

ChemComm

Chemical Communications

Accepted Manuscript

This article can be cited before page numbers have been issued, to do this please use: A. Arun, H. M. Tay and P. D. Beer, *Chem. Commun.*, 2024, DOI: 10.1039/D4CC03916E.



This is an Accepted Manuscript, which has been through the Royal Society of Chemistry peer review process and has been accepted for publication.

Accepted Manuscripts are published online shortly after acceptance, before technical editing, formatting and proof reading. Using this free service, authors can make their results available to the community, in citable form, before we publish the edited article. We will replace this Accepted Manuscript with the edited and formatted Advance Article as soon as it is available.

You can find more information about Accepted Manuscripts in the [Information for Authors](#).

Please note that technical editing may introduce minor changes to the text and/or graphics, which may alter content. The journal's standard [Terms & Conditions](#) and the [Ethical guidelines](#) still apply. In no event shall the Royal Society of Chemistry be held responsible for any errors or omissions in this Accepted Manuscript or any consequences arising from the use of any information it contains.

ARTICLE

Mechanically Interlocked Host Systems for Ion-Pair Recognition**

Arya Arun,^{a,b} Hui Min Tay^a and Paul D. Beer^{*a}Received 00th January 20xx,
Accepted 00th January 20xx

DOI: 10.1039/x0xx00000x

The ever-increasing interest directed towards the construction of host architectures capable of the strong and selective recognition of various ionic species of biological, medical and environmental importance has identified mechanically interlocked molecules (MIMs), such as rotaxanes and catenanes, as potent host systems, owing to their unique three-dimensional topologically preorganised cavity recognition environments. Ion-pair receptors are steadily gaining prominence over their monotopic receptor analogues due to their enhanced binding strength and selectivity demonstrated primarily through acyclic and macrocyclic heteroditopic host systems. Exploiting the mechanical bond for ion-pair recognition through the strategic design of neutral heteroditopic MIMs offer exciting opportunities to accomplish potent and effective binding while mitigating competing interactions from the bulk solvent and counter-ions. This review details the design and ion-pair recognition capabilities of rotaxanes and catenanes employing hydrogen bonding (HB) and halogen bonding (XB) motifs, providing valuable insight into the burgeoning field and inspiration for future research.

Introduction

Driven by the prevalence and importance of charged species in chemistry,^{1, 2} biology³⁻⁵ and the environment,⁶⁻⁸ the design of molecular host systems for ion recognition remains a crucial research focus within supramolecular chemistry, with ongoing efforts aimed at enhancing binding affinities and selectivity profiles through strategic structural host design.

The development of supramolecular anion receptors has lagged behind its sister field of cation coordination, reflecting the greater challenges associated with anion binding, including relatively low charge densities, pH sensitivity, variable geometries and high hydration enthalpies of anions.⁹ Indeed, anion receptors capable of functioning in competitive aqueous-containing media have traditionally utilised multiple positively charged scaffolds.¹⁰⁻¹⁵ Despite the gains in anion affinity elicited by employing favourable Coulombic interactions, their non-directional nature can lead to compromised binding selectivity. Additionally, the nominally 'non-coordinating' counter-anions of the positively charged receptor may compete with the target anion guest. From a functional perspective, the reliance on positively charged binding motifs limits the potential applications of anion receptors; for instance, anion transport across lipid bilayers typically requires lipophilic carrier molecules.^{16, 17} To address these limitations, interest in the design and synthesis of *neutral* anion receptors has been steadily growing in recent years.

In this context, mechanically interlocked molecules (MIMs) offer distinct advantages over traditional acyclic or macrocyclic hosts due to their unique structural and topological features (Figure 1). The pre-organised and solvent-shielded cavities of

MIMs present unique opportunities to design three dimensional binding sites for guest encapsulation, which through non-covalent donor/acceptor group functionalisation can be tuned to achieve size and shape complementarity towards a target guest.¹⁸⁻²⁵ The resulting improvements in binding affinity and selectivity over non-interlocked receptor analogues, termed a 'mechanical bond effect' (MBE),²⁶ have motivated the construction of numerous MIM-based cation²⁷⁻³¹ and anion³²⁻³⁷ receptors over the years. In particular, the development of highly potent charge-neutral binding motifs has enabled the design of neutral MIM-based receptors for various novel applications, broadening the potential functionality of this class of receptors. Pioneering studies on transition metal cation template-directed synthesis of MIMs by Sauvage and co-workers demonstrated the potential of neutral catenanes to bind and stabilise low oxidation states of transition metal cations such as copper(I) *via* encapsulation of the guest within a bis-phenanthroline-functionalised interlocked cavity.³⁸⁻⁴⁴ In more recent years, Goldup and co-workers designed a series of neutral [2]rotaxanes capable of binding transition metal cations in CH₃CN, wherein the confined environment of the interlocked cavity enforces unusual coordination geometries and redox state stabilisation.⁴⁵

