

pubs.acs.org/OrgLett



CB[7]- and CB[8]-Based [2]-(Pseudo)rotaxanes with Triphenylphosphonium-Capped Threads: Serendipitous Discovery of a New High-Affinity Binding Motif

Iago Neira, Carlos Peinador,* and Marcos D. García*



binding motif has been extensively studied experimentally, with results in good agreement with

echanically interlocked molecules (MIMs)^{1,2} are no Monger a chemical curiosity but a solid platform for the development of new functionality.³ Nevertheless, the efficient synthesis of these entities is still challenging, especially when restricted to aqueous media and the limited choice of intermolecular interactions and reactivity that can be used in this setting.⁴ Cucurbit[n]urils $(CB[n]s, n = 5-8, 10, 13-15)^5$ have substantially eased this problem, enlarging the toolbox for the preparation of MIMs in aqueous media,⁶ in particular, when using well-developed synthetic strategies that employ premade axle components (e.g., capping⁷ or slipping⁸). Watersoluble CB[n]s are commercially available, nontoxic, and fairly nonreactive and own flexible inner hydrophobic cavities of different sizes, and their host-guest chemistry is mainly directed by cation-dipole interactions,⁹ the hydrophobic effect¹⁰ and optimization of the host-guest packing coefficient.^{11,12} Furthermore, in the case of CB[8], its polar and large hydrophobic cavity, 1.7 times the volume of CB[7], allows for the preparation of unusual 1:2 heteroternary complexes with aromatic guests of complementary electron acceptor/donor nature, which are stabilized within the cavity by increased charge-transfer interactions.¹³

those obtained by dispersion-corrected DFT methods.

Following our interest in the chemistry of $CB[n]s^{14}$ and pyridinium salts,¹⁵ we designed the synthetic strategy for the construction of CB[7]- and CB[8]-based asymmetric [2]rotaxanes depicted in Scheme 1c, having as key steps (a) the threading of CB[7] and CB[8] into complementary triphenylphosphonium-capped semidumbbells $1_a^{2+}/1_b^{2+}$ and (b) trapping of the interlocked molecule by an unusual kinetically controlled imine bonding reaction recently developed by our group.¹⁶ Scheme 1. (a) Synthesis of Thread 2^{4+} ; (b) Schematic Representation of CB[7] and CB[8]; and (c) Planned Synthesis of Rotaxanes $2^{4+} \subset CB[7]$ and $2^{4+} \subset CB[8]$ by Simultaneous Threading–Capping



Building blocks $1_a \cdot 2Br$ and $1_b \cdot 2Br$ were efficiently prepared by substitution reactions of commercially available (4-(bromomethyl)benzyl)triphenylphosphonium bromide with, respectively, isonicotinaldehyde and 4-hydrazinylpyridine.¹⁷

Received: March 24, 2022 Published: May 6, 2022



© 2022 The Authors. Published by American Chemical Society We then tested the assembly of the axle component 2^{4+} (Scheme 1) by reacting 1_a^{2+} and 1_b^{2+} for 18 h in refluxing water and in the presence of a catalytic amount of TFA. The process produced the expected thread, which was isolated after column chromatography as $2 \cdot 4Cl$ in 69% yield and fully characterized by means of ${}^{31}P/{}^{1}H/{}^{13}C$ 1D/2D NMR and ESI-MS.¹⁷ As previously demonstrated for related hydrazones,¹⁶ cation 2^{4+} displayed an abnormal stability, with no signs of hydrolysis being detected over a period of weeks.¹⁷ Having the assembled thread in our hands, we tested a slipping approach for the synthesis of $2^{4+} \subset CB[7]^8$ by heating equimolar 1.0 mM mixtures of $2 \cdot 4Cl$ and CB[7] in D₂O. This procedure failed to produce the MIM, validating therefore the use of the triphenylphosphonium groups as suitable stoppers for CB[7]-based rotaxanes.^{16c}

