

Cite this: *Chem. Sci.*, 2018, 9, 1050

Binding of anions in triply interlocked coordination catenanes and dynamic allostery for dehalogenation reactions†

Linlin Yang,[‡] Xu Jing,[‡] Bowen An, Cheng He,[✉] * Yang Yang and Chunying Duan[✉] *

By synergistic combination of multicomponent self-assembly and template-directed approaches, triply interlocked metal organic catenanes that consist of two isolated chirally identical tetrahedrons were constructed and stabilized as thermodynamic minima. In the presence of suitable template anions, the structural conversion from the isolated tetrahedral conformers into locked catenanes occurred *via* the cleavage of an intrinsically reversible coordination bond in each of the tetrahedrons, followed by the reengineering and interlocking of two fragments with the regeneration of the broken coordination bonds. The presence of several kinds of individual pocket that were attributed to the triply interlocked patterns enabled the possibility of encapsulating different anions, allowing the dynamic allostery between the unlocked/locked conformers to promote the dehalogenation reaction of 3-bromocyclohexene efficiently, as with the use of dehalogenase enzymes. The interlocked structures could be unlocked into two individual tetrahedrons through removal of the well-matched anion templates. The stability and reversibility of the locked/unlocked structures were further confirmed by the catching/releasing process that accompanied emission switching, providing opportunities for the system to be a dynamic molecular logic system.

Received 17th September 2017
Accepted 30th November 2017

DOI: 10.1039/c7sc04070a

rsc.li/chemical-science

Introduction

Biomolecular machines are fascinating systems in which integrated functions are controlled in a synergistic manner. The constitution of the individual components within the interconnected networks governs their capacity to perform specific functions and to interact with other relevant species in the recognition mediated systems.^{1–5} Inspired by sophisticated structures and mechanisms in biological systems, molecular recognition and self-assembly have been exploited as efficient driving forces for the fabrication of artificial molecular systems that operate far from equilibrium in a preorganized manner through well-controlled molecular design.^{6,7} Of these reported artificial systems, molecules containing mechanical bonds, such as catenanes,^{8–10} knots¹¹ and rotaxanes,^{12–14} have captured the attention of the scientific community because of their intriguing architectures and topologies as well as the ability of their components to undergo controllable intra-molecular movements.

A template-directed approach that preorganizes building blocks into entwined or threaded systems by non-covalent interactions was developed to precisely control the architectures and topologies of the mechanically interlocked molecules.^{15–18} For more complex molecules with non-trivial topologies that rely on dynamic bonds, new strategies are required to ensure that the systems undergo ‘error checking’ reactions to obtain their thermodynamic minima.^{19–22} In these cases, their intrinsic natures, including well-defined geometries and fast exchange kinetics, allow metal–ligand interactions to act as promising dynamic bonds for the creation of interlocked structures that reproduce key structural or functional aspects of natural systems and their artificial analogues.^{23,24} However, compared to the clear understanding of template-directed approaches for the creation of mechanically interlocked systems, the requirements for integrating the self-replication of machines with recognition mediated processes relevant to the revisable locked/unlocked architectures and dynamic control of their optical, electronic and transport properties remain largely unexplored.^{25–27}

On the other hand, enzymes accelerate biochemical reactions by providing scaffolds to recognize and activate substrates and stabilize transition states between reactants and products.^{28,29} It is postulated that strongly halophilic molecular knots can promote anion abstraction reactions, echoing the properties of dehalogenase enzymes which contain halide-binding sites that facilitate the cleavage of carbon–halogen

State Key Laboratory of Fine Chemicals, College of Zhang Dayu, Dalian University of Technology, Dalian, 116024, P. R. China. E-mail: cyduan@dlut.edu.cn

† Electronic supplementary information (ESI) available: Characterization data and additional tables and figures. CCDC 1515722 and 1515723. For ESI and crystallographic data in CIF or other electronic format see DOI: 10.1039/c7sc04070a

‡ These authors contributed equally.



bonds.³⁰ The pathway involving substrate capture surmounts the kinetic barriers of the reaction through a “mild” equilibrium control method. Multiply interlocked catenanes exhibit high stabilities and have the potential to form different kinds of pocket that encapsulate several kinds of guest with different sizes and shapes.^{31–36} In these cases, positively charged, multiply coordinated interlocked structures are promising promoters to selectively and simultaneously bind a large number of guest anions.^{37–39} Inspired by these systems, a new approach based on interlocked catenane was developed to realize the dehalogenation process, simulating the mechanism in dehalogenation enzymes.

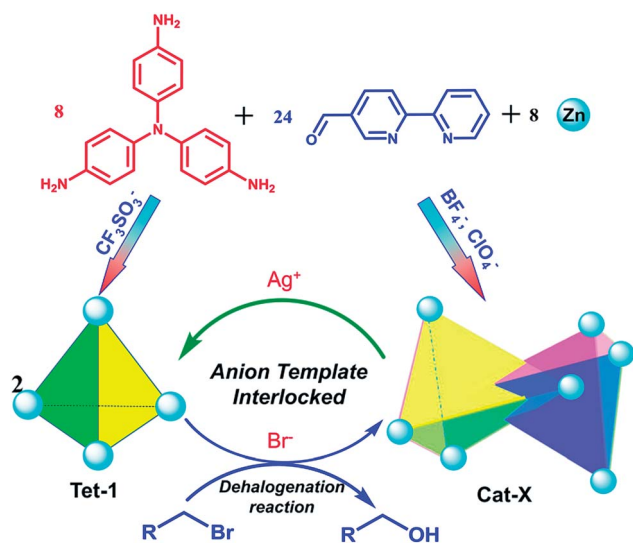
Herein, through the synergistic combination of multi-component self-assembly with a template-directed approach, we report a new approach to construct triply interlocked coordination catenanes from 40 components. The one-pot assembly reaction was well controlled from several types of commercially available fragment^{40–43} including a tris(4-amino phenyl)amine (TPA) group extended by a 2,2'-bipyridine-5-carbaldehyde chelator and the dynamics active zinc(II) ions. These triply interlocked catenanes could also be formed and unlocked into individual tetrahedrons upon the addition or removal of an anion template with a well-matched size and conformational structure. Additionally, the large inner pocket and the multiply interlocked fashion lead to a cavity that is divided into several individual pockets, enabling the encapsulation of different kinds of anion and promoting the dehalogenation reaction of 3-bromo-cyclohexene (Scheme 1).

Results and discussion

The reaction of tris(4-aminophenyl)amine (TPA, 8 equiv.) with 2,2'-bipyridine-5-carbaldehyde (24 equiv.) and $\text{Zn}(\text{BF}_4)_2$ (8 equiv.) provided the complex Cat-BF_4 with a 60% yield. The ESI-

MS spectrum revealed a series of m/z species corresponding to $[\text{Zn}_8\text{L}_8][\text{BF}_4]_n^{(16-n)+}$ (where L is the postulated Schiff-base ligand derived from the condensation of TPA and 2,2'-bipyridine-5-carbaldehyde, Scheme S1; $\dagger n = 7–10$, Fig. S25[†]). The splitting of the signals in the ^{19}F -NMR spectrum of the BF_4^- anions revealed the encapsulation of BF_4^- anions within the molecular architecture (Fig. S3[†]).⁴⁴ The fact that the one-pot self-assembly from 40 individual pieces led to the metal-organic catenane suggested the high thermodynamic stability of Cat-BF_4 in the solution.