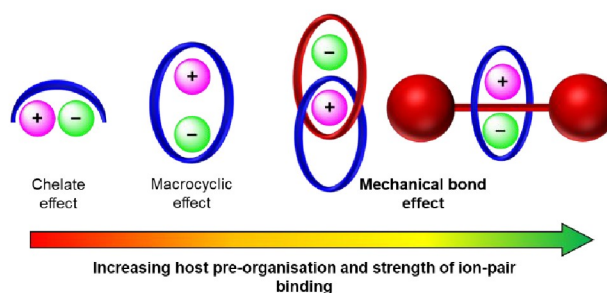


Figure 1. Mechanical bond effect enhanced ion-pair recognition.

^a Department of Chemistry, University of Oxford, Chemistry Research Laboratory Mansfield Road, Oxford OX1 3TA, UK. E-mail: paul.beer@chem.ox.ac.uk

^b Department of Chemistry, University of Oxford, Rodney Porter Building, Sibthorp Road, Oxford OX1 3QU, UK.



Notable advancements have also been made in the area of neutral anion binding MIMs, with Beer and co-workers reporting a template-directed synthetic strategy driven by potent halogen bonding (XB)···Cl⁻ interactions to prepare neutral interlocked hosts capable of halide anion recognition in aqueous-acetone solvent mixtures.⁴⁶ This strategy was recently employed in the preparation of the first hetero[2]catenanes capable of binding and selectively transporting halide anions across a phospholipid bilayer.⁴⁷ The compact geometry and lipophilic nature of the catenane hosts enabled facile membrane transversal while their strong binding affinity and preference for halides translated to an impressive transport selectivity for chloride over nitrate and hydroxide anions. Rotaxane hosts prepared using a similar template-directed approach have also shown promise as selective colorimetric sensors for chloride anions.⁴⁸ Recently, Chmielewski and co-workers reported a charge-neutral homo[2]catenane synthesised *via* a novel sulfate anion template-directed strategy, which showed impressive binding affinity and selectivity for sulfate over other oxoanions and halides, and was capable of functioning as a sensitive fluorescence sensor for sulfate anions in competitive 9:1 DMSO/H₂O solvent mixtures.⁴⁹

While charge-neutral MIM-based receptors are increasingly finding applications in cation and anion recognition, their greatest advantage over charged receptors lies in their propensity to be utilised as heteroditopic receptors capable of simultaneously binding a cation and anion ion-pair. Ion-pair recognition has emerged as a powerful strategy to augment the binding properties of a receptor by exploiting favourable proximal cation-anion electrostatic interactions and, in some cases, conformational allosteric cooperativity associated with binding oppositely charged guests.⁵⁰ Indeed, studies conducted on acyclic/macrocyclic heteroditopic ion-pair receptors generally reveal improved binding affinities for ion-pairs or constituent ions relative to monotopic receptor analogues.⁵¹ This has encouraged their use in a myriad of applications, including salt extraction/solubilisation,⁵²⁻⁶² membrane transport^{63, 64} and the recognition of biologically-relevant zwitterions.^{65, 66}

Mechanically interlocked molecules are promising scaffolds upon which cation *and* anion binding sites can be incorporated in a modular fashion to construct complex heteroditopic receptors. While still in its infancy, the field has gained significant traction in recent years, with a growing diversity of catenane- and rotaxane-based heteroditopic hosts that show promising ion-pair binding properties and potential applications in ion-pair sensing and extraction.⁶⁷⁻⁷²

In this article, we present prominent developments in the field of mechanically interlocked heteroditopic host molecules for ion-pair recognition, ranging from early receptors that utilise traditional hydrogen bonding (HB) interactions for anion binding to recent examples featuring halogen bonding (XB) donor groups. In particular, we highlight the importance of the interlocked receptor topology in creating solvent-shielded binding cavities and its role in dictating MBE enhanced ion-pair affinity and selectivity profiles of the receptors.