As originally intended, we alternatively undertook the synthesis of $2^{4+} \subset CB[7]$ starting from the host and the individual components 1_a^{2+} and 1_b^{2+} (Scheme 1). Nevertheless, we first studied the CB[7]-guest complexation processes by following by ${}^{1}H/{}^{31}P$ NMR the reactions between 1.0 mM solutions in D₂O, of either 1_a^{2+} or 1_b^{2+} and increasing amounts of the host.¹⁷ In both cases, the data recorded were in good agreement with the formation of the expected 1:1 host–guest complexes $1_a^{2+}/1_b^{2+} \subset CB[7]$, which appear in the spectra in a situation of rapid exchange in the NMR time scale. Further 1D/2D NMR and HR ESI-MS experiments additionally validated the formation of the pseudorotaxanes, and $K_a = 1.46 \pm 0.07 \times 10^5 M^{-1}$ could be estimated for $1_b^{2+} \subset CB[7]$ through an UV–vis titration experiment, which showed a good fitting of the data to a 1:1 binding isotherm (Figure 1c).

Surprisingly, despite the semidumbbells having xylylenebased hydrophobic cores flanked with two positive charges, the binding sites observed in both cases for the CB[7] host were not these moieties^{14c} but, instead, the corresponding pyridinium rings. For instance, the ¹H NMR signals corresponding to the pyridinium ring in 1_a^{2+} appear significantly deshielded ($\Delta \delta H_f = 0.62$ ppm and $\Delta \delta H_e = 0.63$ ppm, Figure 1a,b), while $H_{b,c}$ within the xylylene core are only slightly altered. Furthermore, the ³¹P NMR signal for 1_a^{2+} was also downshifted $\Delta \delta P_a = 0.73$ ppm, in good agreement with the discussed insertion mode.¹⁷ Dispersion-corrected calculations carried out for $1_a^{2+} \subset CB[7]$ supported this end,¹⁷⁻²¹ being the local minimum m_1 found for the insertion of the pyridyl group within CB[7], 2.1 kcal/mol more stable than that corresponding to the inclusion of the xylylene moiety (m_2 , Figure 2d).

Once both the formation of $\mathbf{1}_{a}^{2+}/\mathbf{1}_{b}^{2+}\subset CB[7]$ and the axle 2^{4+} were firmly established, we undertook the synthesis of the [2]rotaxane $2^{4+} \subset CB[7]$. Hence, the very same conditions used for the synthesis of 2.4Cl were applied, but in the presence of 2 equiv of CB[7]. After 18 h, the expected CB[7]-based MIM was isolated as $2.4PF_6 \subset CB[7]$ in 55% yield, being thoroughly characterized by ³¹P/¹³C/¹H NMR and ESI-MS (Figure 2b).¹⁷ Additionally, an analytical sample of 2.4ClCCB[7] could be obtained by anion metathesis,¹⁷ allowing the characterization of the [2]rotaxane in aqueous media by 1D/2D NMR techniques. In particular, the ¹H spectrum in D₂O showed an appropriate integration of the asymmetric protons for CB[7] in relation with the axle nuclei, with chemical shifts for H_{f-h} consistent with the expected shielding caused by the positioning of the central bis-pyridinium moiety within the host (Figure 2c). This later observation was also found in good agreement with the DFT-optimized geometry obtained for a





Figure 1. Partial ¹H NMR (500 MHz, D₂O) spectrum of (a) 1 mM solution of $\mathbf{1}_{a}^{2+}$ and (b) 1 mM of $\mathbf{1}_{a}^{2+} + 1$ equiv of CB[7]. (c) Partial UV–vis and titration data (inset) for $\mathbf{1}_{b}^{2+}$ (15 μ M) in the presence of increasing concentrations of CB[7] in water. The red line in the inset shows the fit to a 1:1 binding model. (d) Schematic representation of the two energy minima \mathbf{m}_{1} and \mathbf{m}_{2} found for $\mathbf{1}_{a}^{2+} \subset CB[7]$ using DFT methods.^{17–19}



Figure 2. Rotaxane $2^{4+} \subset CB[7]$: (a) DFT-optimized geometry; (b) LR ESI-MS spectrogram for the hexafluorophosphate salt; (c) partial ¹H NMR (500 MHz, D₂O) of (top) axle 2^{4+} and (bottom) $2^{4+} \subset CB[7]$.

local minimum of the [2]rotaxane (Figure 2a).¹⁷⁻²¹ DOSY NMR also supported the formation of the MIM, with all of the resonances for the compound diffusing as a whole on the recorded spectrum (see Figure S5).

