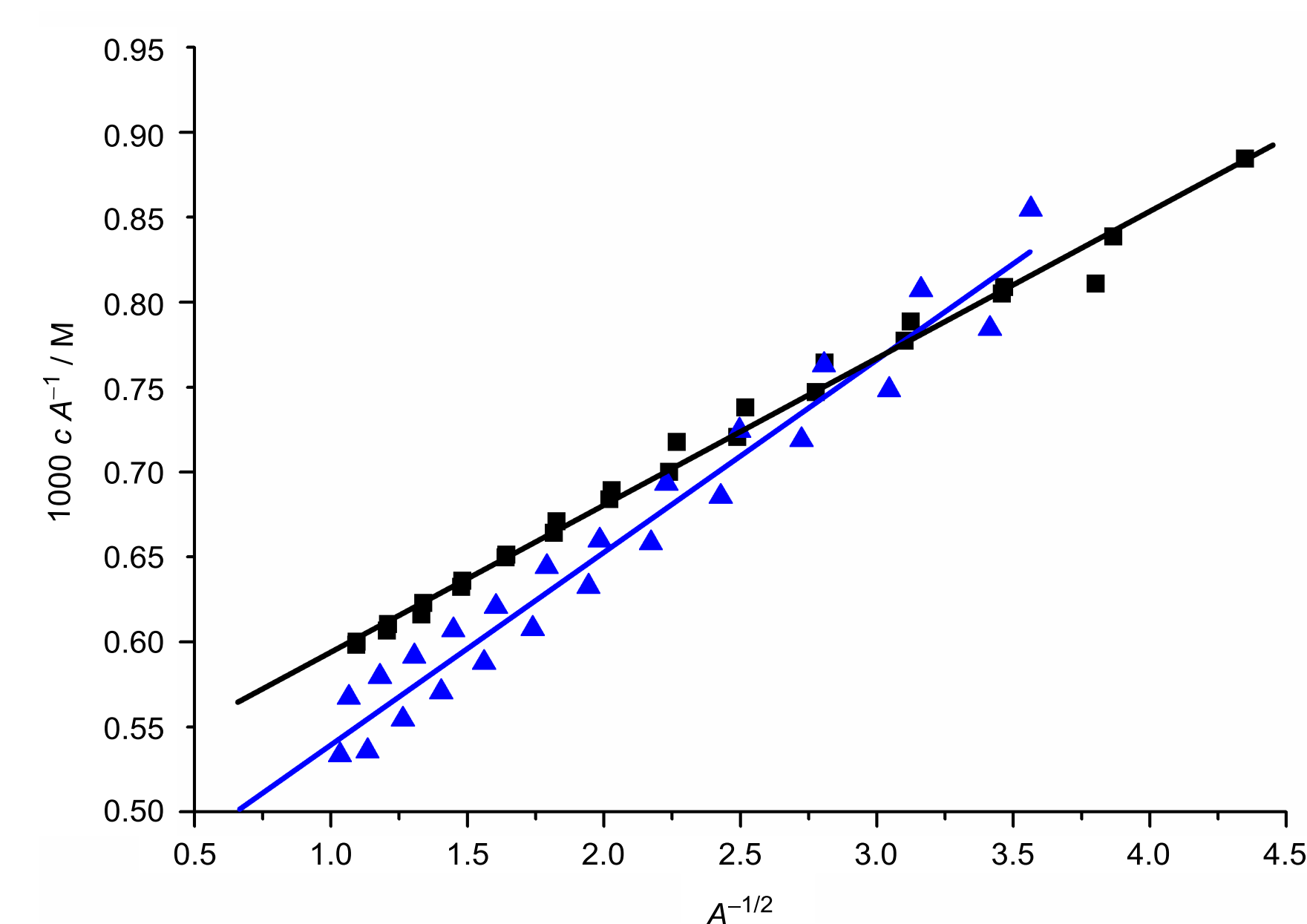


¹H NMR Spectroscopic Studies



Partial ¹H NMR spectra (400 MHz) recorded in CD₃CN (*c* = 1.5 mM) at 298 K of CBPQT⁴⁺, the MPTTF derivatives 1–5, and equilibrated solutions containing 1–5 mixed with equimolar amounts of CBPQT⁴⁺.