Heteroditopic MIM-based Receptors

Heteroditopic receptors typically consist of one or more cation and anion binding sites, which can be located in close proximity to each other or spatially separated. This forms the basis for a broad general classification of ion-pair receptors into contact ion-pair or separated ion-pair receptors (**Figure 2**). Contact ion-pairs are formed when the two bound ions are held proximal to each other within the receptor, as determined by X-ray crystallographic studies. In contrast, when the cation and anion binding sites are distal to each other, the co-bound ion-pair may be separated by solvent molecules or the receptor framework.⁷³

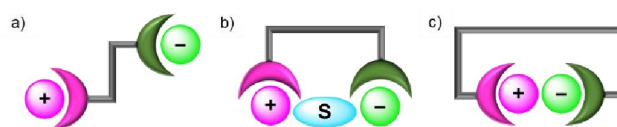


Figure 2. Ion-pair binding modes in heteroditopic receptors: (a) Host-separated; (b) Solvent-separated; (c) Direct contact.

The literature surrounding supramolecular ion-pair recognition has been largely dominated by reports of heteroditopic receptors for alkali metal halide binding,^{52, 61, 63, 64, 74-86} owing to the ubiquity of these salts across a range of biological and environmental systems. For instance, sodium and potassium are crucial in maintaining cellular electrolyte concentrations, while the mis-regulation of chloride and iodide in the body has been associated with various diseases.⁸⁷⁻⁸⁹ Recently, methods for detection and recovery of lithium salts have grown in prominence with the large-scale development of lithium-ion battery technologies.^{90, 91} Heteroditopic receptors capable of binding alkali metal halide ion-pair binding typically utilise hard oxygen donors for metal cation coordination, often introduced as crown ether or poly(ethylene glycol) moieties, alongside a range of neutral hydrogen bonding, and more recently halogen bonding, donor groups for halide binding. The synthetic accessibility of these binding motifs, as well as their proven ability for strong and selective alkali metal halide binding, have encouraged their integration into a range of MIM-based heteroditopic receptors.

MIM Heteroditopic Receptors Utilising Hydrogen Bonding (HB) Interactions

Despite the advantages of their unique topologies and preorganised interlocked cavities that have in particular been shown to enhance anion guest recognition capabilities, examples of neutral heteroditopic MIMs remain relatively scarce. Early examples of neutral hydrogen bonding interlocked frameworks capable of ion-pair recognition include Smith's [2]rotaxane **1a** comprising a macrobicyclic component functionalised with a crown ether cation and isophthalamide



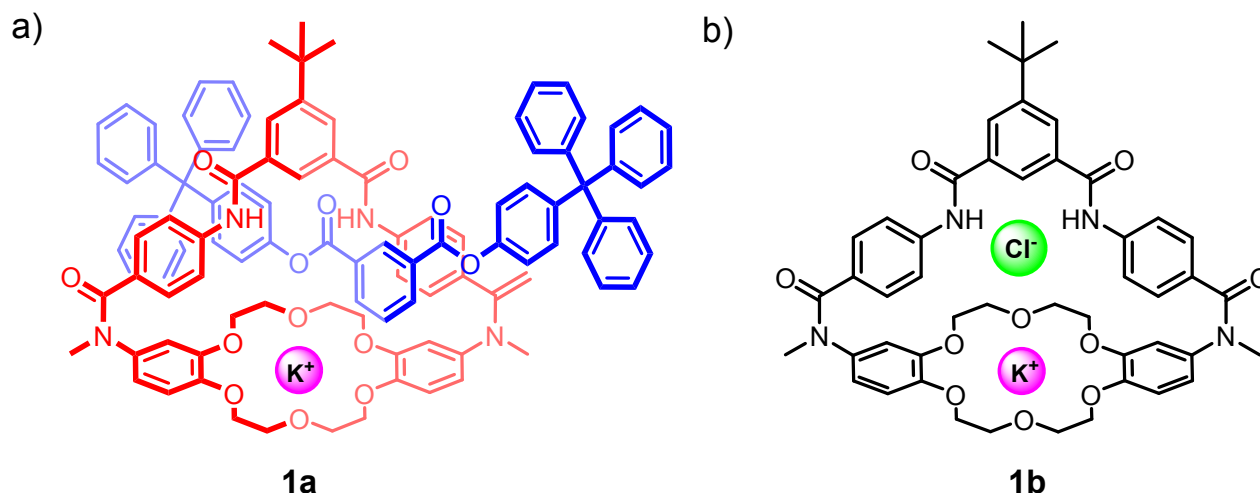


Figure 3. a) Smith's heteroditopic [2]rotaxane **1a** capable of KCl ion-pair binding; b) Macrobicyclic precursor **1b**, which previously also demonstrated cooperative binding of KCl ion-pairs.