X-ray crystallographic analysis of Cat-BF_4 confirmed the formation of the metal-organic catenane and revealed that the triply locked identical cages feature seven pockets, each filled with a BF_4^- anion (Fig. 1). In the asymmetrical unit, four zinc centers with identical Δ or Λ stereochemistries of one tetrahedron are positioned at the terminal apical sites and are bridged by four tripodal ligands that cap the four faces. Within one tetrahedron, the average $\text{Zn}\cdots\text{Zn}$ separation is 19.1 Å, and a tetrahedron has an approximate volume of 820 Å³. Two chirally identical tetrahedrons interlocked in C_2 symmetry with one $\text{Zn}(\text{bipy})_3$ node situated at the center of the internal cavity of the counterpart. Three bidentate chelating arms coordinated to one catenane-coordinated zinc center inter-penetrate across three of the four openings of the counterpart to form a triply interlocked cage.^{45,46}



Scheme 1 The illustration of the reversible assembly and dehalogenation reaction of triply interlocked catenanes via anion template and subcomponent self-assembly.

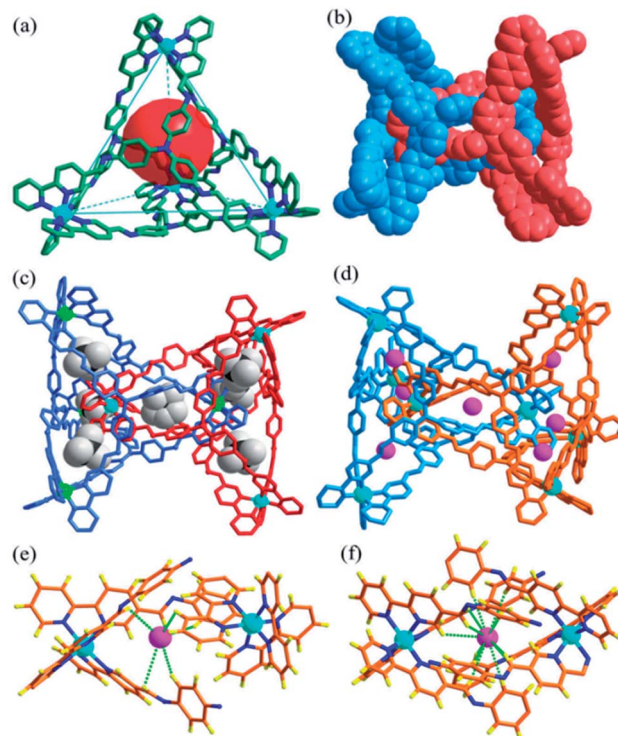


Fig. 1 The structure of the triply interlocked catenane Cat-BF_4 , showing the discrete polyhedrons (a) (the interlocked partner was omitted for clarity) and the space-filling model of the interlocked structure (b). Structures of the interlocked catenanes Cat-BF_4 (c) and Cat-I/BF_4 (d) filled with BF_4^- and I^- anions in the pockets, respectively; and a view of the hydrogen bonding interactions with I^- anions in the outer (e) and central (f) pocket.



The six locked arms face one another to create an inner core – a molecular bowl that is similar to an ‘egg cup’.⁴⁷ One of the BF_4^- anions occupies the center of the molecular bowl. Abundant C–H...F interactions, which stabilize the interlocked structure, are found between the CH=N groups and BF_4^- anions (Fig. S21†). It should also be noted that the large inner pocket and the multiply interlocked fashion lead to a cavity that is divided into 7 individual pockets. Beside the center pocket (inner pocket) that is filled by a BF_4^- anion, the other six pockets (outer pockets) formed each encapsulate a BF_4^- anion. These anions occupy the middle point of the seven shortest separated zinc pairs of the catenane zinc(II) nodes to adhere the charged zinc(II) nodes together through electrostatic interactions and to partly balance the positive charges of the catenane tetrahedrons. These cooperative weak interactions allow the subcomponents of **Cat-BF₄** to recognize each other well and to finely compact in an interlocked nature, benefiting the construction of complicated systems using the same constituent fragments.⁴⁸

Replacing $\text{Zn}(\text{BF}_4)_2$ with $\text{Zn}(\text{CF}_3\text{SO}_3)_2$ under the same assembly conditions gave an isolated tetrahedron **Tet-1**. ESI-MS analysis of the resulting solution of **Tet-1** revealed a family of prominent signals at $m/z = 619.15, 772.77, 1003.2$ and 1387.25 that are assigned to $[\text{Zn}_4\text{L}_4(\text{CF}_3\text{SO}_3)_n]^{(8-n)+}$ ($n = 2, 3, 4$ and 5 , Fig. S27†) species based on a comparison with simulated and natural isotopic abundances (Fig. 4a). The presence of one set of ligand signals in the ^1H NMR (Fig. 2b) and ^{19}F -NMR (Fig. S10†) spectra suggests that the four ligands and eight CF_3SO_3^- anions in **Tet-1** are identical. Important evidence for the interlocked structure of **Cat-BF₄** is provided in the NOESY spectrum (Fig. S5†), where H–H interactions between the bipyridyl protons (H_j) and the **TPA** protons were found. However, no relevant signals were observed for **Tet-1** because of the long distance of these interactions (Fig. S9†). The different self-

assembly products should be impacted by the different sizes and shapes of the template anions.

Generally, the intrinsic reversibility of the coordinated bonds allows the dynamic self-assembly to occur through an ‘error checking’ mode, allowing the system to reach its thermodynamic minimum. Thus, this reversibility allows for the interlocked structure to be assembled from isolated fragments of the components with extra stimulating template anions.^{49,50} Treating the solution containing the isolated tetrahedron **Tet-1** with ClO_4^- at room temperature gave direct evidence for the formation of the inter-locked architecture. As shown in Fig. S49,† the simple and sharp peaks split into several parts corresponding to the loss of symmetry in the interlocked architecture.⁵¹ The yield of the interlocked catenane increased directly with increasing concentration of ClO_4^- (Fig. 3a). This result indicates that the interlocked catenane sites are at the thermodynamic minimum, and could be rapidly transformed if a suitable template anion is present. The moderate and clear formation of **Cat-ClO₄** allowed the conversion process to be monitored in the ^1H -NMR along with the changes of ClO_4^- anions. For **Cat-ClO₄**, the inner binding core contains one rigidly binding perchlorate anion sited in its central pocket. A simple thermodynamics model indicated that two individual tetrahedrons and one ClO_4^- anion are involved, and the mechanism of the conversion process was assumed (see the ESI, Section 5.2†). The well-matched fitting of the ^1H -NMR titration profile generated from the addition of Bu_4NClO_4 at different concentrations to the solution of **Tet-1** (2 mM) with an assumed plot gave an equilibrium constant $K_{\text{ass}}(\text{ClO}_4)$ of $7.0 \times 10^4 \text{ M}^{-2}$ ($\Delta G = -27.63 \text{ kJ mol}^{-1}$) at room temperature (Fig. 3a).