UV-Vis-NIR Studies

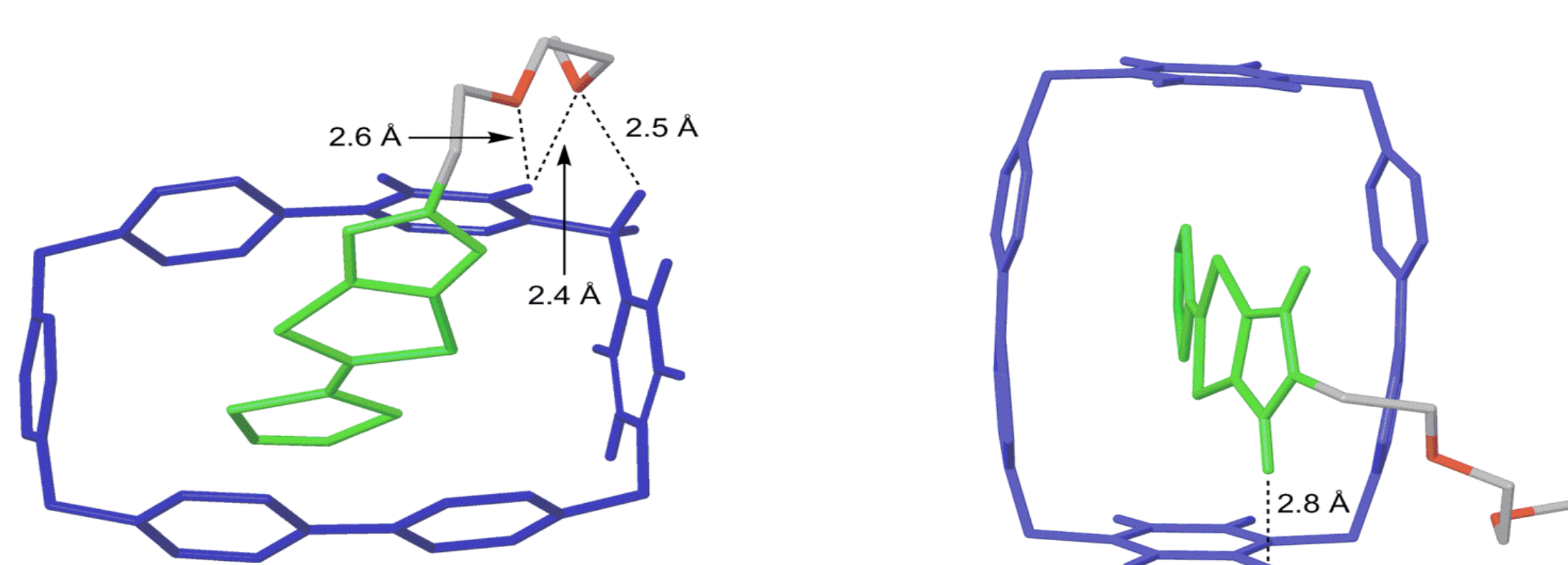


Linear plots of $1000 c A^{-1}/M$ against $A^{-1/2}$ for a 1:1 mixture of CBPQT⁴⁺ and MPTTF derivative 1 (blue line) and 3 (black line).

Binding Constants (K_a Values) and Derived Free Energies of Complexation (ΔG°) between the MPTTF Derivatives 1–5 and CBPQT⁴⁺ Determined by UV-Vis-NIR Absorption Spectroscopy at 298 K in MeCN.

compound	λ_{max} (nm)	data points	correlation coefficient	K_a (M ⁻¹)	ΔG° (kcal mol ⁻¹)
1	877	24	0.960	33000 ± 4000	-6.2 ± 0.1
2	854	28	0.987	41000 ± 4000	-6.3 ± 0.1
3	854	27	0.991	68000 ± 8000	-6.6 ± 0.1
4	856	28	0.986	64000 ± 8000	-6.6 ± 0.1
5	854	29	0.999	63000 ± 7000	-6.6 ± 0.1

Molecular Modelling and Cyclic Voltammetry



M06-L/6-31G** geometry optimized superstructure of the [2]pseudorotaxane 3-CBPQT⁴⁺ (solvent corrected, MeCN) shown at two different angles. The MPTTF unit is colored green and the CBPQT⁴⁺ ring is blue, whereas carbon and oxygen atoms in the glycol chain are colored gray and red, respectively.

The first ($E_{1/2}^1$) and second ($E_{1/2}^2$) redox potentials for the MPTTF derivatives 1–5 obtained by cyclic voltammetry (CV) at 298 K in MeCN.

Compound	$E_{1/2}^1$ (V)	$E_{1/2}^2$ (V)
1	+0.030	+0.400
2	+0.025	+0.415
3	+0.020	+0.415
4	+0.020	+0.415
5	+0.020	+0.415

Conclusions

- The MPTTF-derivatives 1–5 show similar redox potentials, so the change in binding constants is not ascribed to come from a change in electron donor properties
- The [2]pseudorotaxanes 1–5-CBPQT⁴⁺ show a trend towards higher binding energy, the longer the glycol chain. The biggest change in binding energy is seen for compounds 2 and 3
- Investigations suggest that hydrogen bonds can form between the first and second oxygen atoms and CBPQT⁴⁺ in the [2]pseudorotaxanes, providing an extra stabilization of $\Delta G^\circ = -0.3 \text{ kcal mol}^{-1}$ to the [2]pseudorotaxanes