Continuing with our study, we then tackled the synthesis of the CB[8] analogue of the [2]rotaxane, first by employing again a slipping approach between thread 2^{4+} and the host. In this case, sonication of a 2.0 mM solution of 2^{4+} in D₂O with excess macrocycle led to unexpected results: not only the relative integration of the CB[8] signals on the ¹H NMR showed two units of the host strongly interacting with the axle, but the whole species displayed a sole diffusion coefficient on the corresponding DOSY experiment (Figure 3c). Assignment



Figure 3. Pseudo[3]rotaxane $2^{4+} \subset (CB[8])_2$: (a) schematic representation; (b) LR ESI-MS spectrogram for the chloride salt; (c) partial ¹H NMR (500 MHz, D₂O) of (top) axle 2^{4+} , (middle) 2 mM 2^{4+} + 2 equiv of CB[8], and (bottom) DOSY experiment for the precedent solution.

of the ¹H nuclei could be carried out based on 2D NMR experiments,¹⁷ as $H_{a-d,i-m}$ on the xylene moieties consistently shielded compared to the free axle, in good agreement with their inclusion within the hydrophobic cavity of CB[8]. Moreover, the signals for the six phenyl rings on the two nonequivalent $R-P^+Ph_3$ groups on 2^{4+} appeared, in turn, divided into two very differently affected sets of signals on the ¹H spectrum, slowly exchanging on the NMR time scale. In essence, 20 hydrogens accounting for two of the phenyl rings on each moiety appear slightly deshielded, as would be expected for an interaction of those with the outer rims of the CB[8]s. On the other hand, the remaining 10 nuclei $(H_{x-z/x'-z'})$ appear significantly shielded by ca. 1 ppm, a situation that would imply their surprising inclusion within the cavity of the macrocycles forming an heteroternary binding motif in conjunction with the xylyl moiety. In contraposition with what discussed above for $2^{4+} \subset CB[7]$, the chemical shifts observed in this case for the nuclei of the pseudoviologen moiety appear more erratically affected by CB[8], as it would derive from their positioning outside of the host. Moreover, ³¹P NMR showed as well data consistent with the binding of the host to the phosphonium ends of the nonsymmetric axle $(\Delta\delta P = -1.61 \text{ and } -1.77 \text{ ppm})$. Finally, ESI-MS experiments

could be recorded for the pseudorotaxane $2 \cdot 4 \text{Cl} \subset (\text{CB}[8])_2$, showing a clear peak for the proposed complex $2^{4+} \subset (\text{CB}[8])_2$ at m/z 897.0462 (calcd 897.0461) (Figure 3b).

To further validate our hypothesis of the CB[8] host introducing within its cavity two of the four aromatic rings of the benzyltriphenylphosphonium moiety, we proceed to study the interaction of $\mathbf{1}_{a}^{2+}$ and $\mathbf{1}_{b}^{2+}$ with the macrocycle in aqueous media. Hence, we recorded the ¹H NMR spectrum of equimolar 2 mM solutions of the guests, saturating with excess CB[8] by sonication.¹⁷ These experiments showed clear indications of interaction between the components, showing nearly the same patterns of integration, splitting of the CB[8] signals and complexation induced shifts, discussed above for the pseudo[3]rotaxane $2^{4+} \subset (CB[8])_2$. HR ESI-MS experiments further confirmed the identity of the $l_a^{2+}/l_b^{2+} \subset CB[8]$ 1:1 complexes, showing diagnostic peaks for $[1_a^{2+} \subset CB[8]]^{2+}$ at m/z 909.7973 (calcd 909.7965), and for $[\mathbf{1}_{b}^{2+} \subset CB[8]]^{2+}$ at m/z 901.8046 (calcd 901.8064). Interestingly, 2D ROESY NMR experiment recorded for $l_a^{2+}/l_b^{2+} \subset CB[8]$ allowed us to obtain further information, as we observed three EXSY exchange peaks associated to each of the nonequivalent protons of the phenyl groups, implying its slow exchange in and *out* of the macrocycle in the NMR time scale (Figure 4).