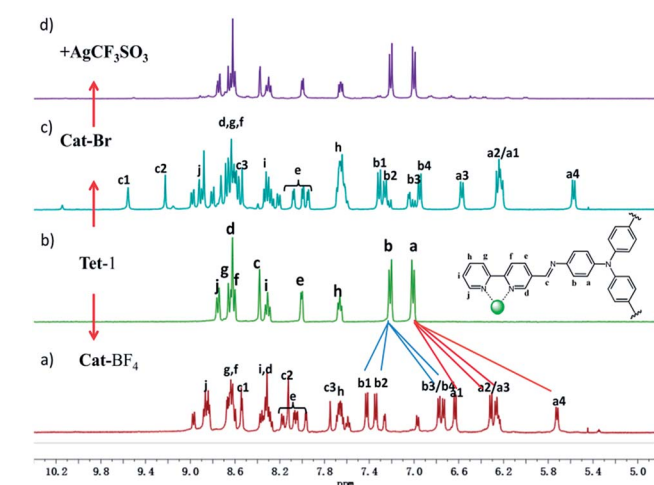


Fig. 2 ^1H -NMR spectra of (a) the interlocked cage **Cat-BF₄** and (b) the monomeric tetrahedron **Tet-1**. ^1H -NMR spectra of the monomeric tetrahedron after addition of 7.0 equiv. Br^- ions (c); followed by subsequent addition of excess (11 equiv.) AgCF_3SO_3 (d) which unlocks the assembly and restores the structure to a discrete tetrahedron.

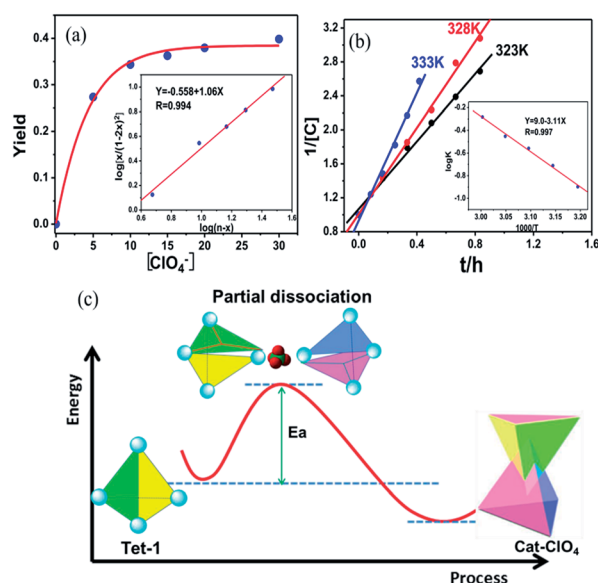


Fig. 3 (a) The fitting curve of the conversion of **Cat-ClO₄** with the addition of different equiv. of ClO_4^- . (b) The kinetic plot fitting for the dimerization process at different temperatures in the presence of 20 equiv. of ClO_4^- . (c) The schematic representation of a putative partial dissociation mechanism for the formation of the interlocked Zn_8L_8 catenane.



Meanwhile, with BF_4^- as the template anion (Fig. S51[†]), both the rate and conversion of **Tet-1** to catenane were quite low compared to those in the presence of ClO_4^- , whereas the addition of the PF_6^- anion could not lead to the formation of the interlocked structure (Fig. S47[†]), further confirming that the size and shape of the template anions were essential to the interlocked system.

The formation of interlocked **Cat-ClO₄** was dynamically active, and the well-fitted linear relationship of $1/[\text{Tet-1}]$ with the reaction time at a fixed temperature suggested a second-order kinetic behaviour (Fig. 3b). The efficient collision between the two tetrahedrons was assumed to be the rate-determining step, which agreed well with the aforementioned thermodynamics assumption for the interlocked transformation. From the determination of kinetic constants at different temperatures, the activation energy of the catenane construction reaction from two isolated tetrahedrons was calculated to be 59.5 kJ mol^{-1} , according to the Arrhenius equation (Table S3[†]). Generally, the interlocked transformation involves the breaking of at least two of the interacting bonds followed by the recovery of these bonds to interlock together. The fact that the activation energy is lower than the cleavage energy of one set of the chelating Zn–N (380 kJ mol^{-1}) coordination bonds⁵² suggests that the structural conversion likely occurs through the random cleavage of one high-probability coordination bond out of the twelve active positions of each tetrahedron, followed by the reengineering of the two colloidal activated tetrahedrons with the formation of new chelating coordination bonds (Fig. 3c).⁵³ Control experiments involving the use of more inert metal ions, with Fe(II) or Co(II) replacing Zn(II) in the subcomponent assembly, gave only the isolated tetrahedron (Fig. S31–S34[†]). The inactivity of the interlocked transformation with the kinetically inert metal ions confirmed that the structural formation was controlled by the partial dissociation dynamics of the Werner-type interactions and the suitable metal–ligand interactions.^{54–56}

The kinetic study suggested that the addition of anions with a similar size which could weakly coordinate to the Zn centers would lower the reaction energy barrier and accelerate the conversion. In this case, the addition of halide anions as NBU_4^+ salts to the solution of **Tet-1** resulted in the quickly exclusive detection of signals corresponding to the interlocked species $[\text{X} \subset \text{Zn}_8\text{L}_8]$ ($\text{X} = \text{I}^-$ and Br^-) and a number of CF_3SO_3^- anions as counter anions in the ESI-MS spectra (Fig. 4). This result demonstrates that the halide anions I^- and Br^- have suitable sizes and trigger hydrogen bonding similarly to BF_4^- . In addition, the driving forces for structure allostery were strong enough to direct the equilibrium controlled transformation to the thermodynamic minimum of the system. When excess Bu_4NBF_4 salt was added to the solution of the interlocked structure $[\text{I}^- \subset \text{Zn}_8\text{L}_8]$, crystals of **Cat-I/BF₄** were obtained and characterized using X-ray diffraction. As shown in Fig. 1d, the cage morphology of **Cat-I/BF₄** is slightly altered in the presence of I^- . Clearly, one I^- anion occupies the ‘egg cup’-like inner core, and six I^- or BF_4^- anions statistically occupy the middle points of the shortest separated zinc pairs to adhere the charged nodes together *via* electrostatic interactions.^{57,58} In this