anion binding units (Figure 3a).⁷⁸ The synthesis of the rotaxane **1a** utilises a discrete potassium cation template bound to the dibenzo-18-crown-6 ether group to direct the association of an anionic phenolate half-axle through the isophthalamide cavity. Subsequent stoppering esterification reactions afforded the [2]rotaxane in 20% yield. The crucial role played by the potassium cation template was highlighted by the failure to form the desired rotaxane product in the absence of the potassium cation as well as in the presence of a larger cesium cation. ¹H-NMR anion binding studies conducted with the neutral [2]rotaxane **1a** in DMSO-*d*₆/CD₃CN solvent mixtures reveal enhanced chloride anion affinities relative to the free heteroditopic macrobicyclic receptor **1b** (Figure 3b) ($K_a = 300 \text{ M}^{-1}$ and $K_a = 50 \text{ M}^{-1}$ respectively), attributable to the steric shielding effect of the interlocked binding cavity inhibiting the competing DMSO complexation previously reported to occur within the free macrocycle **1b**. Interestingly, whilst macrobicyclic **1b** exhibits a 7-fold enhancement in chloride binding affinity in the presence of potassium, no enhancement in chloride binding is observed in the case of the [2]rotaxane **1a**. This lack of cooperativity is ascribed to the aforementioned steric effects engendered by the interpenetrating axle which disrupt the electrostatic interactions between the co-bound ions. Furthermore, ¹H-NMR studies in polar organic solvents suggest that the presence of a K⁺ cation serves to 'lock' the dynamic co-conformations assumed by the components in the free state, as evidenced by the sharpening of the proton signals in the ¹H-NMR spectrum. By contrast, chloride complexation fails to disrupt the dynamic inter-component motions likely due to the lack of any axle-anion interactions.

Since then only a handful of neutral hydrogen bond donor containing heteroditopic MIMs have been reported, including Beer's neutral [2]rotaxane **2** consisting of a pyridine *N*-oxide axle and an isophthalamide bridged calix[4]diquinone macrocycle operating as an axle-separated ion-pair receptor (Figure 4a).⁶⁷ Synthesised using a sodium cation-template driven threading methodology followed by copper(I) catalysed-mediated CuAAC stoppering reactions, the resulting [2]rotaxane demonstrates remarkable alkali metal halide ion-pair recognition in 4:1 CDCl₃/CD₃OD solvent media. ¹H-NMR based characterisation studies suggest that the free rotaxane **2** adopts a co-conformation where the axle pyridine *N*-oxide forms intercomponent hydrogen bonds with the macrocycle bis-amide functionalised pyridine unit. Sodium cation complexation thereafter pre-organises the *N*-oxide motif to pirouette within the macrocycle towards the calixdiquinone unit. Quantitative ¹H-NMR studies conducted on **2** reveal significant amplification in the halide binding constants in the presence of co-bound sodium and potassium cations, displaying a striking 15-fold cooperativity factor in the case of the NaCl ion-pair. This enhancement is accredited to favourable axle-separated electrostatic interactions between the co-bound ions and receptor preorganisation in the presence of the alkali metal cation. More recently, the same authors prepared an analogous [2]catenane **3** via a sodium cation-template activated ester amide condensation clipping methodology (Figure 4b), wherein the [2]catenane was serendipitously isolated with a NaCl ion-pair endotopically bound within the binding cavities separated by the macrocycles, characterised by single crystal X-ray structural analysis (Figure 4c).⁹²