Figure 4. Partial 2D ROESY NMR (500 MHz, D_2O) showing EXSY exchange cross peaks between phenyl groups in the complex $l_a^{2+} \subset CB[8]$.

Consequently, the energy barrier for the exchange (ΔG^{\ddagger}) could be estimated from VT-NMR experiments on both $\mathbf{1}_{a}^{2+}/\mathbf{1}_{b}^{2+}\subset CB[8]$ (see Figures S8 and S9), showing similar values of approximately 15.0 kcal/mol. Finally, the strength of the association could be estimated in the case of $\mathbf{1}_{a}^{2+}\subset CB[8]$, by using NMR competitive experiments with (trimethyl-ammonium)methylferrocene as a standard for the calculation $(\log K_{a} = 9.49).^{4,17}$ Thus, the host–guest interaction was observed to be stronger than that of the standard, and that the process was under no kinetic barriers, showing a quite impressive binding constant $K_{a} = (3.6 \pm 0.7) \cdot 10^{10} \text{ M}^{-1}$,

comparable to that found to other high-affinity substrates such as adamantane derivatives. $^{\rm 22}$

To gain more insight on the structural characteristics of these atypical heteroternary complexes, DFT calculations were carried out on $1_a^{2+} \subset CB[8]$. Among the four local minima found on the potential energy surface for the pseudorotaxane $(m'_{1-4}, Figure 5)$,¹⁷ the two lowest energy conformers show



Figure 5. Schematic representation of the four energy minima m'_1 - m'_4 found for complexes $1_a^{2+} \subset CB[8]$ using DFT methods.¹⁷⁻¹⁹

the proposed heteroternary binding mode, differing on the syn- $(\mathbf{m'}_1)$ or *anti*- $(\mathbf{m'}_2)$ relative disposition of the pyridinium moiety and the inserted phenyl ring, and being separated by $\Delta E = Em'_1 - Em'_2 = 7.7$ kcal/mol. Furthermore, the two other minima identified for the complex display the same binding modes discussed before for $\mathbf{1}_a^{2+} \subset CB[7]$, but being in this case significantly higher in relative energy ($\Delta E = Em'_1$ - $Em'_{4} = 22.7 \text{ kcal/mol}, \Delta E = Em'_{1} - Em'_{3} = 19.3 \text{ kcal/mol}.$ As can be seen in Figure 5, not only is the cavity of the macrocycle able to accommodate one of the phenyl rings and the benzyl group of 1_a^{2+} but also this insertion mode significantly optimizes the occupied volume with respect to the large volume of the CB[8] cavity with a p.c. of 58%, very close to the optimal occupancy value.^{12,13} Furthermore, the two positive charges of the binding motif in $\mathbf{m'}_{1,2}$ (N⁺ and P⁺) are located approximately in the center of the polygons defined by the oxygen atoms flanking the CB[8] portals, somehow optimizing as well the cation-dipole interactions.

In summary, we have discussed here our new results on the synthesis of tryphenylphosphonium-capped CB[7]- and CB[8]-based rotaxanes. The intended strategy has been studied in detail, planned to concomitantly produce CB[7] and CB[8] complexes with complementary aldehyde and hydrazone reactive ends, and their covalent irreversible junction by imine bond formation of the thread component on the MIM. While the method efficiently produced the expected CB[7]-based [2]-rotaxane, the use of CB[8] as a wheel component resulted in the formation of unexpected pseudo-heteroternary complexes between the host and the

capped ends of the thread. This unusual result has been studied in depth, with the experimental data and dispersioncorrected DFT calculations being in excellent agreement with the proposed binding mode. The reported results open the door not only for the development of new water-soluble rotaxanes but also for the design of new high-affinity binding guests for CB[8] inspired by the pseudo-heteroternary complexation mode discussed herein.^{22,23}

ASSOCIATED CONTENT

Supporting Information

The Supporting Information is available free of charge at https://pubs.acs.org/doi/10.1021/acs.orglett.2c01028.