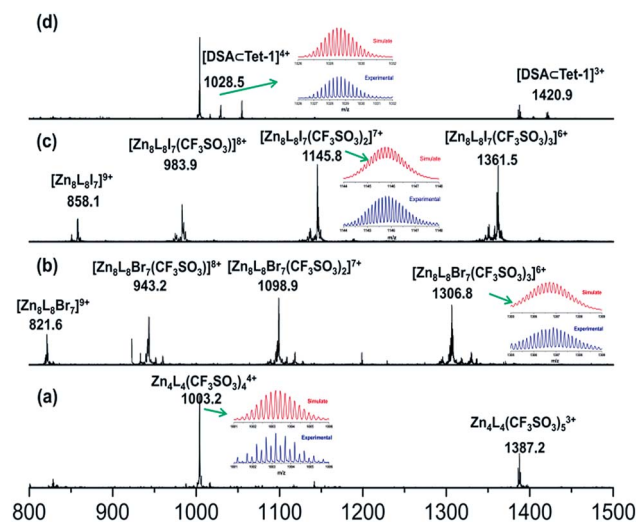


Fig. 4 ESI-MS spectra of the solution containing the isolated tetrahedron **Tet-1** (a). The aforementioned solution upon the addition of (b) Bu_4NBr , (c) Bu_4NI and (d) Bu_4NDSA , showing the formation of interlocked structures with a $7\text{Br}^- \subset \text{Zn}_8\text{L}_8$ core, $7\text{I}^- \subset \text{Zn}_8\text{L}_8$ core and host–guest $\text{DSA} \subset \text{Tet-1}$ species, respectively.

structure, the I^- anion in the central cavity was strongly fixed, while the outer six I^- anions could be randomly exchanged with the added BF_4^- anions. The results showed that there are two kinds of micro-environment in this catenane. The central cavity can bind anions more powerfully than the other six ‘outer’ cavities, which is likely due to the presence of more hydrogen bonding interaction sites (Fig. 1f).

To elucidate the driving force for dimerization and the origin of the thermodynamic stability of the interlocked tetrahedrons, we evaluated the thermodynamic parameters using isothermal titration calorimetry (ITC).⁵⁹ A typical titration curve is shown in Fig. 5a. The observed inclusion number between **Tet-1** and Br^- anions was 3.5, which is in good agreement with the seven Br^- anions in the ESI-MS spectra of **Cat-Br**. Curve fitting by the computer simulation using an ‘independent’ model reveals the activation enthalpy $\Delta H = 8.1 \text{ kJ mol}^{-1}$, the activation entropy $\Delta S = 123.8 \text{ J mol}^{-1}$, and the Gibbs free energy $\Delta G = -28.77 \text{ kJ mol}^{-1}$ (Fig. S69[†]).^{60,61} These results demonstrate that the dimerization of **Tet-1** is an entropy-driven endothermic reaction, which is attributed to the increase of molecular disorder and release of solvents.⁶² Similarly, the ΔG between the dimerization of **Tet-1** with I^- anions is $-30.15 \text{ kJ mol}^{-1}$ (Fig. S70[†]). The sequence of the Gibbs free energy in the dimerization process seems to be ranked as $\text{BF}_4^- < \text{ClO}_4^- < \text{Br}^- < \text{I}^-$.

Since silver salts can remove halide anions from solution *via* the formation of an insoluble AgX precipitate, we identified that the bound halide anions in **Cat-X** could also be precipitated by the addition of AgCF_3SO_3 with the catenane unlocked. As shown in Fig. 2d, the $^1\text{H-NMR}$ titration of AgCF_3SO_3 into a solution containing **Cat-Br** resulted in the deterioration of the splitting of the signals of the interlocked structure. The simple NMR pattern is attributed to the tetrahedron **Tet-1**, and the ESI-MS



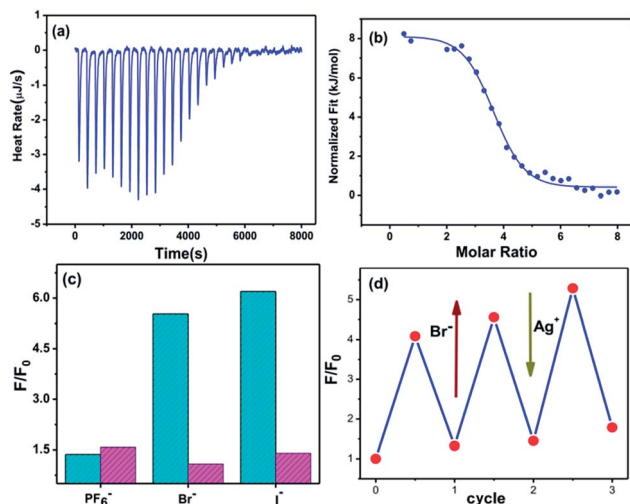


Fig. 5 (a) The isothermal titration calorimetry (ITC) data for sequential injections (10 μL per injection) of Br^- solution (5 mM) into **Tet-1** solution (0.125 mM). (b) The apparent reaction heat obtained from the integration of calorimetric traces. (c) The fluorescence of $\text{DSA} \subset \text{Tet-1}$ species with the addition of 10 equiv. of different template ions (PF_6^- , Br^- and I^- , cyan column) followed by titration of 20 equiv. of AgCF_3SO_3 (red column) to the aforementioned solution (F_0 corresponds to the fluorescence of $\text{DSA} \subset \text{Tet-1}$). (d) The reversible Br^-/Ag^+ controlled uptake and release of DSA monitored by fluorescence.

spectrum (Fig. S42[†]) confirmed the formation of the isolated tetrahedron from the locked/unlocked equilibrium. The conformational flexibility and dynamic reversibility of the interlocked catenane from the isolated tetrahedron with a large inner cavity provided the possibility of introducing photo or electro-active elements to functionalize the transformation processes.^{63,64} Since emission is a sensitive method to recognize responsive behavior, Bu_4NDSA (where **DSA** is dansyl acid), a widely used luminescent indicator, was chosen as a guest to visually show the reversibility of the allostery of the structure. As shown in Fig. 4d, upon the addition of NBu_4DSA , the ESI-MS spectrum of a **Tet-1** solution exhibited new peaks at $m/z = 1028.5$ and 1420.9 that were assigned to $[\text{DSA} \subset \text{Tet-1}]^{4+}$ and $[(\text{DSA} \subset \text{Tet-1})(\text{CF}_3\text{SO}_3)]^{3+}$ species, respectively. The ^1H NMR spectrum of **Tet-1** was largely affected upon the addition of 5 equiv. of **DSA**, demonstrating the formation of the complexes of **DSA** in solution (Fig. S81[†]). Preliminary results also indicated that the formation of the host-guest species $\text{DSA} \subset \text{Tet-1}$ led to a strong decrease in luminescence.⁶⁵

Furthermore, the addition of Br^- and I^- to a solution containing both NBu_4DSA and **Tet-1** resulted in the direct enhancement of the luminescence (Fig. 5c). The recovered fluorescence was quenched and returned to its initial value when Ag^+ ions were added to the solution containing the interlocked structure, and the same cycle could be repeated several times. These results showed that, during the locked/unlocked process, the individual cage as well as the interlocked structure in solution are quite stable and exhibit excellent structure reversibility. The halide-communicated catching/releasing conformational switching resembles a key feature of

dehalogenase enzymes in which the halide anion binding pathway efficiently surmounts the kinetic barriers associated with the carbon-halide bonds through a “mild” equilibrium control method. The locked/unlocked behavior allowed the system to become a dynamic molecular logic system, enabling it to adopt what are otherwise thermodynamically inaccessible functional conformations.