ARTICLE

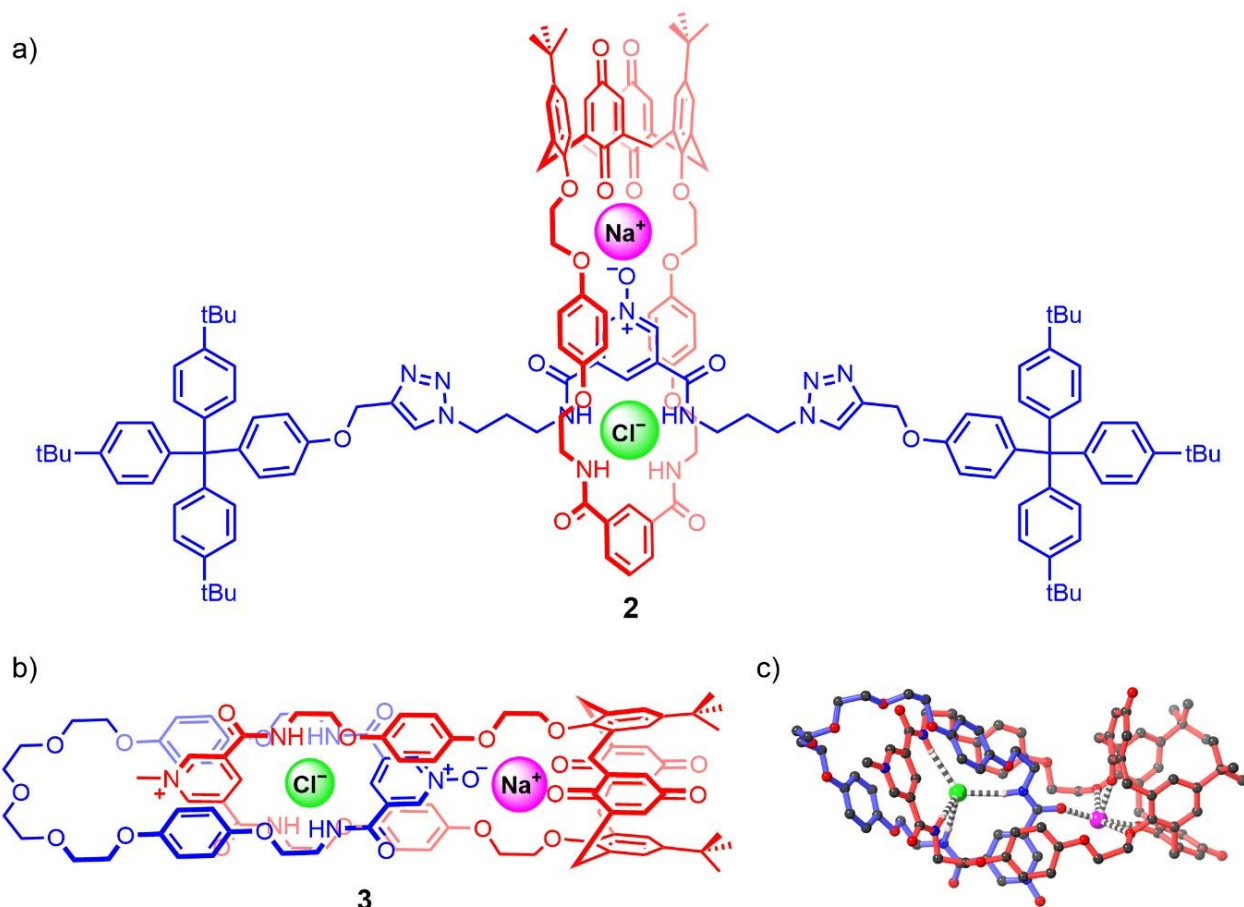


Figure 4. Knighton and Beer's heteroditopic a) [2]rotaxane **2** and b) [2]catenane **3**, based on interlocked calix[4]diquinone and pyridine-N-oxide components. c) Single crystal XRD structure of [2]catenane bound to an NaCl ion-pair.

Another recent example includes Goldup's fluorescent [2]rotaxane **4** comprising a urea-based naphthalimide axle component and bipyridine functionalised macrocycle, synthesised using active metal template methodology in 92% yield (Figure 5).⁷¹ Analysis of ¹H-NMR, UV-vis and fluorescence titration data of **4** reveals that in the free state, the bipyridine group in the macrocycle interacts with the anion binding urea motif in the axle *via* strong HB interactions, precluding it from participating in anion binding even in the presence of highly basic anions such as AcO⁻ and F⁻. By contrast, the free axle exhibits similar binding trends albeit with higher binding affinities. Notably, protonating the bipyridine motif enables macrocyclic translocation away from the urea unit in the axle, consequently enabling exogenous anion recognition at the urea station in 1:1 CDCl₃/CD₃CN solvent mixtures ($K_a(\text{Cl}^-) > 10^4$, $K_a(\text{Br}^-) = 4.7 \times 10^3$, $K_a(\text{HSO}_4^-) = 2.3 \times 10^3$). Interestingly, while the protonated rotaxane **4-H⁺** shows a greater affinity towards less

basic anions in comparison to the free axle, presumably due to favourable electrostatic interactions between the co-bound ions, addition of basic anions like AcO⁻ and F⁻ predictably results in deprotonation of the interlocked system.