Experimental procedures, characterization data for new compounds, additional figures and computational details including the Cartesian coordinates for the DFT structures discussed in the text (PDF)

AUTHOR INFORMATION

Corresponding Authors

- Carlos Peinador Departamento de Química and Centro de Investigaciones Científicas Avanzadas (CICA). Facultad de Ciencias, Universidade da Coruña, 15071 A Coruña, Spain; orcid.org/0000-0001-5823-6217; Email: carlos.peinador@udc.es
- Marcos D. García Departamento de Química and Centro de Investigaciones Científicas Avanzadas (CICA). Facultad de Ciencias, Universidade da Coruña, 15071 A Coruña, Spain; orcid.org/0000-0002-3189-740X; Email: marcos.garcia1@udc.es

Author

 Iago Neira – Departamento de Química and Centro de Investigaciones Científicas Avanzadas (CICA). Facultad de Ciencias, Universidade da Coruña, 15071 A Coruña, Spain;
 orcid.org/0000-0002-1297-6938

Complete contact information is available at: https://pubs.acs.org/10.1021/acs.orglett.2c01028

Author Contributions

All authors have given approval to the final version of the manuscript.

Funding

Funding for open access charge: Universidade da Coruña/ CISUG.

Notes

The authors declare no competing financial interest.

ACKNOWLEDGMENTS

This research was supported by the Agencia Estatal de Investigación (PID2019-105272GB-I00) and Xunta de Galicia (ED431C 2018/39). I.N. thanks the MECD (FPU program). The authors are indebted to Centro de Supercomputación de Galicia (CESGA) for providing the computer facilities.

REFERENCES

 Bruns, C. J.; Stoddart, J. F. The Nature of the Mechanical Bond: From Molecules to Machines; John Wiley & Sons: Hoboken, NJ, 2016.
 Xue, M.; Yang, Y.; Chi, X.; Yan, X.; Huang, F. Development of Pseudorotaxanes and Rotaxanes: From Synthesis to StimuliResponsive Motions to Applications. Chem. Rev. 2015, 115, 7398-7501.

(3) For selected recent reviews on rotaxanes and rotaxane-based functionality see: (a) Jamieson, E. M. G.; Modicom, F.; Goldup, S. M. Chirality in rotaxanes and catenanes. Chem. Soc. Rev. 2018, 47, 5266-5311. (b) Mena-Hernando, S.; Pérez, E. M. Mechanically interlocked materials. Rotaxanes and catenanes beyond the small molecule. Chem. Soc. Rev. 2019, 48, 5016-5032. (c) Takata, T. Switchable Polymer Materials Controlled by Rotaxane Macromolecular Switches. ACS Cent. Sci. 2020, 6, 129-143. (d) Zhou, H.-Y.; Zong, Q.-S.; Han, Y.; Chen, C.-F. Recent advances in higher order rotaxane architectures. Chem. Commun. 2020, 56, 9916-9936. (e) Schroeder, H. V.; Schalley, C. A. Electrochemically switchable rotaxanes: recent strides in new directions. Chem. Sci. 2019, 10, 9626-9639. (f) McGonigal, P. R. Multiply threaded rotaxanes. Supramol. Chem. 2018, 30, 782-794. (4) Oshovsky, G. V.; Reinhoudt, D. N.; Verboom, W. Supramolecular Chemistry in Water. Angew. Chem., Int. Ed. 2007, 46, 2366-2393.

(5) Barrow, S. J.; Kasera, S.; Rowland, M. J.; del Barrio, J.; Scherman, O. A. Cucurbituril-based molecular recognition. *Chem. Rev.* 2015, 115, 12320–12406.