We wondered if such a strong halide allostery process might be applied to the dehalogenation reaction, which has been traditionally performed by halophilic silver salts. We envisaged that the seven pockets in the catenane might be able to promote the formation of a carbocation intermediate by abstracting the Br^- anions from the C-Br bond cleavage reactions.³⁰ In a typical procedure, **Tet-1** was mixed with 3-bromocyclohexene (3.6 μM) in CD_3CN in the presence of 2,4,6-collidine and MeOTf at room temperature, resulting in conversion to 2-cyclohexene-1-ol after stirring for 3 hours. A loading of 10% mole ratio of the **Tet-1** tetrahedron resulted in 65% conversion to 2-cyclohexene-1-ol. The bromide ions bound within the cavities of **Cat-Br** react with methyl triflate to form bromomethane gas. Clearly, the presence of MeOTf restores the formed **Cat-Br** to the unlocked **Tet-1** as shown in Fig. S73.[†] However, **Cat-BF₄** only reacts with a stoichiometric amount of 3-bromo-cyclohexene, and peaks of **Cat-Br** were detected in the dehalogenation of 3-bromo-cyclohexene in the absence of MeOTf (Fig. S90[†]). Control experiments showed that neither the ligand components (**TPA** + 2,2'-bipyridine-5-carbaldehyde), the zinc(II) salt, nor $\text{Zn}(\text{bpy})_3$ could promote this reaction in a decent yield (Table S5[†]).

As shown in Fig. 6, the transformation exhibited a first order kinetic behavior. When the concentration of the 3-bromo-cyclohexene was fixed, the initial reaction rate of the dehalogenation reaction exhibited a linear relationship with the

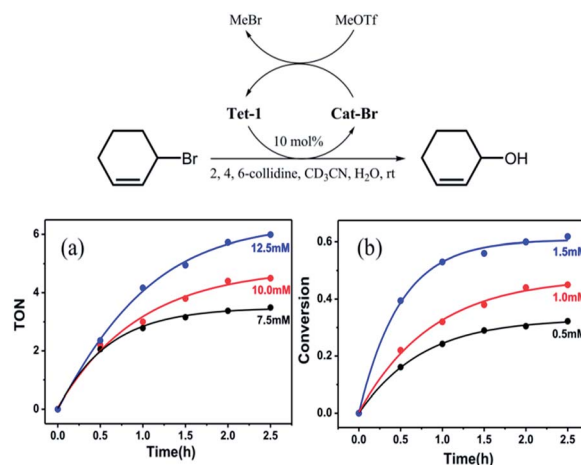


Fig. 6 The kinetics study of the dehalogenation reaction of 3-bromocyclohexene. (a) The variation in turnover number (per mole of the catalyst **Tet-1**) with the concentration of the substrate 3-bromocyclohexene varies in the system containing 2,4,6-collidine (20 mM), MeOTf (20 mM), and **Tet-1** (1 mM), and (b) the conversion of the substrate 3-bromo-cyclohexene (10.0 mM) with the concentration of **Tet-1** varies in the system containing 2,4,6-collidine (20 mM) and MeOTf (20 mM).



concentration of **Tet-1**. When the concentration of **Tet-1** was fixed, the initial turnover frequency of the reaction hardly changed with the concentration of the substrate. It seems that the rate of the reaction only depends on the concentration of the bromide encapsulated interlocked complex. The formation of the bromide encapsulated interlocked complex was possibly the rate-limited step and the dehalogenation reaction took place through the direct SN1 cleavage of the C–Br bond.

Conclusions

In summary, we report a new approach to construct triply interlocked coordination catenanes that consist of two chirally identical tetrahedrons. The new approach included the synergic combination of multi-component self-assembly and molecular recognition, and at the same time also involved seeking the thermodynamic minima of the interlocked/unlocked equilibrium based on the intrinsic dynamic active nature of the coordination assembly. The presence of several kinds of individual pocket that were attributed to the triply interlocked patterns provided the possibility to encapsulate different kinds of anion, allowing the dynamic allostery between the locked/unlocked conformers to efficiently promote the elimination of 3-bromocyclohexene.

Experimental

Preparation of the interlocked tetrahedron Cat-BF₄ [Zn₈L₈] [BF₄]₁₆

TPA (29 mg, 0.1 mmol, 4 equiv.), 2,2'-bipyridine-5-carbaldehyde (56 mg, 0.3 mmol, 12 equiv.), Zn(BF₄)₂ (34 mg, 0.1 mmol, 4 equiv.), and dry acetonitrile (30 mL) were added to a Schlenk tube. The solution was degassed for three evacuation/nitrogen fill cycles. The tube was kept at 343 K for 24 h. Diethyl ether was diffused into an acetonitrile solution and the desired product **Cat-BF₄** was isolated by filtration as red square crystals in 60% yield. ¹H NMR (500 MHz, 298 K, CD₃CN): δ = 9.00(6H, d), 8.86(24H, m), 8.59–8.70(36H, m), 8.56(12H, s), 8.27–8.42(36H, m), 8.15(12H, s), 7.9–8.1(18H, d), 7.77(6H, s), 7.6–7.7(24H, m), 7.45(12H, d), 7.37(12H, d), 7.29(6H, d), 7.0(6H, d), 6.65–6.79(24H, dd), 6.66(12H, d), 6.25–6.36(24H, dd), 5.7(12H, d). ESI-MS: *m/z* [Zn₈L₈(BF₄)₇]⁹⁺ 826.8, [Zn₈L₈(BF₄)₈]¹⁸⁺ 941.0, [Zn₈L₈(BF₄)₉]²⁷⁺ 1087.9, [Zn₈L₈(BF₄)₁₀]³⁶⁺ 1283.7, [Zn₈L₈(BF₄)₁₁]⁴⁵⁺ 1557.8. Elemental analysis (Zn₈C₄₀₈H₂₈₈N₈₀B₁₆F₆₄·14H₂O) calcd (%): C 57.82, H 3.76, N 13.22; found (%): C 57.91, H 3.35, N 13.48. IR (KBr, cm⁻¹), ν 3547, 3074, 1605, 1498, 1474, 1440, 1318, 1286, 1060, 835, 796, 750, 736.