Calix[4]pyrroles and their derivatives have been formerly known to function as heteroditopic receptors, binding anions through the cone-shaped pyrrole NH groups and cations *via* the electron rich bowl-shaped cavity formed by the calix[4]pyrrole anion complex.⁹³ Looking to exploit the ambidentate nature of calix[4]pyrroles, Ballester and co-workers sought to design a rigid three-dimensional macrocyclic scaffold containing two calix[4]pyrrole units, envisioned to bind ion-pairs by forming cascade complexes.⁹⁴ Encouraged by the binding behaviour exhibited by the macrocycle, the authors synthesised a bis-calix[4]pyrrole functionalized [2]rotaxane **5** prepared using CuAAC 'click' reactions to stopper a pseudo-[2]rotaxane assembly involving a N-oxide axle thread and the bis-



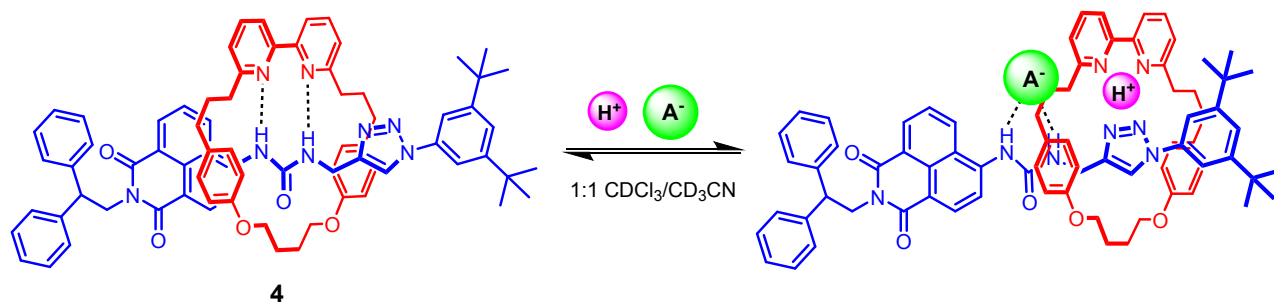


Figure 5. Goldup's dynamic [2]rotaxane shuttle **4** capable for cooperative HCl ion-pair binding.

calix[4]pyrrole containing cylindrical macrocycle component (**Figure 6**).⁷⁰ ¹H-NMR based binding studies indicated that **5** strongly bound tetraalkylammonium salts of Cl⁻, OCN⁻ and NO₃⁻ in a 1:1 binding stoichiometry in CDCl₃. Quantitative ITC titration experiments further revealed that the interlocked system demonstrated a preference for TBANCO ($K_a = 7.9 \times 10^5 \text{ M}^{-1}$) over TBANO₃ ($K_a = 4 \times 10^4 \text{ M}^{-1}$) and TBACl ($K_a = 5 \times 10^4 \text{ M}^{-1}$) in chloroform, likely due to host-guest size complementarity between the cylindrical cavity and linear NCO⁻ anion. Notably, replacing the organic cation to methyltriethylammonium (MTEO⁺) resulted in a 20-fold increase in the binding of MTEOCl relative to TBANCO, presumably due to better encapsulation of the quaternary ammonium cation within the calix[4]pyrrole unit.

Squaramides are another well-known class of neutral, potent hydrogen bonding anion receptors. However, the unique ambidentate properties of the squaramide motif allows for the potential simultaneous binding of anions through the Lewis acidic NH donors and cations *via* bidentate chelation of the Lewis basic carbonyl groups. Beer and co-workers recently demonstrated for the first time ambidentate squaramide ion-pair binding behaviour through a series of HB functionalized heteroditopic [2]rotaxanes **6a-c** wherein the squaramide-based axle component played a pivotal role in achieving axle-separated sodium halide ion-pair recognition (**Figure 7**).⁹⁵

Taking inspiration from Chiu's alkali metal template directed MIM synthesis,⁹⁶ an unprecedented sodium cation-template directed strategy driven by orthogonal coordination of the sodium cation by the Lewis basic carbonyls of the axle squaramide unit and the polyether/pyridyl region of the macrocycle was employed to facilitate threading and subsequent mechanical bond formation.