(6) (a) Han, Z.; Zhou, Q.; Li, Y. Self-assembled (pseudo)rotaxane and polyrotaxane through host-guest chemistry based on the cucurbituril family. J. Incl. Phenom. Macrocycl. Chem. 2018, 92, 81– 101. (b) Kim, K. Mechanically interlocked molecules incorporating cucurbituril and their supramolecular assemblies. Chem. Soc. Rev. 2002, 31, 96–107. (c) Hou, X.; Ke, C.; Stoddart, J. F. Cooperative capture synthesis: yet another playground for copper-free click chemistry. Chem. Soc. Rev. 2016, 45, 3766–3780.

(7) Wu, C.; Lecavalier, P. R.; Shen, Y. X.; Gibson, H. W. Synthesis of a Rotaxane via the Template Method. *Chem. Mater.* **1991**, *3*, 569–572.

(8) Asakawa, M.; Ashton, P. R.; Ballardini, R.; Balzani, V.; Bělohradský, M.; Gandolfi, M. T.; Kocian, O.; Prodi, L.; Raymo, F. M.; Stoddart, J. F.; Venturi, M. The Slipping Approach to Self-Assembling [n]Rotaxanes. J. Am. Chem. Soc. **1997**, 119, 302–310.

(9) Mock, W. L.; Shih, N.-Y. Structure and Selectivity in Host-Guest Complexes of Cucurbituril1. J. Org. Chem. **1986**, 51, 4440–4446.

(10) Biedermann, F.; Uzunova, V. D.; Scherman, O. A.; Nau, W. M.; De Simone, A. Release of High-Energy Water as an Essential Driving Force for the High-Affinity Binding of Cucurbit[n]urils. *J. Am. Chem. Soc.* **2012**, *134*, 15318–15323.

(11) The well-known Rebek's "55% solution" establishes the correlation between optimized host-guest binding and the packing coefficient that is the ratio of the guest to host volume and is typically in the range of 0.55 ± 0.09 : Mecozzi, S.; Rebek, J., Jr. The 55% Solution: A Formula for Molecular Recognition in the Liquid State. *Chem.*—*Eur. J.* **1998**, *4*, 1016–1022.

(12) Values of the optimal packing coefficients for CB[n]s have been found to be in good agreement with the above-mentioned "55% solution" rule:¹¹ Nau, W. M.; Florea, M.; Assaf, K. I. Deep Inside Cucurbiturils: Physical Properties and Volumes of their Inner Cavity Determine the Hydrophobic Driving Force for Host–Guest Complexation. *Isr. J. Chem.* **2011**, *51*, 559–577.

(13) Biedermann, F.; Scherman, O. A. Cucurbit[8]uril Mediated Donor–Acceptor Ternary Complexes: A Model System for Studying Charge-Transfer Interactions. J. Phys. Chem. B 2012, 116, 2842–2849.

(14) (a) Pazos, E.; Novo, P.; Peinador, C.; Kaifer, A. E.; García, M. D. Cucurbit[8]uril (CB[8])-Based Supramolecular Switches. Angew. Chem., Int. Ed. 2019, 58, 403-416. (b) Neira, I.; García, M. D.; Peinador, C.; Kaifer, A. E. Terminal Carboxylate Effects on the Thermodynamics and Kinetics of Cucurbit[7]uril Binding to Guests Containing a Central Bis(Pyridinium)-Xylylene Site. J. Org. Chem. 2019, 84, 2325-2329. (c) Neira, I.; Domarco, O.; Barriada, J. L.; Franchi, P.; Lucarini, M.; García, M. D.; Peinador, C. An electrochemically controlled supramolecular zip tie based on host-guest chemistry of CB[8]. Org. Biomol. Chem. 2020, 18, 5228-5233.
(d) Novo, P.; García, M. D.; Peinador, C.; Pazos, E. Reversible Control of DNA Binding with Cucurbit[8]uril-Induced Supra-

molecular 4,4'-Bipyridinium-Peptide Dimers. *Bioconjugate Chem.* **2021**, 32, 507–511. (e) Neira, I.; García, M. D.; Peinador, C.; Kaifer, A. E. Cucurbiturils as Effectors on the Self-Assembly of Pd(II) and Pt(II) Metallacycles. *J. Org. Chem.* **2021**, *86*, 14608–14616.