Preparation of the discrete tetrahedron Tet-1

TPA (29 mg, 0.1 mmol, 4 equiv.), 2,2'-bipyridine-5-carbaldehyde (56 mg, 0.3 mmol, 12 equiv.), Zn(CF₃SO₃)₂ (36 mg, 0.1 mmol, 4 equiv.), and dry acetonitrile (30 mL) were added to a Schlenk tube. The solution was degassed by three evacuation/nitrogen fill cycles. The tube was kept at 343 K for 24 h. Diethyl ether was diffused into the acetonitrile solution and the desired product **Tet-1** was isolated by centrifugation as a red solid. The

isolated yield was 75% (based on the product dried under vacuum). ¹H NMR (500 MHz, 298 K, CD₃CN): δ = 8.75(12H, d), 8.58–8.67(36H, m), 8.38(12H, s, *J* = 1.3 Hz, imine), 8.31(12H, m), 8.0(12H, d), 7.6(12H, m), 7.22(24H, d, *J* = 8.8 Hz), 7.01(24H, d, *J* = 8.8 Hz). ¹³C NMR (500 MHz, 298 K, CD₃CN): δ = 153.3, 149.9, 148.6, 148.5, 148.1, 146.1, 144.6, 141.7, 138.9, 134.9, 127.5, 124.2, 123.7, 123.3, 122.6. ESI-MS: *m/z*, [Zn₄L₄(CF₃SO₃)₂]⁶⁺ 619.15, [Zn₄L₄(CF₃SO₃)₃]⁹⁺ 772.77, [Zn₄L₄(CF₃SO₃)₄]¹²⁺ 1003.2 and [Zn₄L₄(CF₃SO₃)₅]¹⁵⁺ 1387.25. Elemental analysis (Zn₄C₂₁₂H₁₄₄N₄₀F₂₄S₈O₂₄·2CH₃CN·3H₂O) calcd (%): C 54.66, H 3.31, N 12.39; found (%): C 54.41, H 3.25, N 12.48. IR (KBr, cm⁻¹), ν 3378, 3063, 1605, 1499, 1474, 1440, 1317, 1280, 1164, 1030, 833, 796, 751, 638.

Preparation of the interlocked tetrahedron Cat-ClO₄ [Zn₈L₈] [ClO₄]₁₆

TPA (29 mg, 0.1 mmol, 4 equiv.), 2,2'-bipyridine-5-carbaldehyde (56 mg, 0.3 mmol, 12 equiv.), Zn(ClO₄)₂ (37 mg, 0.1 mmol, 4 equiv.), and dry acetonitrile (30 mL) were added to a Schlenk tube. The solution was degassed by three evacuation/nitrogen fill cycles. The tube was kept at 343 K for 24 h. Diethyl ether was diffused into an acetonitrile solution and the desired product **Cat-ClO₄** was isolated by filtration as red square crystals in 65% yield (based on the product dried under vacuum). ¹H NMR (500 MHz, 298 K, CD₃CN): δ = 8.88(24H, m), 8.60–8.64(64H, m), 8.52(6H, s), 8.37(6H, s), 8.28(36H, m), 8.19(12H, m), 8.06–7.97(24H, d), 7.62(36H, m), 7.47(12H, d), 7.38(12H, d), 6.8(12H, d), 6.67(12H, d), 6.6(12H, d), 6.31–6.36(24H, dd), 5.9(12H, d). ESI-MS: *m/z* [Zn₈L₈(ClO₄)₇]⁹⁺ 836.8, [Zn₈L₈(ClO₄)₈]¹⁸⁺ 953.9, [Zn₈L₈(ClO₄)₉]²⁷⁺ 1104.3, [Zn₈L₈(ClO₄)₁₀]³⁶⁺ 1304.9, [Zn₈L₈(ClO₄)₁₁]⁴⁵⁺ 1585.2. Elemental analysis: (Zn₈C₄₀₈H₂₈₈N₈₀Cl₁₆·O₆₄·2CH₃CN·7H₂O) calcd (%): C 57.3, H 3.59, N 13.3; found (%): C 56.8, H 3.65, N 13.42. IR (KBr, cm⁻¹), ν 3520, 3065, 1604, 1498, 1474, 1439, 1317, 1280, 1088, 834, 795, 750, 624.

Preparation of the interlocked tetrahedron Cat-X (X = Br or I)

TPA (29 mg, 0.1 mmol, 4 equiv.), 2,2'-bipyridine-5-carbaldehyde (56 mg, 0.3 mmol, 12 equiv.), Zn(CF₃SO₃)₂ (36 mg, 0.1 mmol, 4 equiv.), and dry acetonitrile (30 mL) were added to a Schlenk tube. The solution was degassed by three evacuation/nitrogen fill cycles. The tube was kept at 343 K for 24 h. Then Bu₄NX (X = Br or I) (7 equiv.) was added to give a solution containing the interlocked cage [Zn₈L₈] with X⁻ inclusion.

Preparation of Cat-I/BF₄ [Zn₈L₈I₉(BF₄)₇]

TPA (29 mg, 0.1 mmol, 4 equiv.), 2,2'-bipyridine-5-carbaldehyde (56 mg, 0.3 mmol, 12 equiv.), Zn(CF₃SO₃)₂ (36 mg, 0.1 mmol, 4 equiv.), and dry acetonitrile (30 mL) were added to a Schlenk tube. The solution was degassed by three evacuation/nitrogen fill cycles. The tube was kept at 343 K for 24 h. Bu₄NI (7 equiv.) was added to give a solution containing the interlocked cage, and subsequently the salt Bu₄NBF₄ (10 equiv.) was added to replace the CF₃SO₃⁻ anions. Then diethyl ether was diffused into an acetonitrile solution and the desired product **Cat-I/BF₄** was isolated by filtration as red square crystals in 55% yield. Elemental analysis (Zn₈C₄₀₈H₂₈₈N₈₀I₉B₇F₂₈·2CH₃CN·16H₂O)



calcd (%): C 54.26, H 3.67, N 12.82; found (%): C 54.31, H 3.45, N 12.98. IR (KBr, cm^{-1}), ν 3381, 3073, 1605, 1503, 1474, 1440, 1318, 1281, 1060, 835, 796, 750, 736.

X-ray crystallography

Crystals of the interlocked **Cat-BF₄** suitable for X-ray diffraction were obtained by slow diffusion of diethyl ether into an acetonitrile solution of the complex over a few days. Crystals of the interlocked **Cat-I/BF₄** suitable for X-ray diffraction were obtained by slow diffusion of diethyl ether into an acetonitrile solution of complex **Cat-I** in the presence of 10 equivalents of BF₄⁻ over a few days. The crystals were very susceptible to loss of solvent. Despite rapid handling times and a low temperature collection, the quality of data was less than ideal.

X-ray intensity data were measured on a Bruker SMART APEX CCD-based diffractometer (Mo-K α radiation, $\lambda = 0.71073 \text{ \AA}$) using the SMART and SAINT programs.^{66,67} The crystal data was solved by direct methods and further refined by full-matrix least-squares refinements on F^2 using the SHELXL-97 software.⁶⁸ Non-H atoms were refined with anisotropic displacement parameters. The hydrogen atoms within the ligand backbones and the solvent CH₃CN molecules were fixed geometrically at calculated distances and allowed to ride on the parent non-hydrogen atoms.

Crystal data for **Cat-BF₄**: Zn₄C₂₀₄H₁₅₈N₄₀O₇B₈F₃₂, $M = 4237.66$, monoclinic, space group $C2/c$, red block, $a = 47.616(2) \text{ \AA}$, $b = 31.748(1) \text{ \AA}$, $c = 56.289(2) \text{ \AA}$, $\beta = 111.304(2)^\circ$, $V = 79\,279(5) \text{ \AA}^3$, $Z = 8$, $\rho_{\text{calc}} = 0.712 \text{ g cm}^{-3}$, $\mu(\text{Mo-K}\alpha) = 0.289 \text{ mm}^{-1}$, $T = 100(2) \text{ K}$ [$R_{\text{int}} = 0.1629$]. Final R_1 [with $I > 2\sigma(I)$] = 0.0985, wR_2 (all data) = 0.2896. CCDC number 1515722.