In order to investigate the factors governing the ion-pair association capabilities of the interlocked squaramide axle containing [2]rotaxane hosts, macrocycles with varying cation and anion binding units were chosen, functionalized with either crown-ether like arrangements of oxygen donors or a combination of polyether and pyridyl isophthalamide linkages. ¹H-¹H ROESY NMR characterisation indicated that the free squaramide rotaxanes adopted co-conformations where the axle squaramide carbonyls were engaged in hydrogen bonding interactions with the bis-amide isophthalamide unit of the macrocycle in concert with analogous interactions between the macrocycle polyether/pyridyl units and axle squaramide NH groups. Notably, sodium cation binding co-conformationally pre-organises the macrocycle such that amide donors from both the macrocycle and axle form an intercomponent convergent hydrogen bond donor site for cooperative anion recognition. Extensive cation, anion and ion-pair ¹H-NMR titration studies in 3:7 CD₃CN/CDCl₃ revealed the [2]rotaxane hosts to

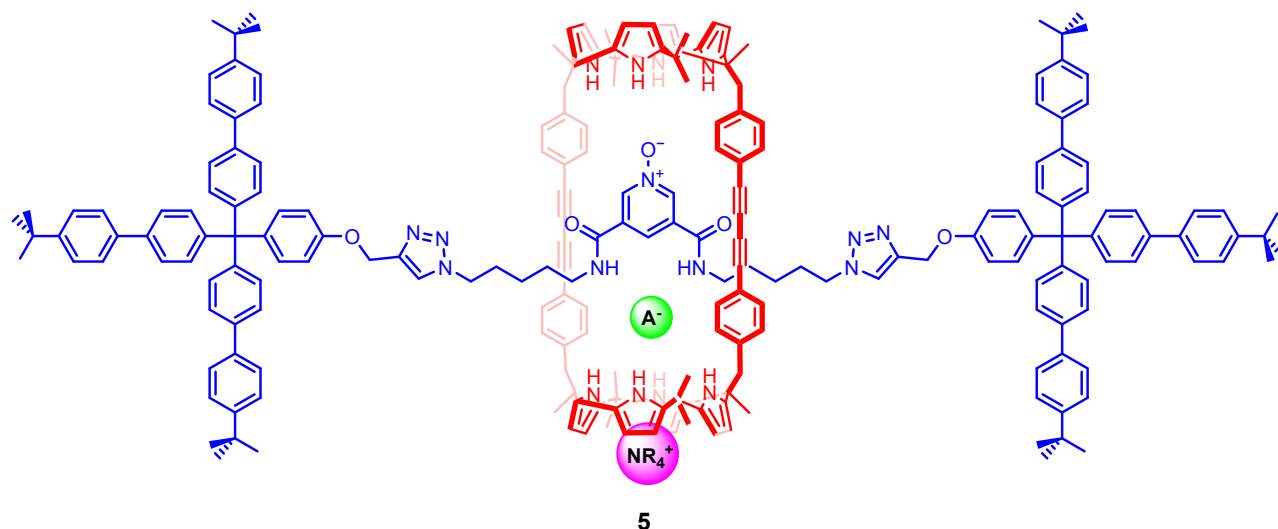


Figure 6. Ballester's [2]rotaxane **5** for the recognition of quaternary ammonium salts.



cooperatively bind and extract sodium halide salts, engendering up to a *ca.* 20-fold enhancement in bromide and iodide binding strength in the presence of an equivalent of sodium cation ($K_a = 494$ to 10^4 M^{-1} and $K_a = 67$ to 1298 M^{-1} respectively for the pyridyl functionalized [2]rotaxane **6c**). This remarkable amplification in association constant magnitudes was attributed to favourable coulombic electrostatic interactions between the co-bound ions and co-conformational allosteric

effects instigated by the ion-binding event. In the case of the NaCl ion-pair, exogenous ion-pairing salt recombination equilibrium necessitated the calculation of % bound sodium to the [2]rotaxane systems in order to rationalise the differing chloride binding properties of the sodium complexed receptors. While systems with macrocycles containing the crown-ether arrangement (**6a**) and polyether linkages (**6b**) failed to bind the NaCl ion-pair, the pyridyl-functionalised rotaxane **6c**