(15) Neira, I.; Blanco-Gómez, A.; Quintela, J. M.; García, M. D.; Peinador, C. Dissecting the "Blue Box": Self-Assembly Strategies for the Construction of Multipurpose Polycationic Cyclophanes. *Acc. Chem. Res.* **2020**, *53*, 2336–2346.

(16) (a) Blanco-Gómez, A.; Fernández-Blanco, A.; Blanco, V.; Rodríguez, J.; Peinador, C.; García, M. D. Thinking Outside the "Blue Box": Induced Fit within a Unique Self-Assembled Polycationic Cyclophane. J. Am. Chem. Soc. 2019, 141, 3959–3964. (b) Blanco-Gómez, A.; Neira, I.; Barriada, J. L.; Melle-Franco, M.; Peinador, C.; García, M. D. Thinking outside the "Blue Box": from molecular to supramolecular pH-responsiveness. Chem. Sci. 2019, 10, 10680– 10686. (c) Neira, I.; Blanco-Gómez, A.; Quintela, J. M.; Peinador, C.; García, M. D. Adjusting the Dynamism of Covalent Imine Chemistry in the Aqueous Synthesis of Cucurbit[7]uril-based [2]Rotaxanes. Org. Lett. 2019, 21, 8976–8980.

(17) See the Supporting Information for details.

(18) The dispersion-corrected BLYP-D3(BJ) functional, in combination with the valence double- ζ size polarization basis set def2-SVP, was used for the energy minimizations reported in this work,¹⁷ as it has been recently proven to afford reliable optimized geometries for CB[n]-based complexes at a reasonable computational cost.¹⁹ Truhlar's SMD model was used to account for solvation effects in water,²⁰ and the quadruple- ζ valence quality basis set def2-QZVP was used in single-point calculations to more accurately evaluate the energy differences between isomeric species.²¹

(19) (a) Hostaš, J.; Sigwalt, D.; Šekutor, M.; Ajani, H.; Dubecký, M.; Řezáč, J.; Zavalij, P. Y.; Cao, L.; Wohlschlager, C.; Mlinarić-Majerski, K.; Isaacs, L.; Glaser, R.; Hobza, P. A Nexus between Theory and Experiment: Non-Empirical Quantum Mechanical Computational Methodology Applied to Cucurbit[n]uril·Guest Binding Interactions. *Chem.—Eur. J.* **2016**, *22*, 17226. (b) Fianchini, M.; Llorens, L.; Pericàs, M. A. Separating Enthalpic, Configurational, and Solvation Entropic Components in Host–Guest Binding: Application to Cucurbit[7]uril Complexes through a Full In Silico Approach via Water Nanodroplets. *J. Phys. Chem. B* **2020**, *124*, 10486–10499.

(20) Marenich, A. V.; Cramer, C. J.; Truhlar, D. G. Universal Solvation Model Based on Solute Electron Density and on a Continuum Model of the Solvent Defined by the Bulk Dielectric Constant and Atomic Surface Tensions. *J. Phys. Chem. B* 2009, *113*, 6378–6396.

(21) Weigend, F.; Ahlrichs, R. Balanced basis sets of split valence, triple zeta valence and quadruple zeta valence quality for H to Rn: Design and assessment of accuracy. *Phys. Chem. Chem. Phys.* **2005**, *7*, 3297–3305.

(22) Liu, S.; Ruspic, C.; Mukhopadhyay, P.; Chakrabarti, S.; Zavalij, P. Y.; Isaacs, L. The Cucurbit[n]uril Family: Prime Components for Self-Sorting Systems. J. Am. Chem. Soc. 2005, 127, 15959–15967.

(23) Houk, K. N.; Leach, A. G.; Kim, S. P.; Zhang, X. Binding Affinities of Host-Guest, Protein-Ligand, and Protein-Transition-State Complexes. *Angew. Chem., Int. Ed.* **2003**, *42*, 4872–4897.