Crystal data for **Cat-I/BF₄**: Zn₄C₂₀₆H₁₆₃N₄₁O₈B_{3.5}I_{4.5}F₁₄, $M = 4477.14$, monoclinic, space group $C2/c$, red block, $a = 47.061(2) \text{ \AA}$, $b = 32.042(1) \text{ \AA}$, $c = 55.568(3) \text{ \AA}$, $\beta = 110.4(1)^\circ$, $V = 78\,535(7) \text{ \AA}^3$, $Z = 8$, $\rho_{\text{calc}} = 0.757 \text{ g cm}^{-3}$, $\mu(\text{Mo-K}\alpha) = 0.637 \text{ mm}^{-1}$, $T = 150(2) \text{ K}$ [$R_{\text{int}} = 0.1617$]. Final R_1 [with $I > 2\sigma(I)$] = 0.0885, wR_2 (all data) = 0.2659. CCDC number 1515723.

For the refinement of **Cat-BF₄**, several atoms on some pyridine rings of the ligands were disordered into two parts, with the site occupancy factors (s.o.f.) of each part being fixed at 0.5. Several fluorine atoms in the BF₄⁻ anions were disordered into two parts with the s.o.f. of each part being refined with a fixed value. Except for the disordered parts, the partially occupied solvent molecules and the partially occupied anions, the other non-hydrogen atoms were refined anisotropically. The hydrogen atoms within the ligand backbones were fixed geometrically at calculated distances and allowed to ride on the parent non-hydrogen atoms. The bond distance in some of the pyridine and benzene rings, as well as in several BF₄⁻ anions, was confined as idealized values. The thermal parameters of adjacent atoms of the ligand backbone and the BF₄⁻ groups were confined to be similar. The SQUEEZE subroutine in PLATON was used.

For the refinement of **Cat-I/BF₄**, except for the half occupied iodide ion sited at the center of the catenane, the other iodide ions share the site with the boron atoms of the BF₄⁻ anions, with the s.o.f. of each part being fixed at a suitable value. Some

of the pyridine rings of the ligands were disordered into two parts, with the site occupancy factors (s.o.f.) of each part being fixed as a certain value. Several of the fluoride atoms in a partially occupied BF₄⁻ anion were disordered into two parts with the s.o.f. of each part being refined with a fixed value. Except for the disordered parts, the partially occupied solvent molecules and the partially occupied anions, the other non-hydrogen atoms were refined anisotropically. The hydrogen atoms within the ligand backbones were fixed geometrically at calculated distances and allowed to ride on the parent non-hydrogen atoms. The bond distance in some of the pyridine and benzene rings, as well as in several BF₄⁻ anions, was confined as idealized values. The thermal parameters of adjacent atoms of the disordered parts of the ligand backbone and the particularly occupied BF₄⁻ groups were confined to be similar. The SQUEEZE subroutine in PLATON was used.

Conflicts of interest

The authors declare no competing financial interests.

Acknowledgements

This study was supported by the National Natural Science Foundation of China (21231003, 21531001 and 21501041).

References

- 1 M. Yoshizawa, K. Kumazawa and M. Fujita, *J. Am. Chem. Soc.*, 2005, **127**, 13456.
- 2 G. H. Clever and M. Shionoya, *Chem.–Eur. J.*, 2010, **16**, 11792.
- 3 H. Dube, D. Ajami and J. Rebek Jr, *Angew. Chem.*, 2010, **122**, 3260.
- 4 G. H. Clever, S. Tashiro and M. Shionoya, *J. Am. Chem. Soc.*, 2010, **132**, 9973.
- 5 J. D. Barrio, S. T. J. Ryan, P. G. Jambrina, E. Rosta and O. A. Scherman, *J. Am. Chem. Soc.*, 2016, **138**, 5745.
- 6 R. Vilar, *Angew. Chem., Int. Ed.*, 2003, **42**, 1460.
- 7 G. T. Spence and P. D. Beer, *Acc. Chem. Res.*, 2013, **46**, 571.
- 8 T. J. Hubin and D. H. Busch, *Coord. Chem. Rev.*, 2000, **200**, 5.
- 9 G. Zhang, O. Presly, F. White, I. M. Oppel and M. Mastalerz, *Angew. Chem., Int. Ed.*, 2014, **126**, 5226.
- 10 G. Gil-Ramirez, D. A. Leigh and A. J. Stephens, *Angew. Chem., Int. Ed.*, 2015, **127**, 6110.
- 11 J. F. Ayme, J. E. Beves, C. J. Campbella and D. A. Leigh, *Chem. Soc. Rev.*, 2013, **42**, 1700.
- 12 C. J. Bruns and J. F. Stoddart, *Acc. Chem. Res.*, 2014, **47**, 2186.
- 13 M. J. Langton and P. D. Beer, *Acc. Chem. Res.*, 2014, **47**, 1935.
- 14 A. Joosten, Y. Trolez, J. P. Collin, V. Heitz and J. P. Sauvage, *J. Am. Chem. Soc.*, 2012, **134**, 1802.
- 15 K. S. Chichak, S. J. Cantrill, A. R. Pease, S.-H. Chiu, G. W. V. Cave, J. L. Atwood and J. F. Stoddart, *Science*, 2004, **304**, 1308.
- 16 M. S. Vickers and P. D. Beer, *Chem. Soc. Rev.*, 2007, **36**, 211.
- 17 F. L. Thorp-Greenwood, A. N. Kulak and M. J. Hardie, *Nat. Chem.*, 2015, **7**, 526.



- 18 S. L. Huang, T. S. A. Hor and G. X. Jin, *Coord. Chem. Rev.*, 2017, **333**, 1.
- 19 S. J. Loeb, *Chem. Soc. Rev.*, 2007, **36**, 226.
- 20 S. M. Goldup, D. A. Leigh, P. J. Lusby, R. T. McBurnery and A. M. Z. Slawin, *Angew. Chem., Int. Ed.*, 2008, **47**, 6999.
- 21 S. Yi, V. Brega, B. Captaina and A. E. Kaifer, *Chem. Commun.*, 2012, **48**, 10295.
- 22 E. M. Lewis, E. L. Gavey, S. A. Cameron and J. D. Crowley, *Chem. Sci.*, 2012, **3**, 778.
- 23 C. S. Wood, T. K. Ronson, A. M. Belenguer, J. J. Holstein and J. R. Nitschke, *Nat. Chem.*, 2015, **7**, 354.
- 24 J. E. M. Lewis, P. D. Beer, S. J. Loeb and S. M. Goldup, *Chem. Soc. Rev.*, 2017, **46**, 2577.
- 25 L. Fang, M. A. Olson, D. Benítez, E. Tkatchouk, W. A. Goddard III and J. F. Stoddart, *Chem. Soc. Rev.*, 2010, **39**, 17.
- 26 V. Balzani, A. Credi, S. Silvi and M. Venturi, *Chem. Soc. Rev.*, 2006, **35**, 1135.
- 27 A. Coskun, M. Banaszak, R. D. Astumian, J. F. Stoddart and B. A. Grzybowski, *Chem. Soc. Rev.*, 2012, **41**, 19.
- 28 M. D. Levin, D. M. Kaphan, C. M. Hong, R. G. Bergman, K. N. Raymond and F. D. Toste, *J. Am. Chem. Soc.*, 2016, **138**, 9682.
- 29 T. H. Kim, P. Mehrabi, Z. Ren, A. Sljoka, C. Ing, A. Bezginov, L. Ye, R. Pomès, R. S. Prosser and E. F. Pai, *Science*, 2017, **355**, eaag2355.
- 30 V. Marcos, A. J. Stephens, J. Jaramillo-Garcia, A. L. Nussbaumer, S. L. Woltering, A. Valero, J. F. Lemonnier, I. J. Vitorica-Yrezabal and D. A. Leigh, *Science*, 2016, **352**, 1555.
- 31 M. Frank, M. D. Johnstone and G. H. Clever, *Chem.–Eur. J.*, 2016, **22**, 14104.
- 32 S. Löffler, J. Lubben, L. Krause, D. Stalke, B. Dittrich and G. H. Clever, *J. Am. Chem. Soc.*, 2015, **137**, 1060.
- 33 J. F. Ayme, J. E. Beves, C. J. Campbell, G. Gil-Ramírez, D. A. Leigh and A. J. Stephens, *J. Am. Chem. Soc.*, 2015, **137**, 9812.
- 34 D. Samanta and P. S. Mukherjee, *J. Am. Chem. Soc.*, 2014, **136**, 17006.
- 35 A. K. Bar, S. Raghothama, D. Moon and P. S. Mukherjee, *Chem.–Eur. J.*, 2012, **18**, 3199.
- 36 S. Löffler, J. Lubben, A. Wuttke, R. A. Mata, M. John, B. Dittrich and G. H. Clever, *Chem. Sci.*, 2016, **7**, 4676.
- 37 S. Freye, R. Michel, D. Stalke, M. Pawliczek, H. Frauendorf and G. H. Clever, *J. Am. Chem. Soc.*, 2013, **135**, 8476.
- 38 D. Preston, J. E. M. Lewis and J. D. Crowley, *J. Am. Chem. Soc.*, 2017, **139**, 2379.
- 39 D. Preston, A. Fox-Charles, W. K. C. Lo and J. D. Crowley, *Chem. Commun.*, 2015, **51**, 9042.
- 40 T. K. Ronson, S. Zarra, S. P. Black and J. R. Nitschke, *Chem. Commun.*, 2013, **49**, 2476.
- 41 M. M. J. Smulders, I. A. Riddell, C. Browne and J. R. Nitschke, *Chem. Soc. Rev.*, 2013, **42**, 1728.
- 42 D. A. Leigh, R. G. Pritchard and A. J. Stephens, *Nat. Chem.*, 2014, **6**, 978.
- 43 L. Yang, X. Jing, C. He, Z. Chang and C. Y. Duan, *Chem.–Eur. J.*, 2016, **22**, 18107.
- 44 S. Freye, D. M. Engelhard, M. John and G. H. Clever, *Chem.–Eur. J.*, 2013, **19**, 2114.
- 45 M. Fujita, N. Fujita, K. Ogura and K. Yamaguchi, *Nature*, 1999, **400**, 52.
- 46 A. Westcott, J. Fisher, L. P. Harding, P. Rizkallah and M. J. Hardie, *J. Am. Chem. Soc.*, 2008, **130**, 2950.
- 47 T. Hasell, X. Wu, T. A. Jones, J. Basca, A. Steiner, T. Mitra, A. Trewin, D. J. Adams and A. I. Cooper, *Nat. Chem.*, 2010, **2**, 750.
- 48 J. Mosquera, T. K. Ronson and J. R. Nitschke, *J. Am. Chem. Soc.*, 2016, **138**, 1812.
- 49 R. Sekiya, M. Fukuda and R. Kuroda, *J. Am. Chem. Soc.*, 2012, **134**, 10987.
- 50 H. Lee, P. Elumalai, N. Singh, H. Kim, S. U. Lee and K.-W. Chi, *J. Am. Chem. Soc.*, 2015, **137**, 4674.
- 51 Y. H. Li, J. J. Jiang, Y. Z. Fan, Z. W. Wei, C. X. Chen, H. J. Yu, S. P. Zheng, D. Fenske, C. Y. Su and M. Barboiu, *Chem. Commun.*, 2016, **52**, 8745.
- 52 Y. Luo, *Comprehensive Handbook of Chemical Bond Energies*, CRC Press, 2007.
- 53 M. D. Pluth and K. N. Raymond, *Chem. Soc. Rev.*, 2007, **36**, 161.
- 54 W. Wang, Y. X. Wang and H. B. Yang, *Chem. Soc. Rev.*, 2016, **45**, 2656.
- 55 A. V. Davis, D. Fiedler, G. Seeber, A. Zahl, R. van Eldik and K. N. Raymond, *J. Am. Chem. Soc.*, 2006, **128**, 1324.
- 56 A. V. Davis and K. N. Raymond, *J. Am. Chem. Soc.*, 2005, **127**, 7912.
- 57 S. Freye, J. Hey, A. Torras-Galan, D. Stalke, R. Herbsy-Irmer, M. John and G. H. Clever, *Angew. Chem., Int. Ed.*, 2012, **51**, 2191.
- 58 R. Zhu, J. Lubben, B. Dittrich and G. H. Clever, *Angew. Chem., Int. Ed.*, 2015, **54**, 2796.
- 59 G. B. Huang, S. H. Wang, H. Ke, L. P. Yang and W. Jiang, *J. Am. Chem. Soc.*, 2016, **138**, 14550.
- 60 J.-P. Demers and A. Mittermaier, *J. Am. Chem. Soc.*, 2009, **131**, 4355.
- 61 C.-F. Zhang, S.-H. Li, C.-C. Zhang and Y. Liu, *Org. Biomol. Chem.*, 2015, **13**, 10808.
- 62 Y. M. Zhang, X. J. Zhang, X. Xu, X. N. Fu, H. B. Hou and Y. Liu, *J. Phys. Chem. B*, 2016, **120**, 3932.
- 63 H. Dube, M. R. Ams Jr and J. Rebek, *J. Am. Chem. Soc.*, 2010, **132**, 9984.
- 64 D. H. Qu, Q. C. Wang, Q. W. Zhang, X. Ma and H. Tian, *Chem. Rev.*, 2015, **115**, 7543.
- 65 X. Jing, C. He, Y. Yang and C. Y. Duan, *J. Am. Chem. Soc.*, 2015, **137**, 3967.
- 66 *SMART Data collection software, version 5.629*, Bruker AXS Inc., Madison, WI, 2003.
- 67 *SAINTE Data reduction software, version 6.45*, Bruker AXS Inc., Madison, WI, 2003.
- 68 G. M. Sheldrick, *SHELX-97: Program for crystal structure analysis*, University of Göttingen, Göttingen, Germany, 1997.