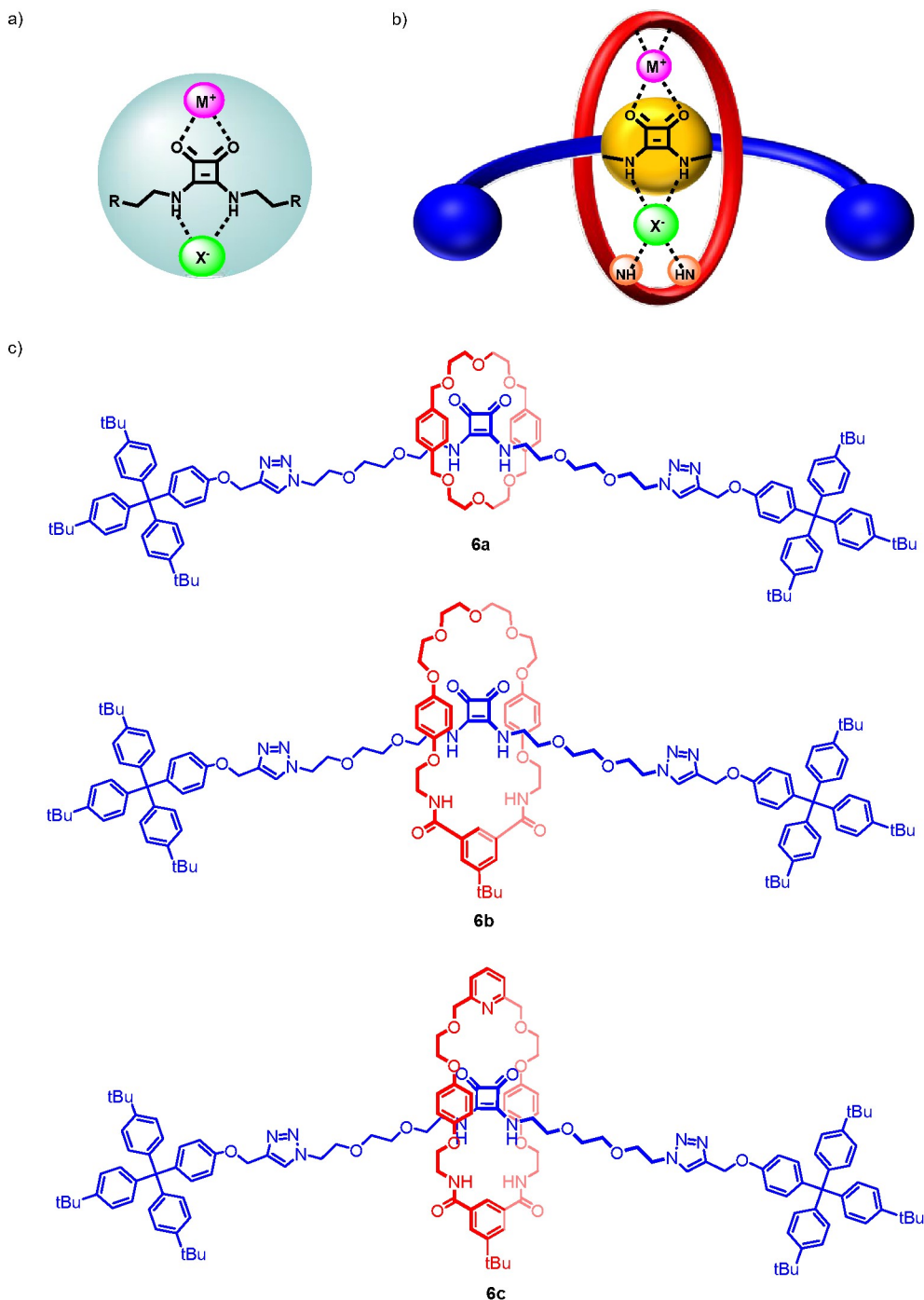


Figure 7. a) Propensity of the squaramide motif for metal cation and ion-pair recognition, highlighting its ambidentate coordination mode, b) Cartoon showing ion-pair recognition in target squaramide axle containing [2]rotaxane host systems, c) Beer's squaramide-based [2]rotaxanes **6a-c** for the recognition of sodium halide ion-pairs.



successfully overcame the NaCl contact ion-pair lattice enthalpy, underpinning the crucial interplay of strong concomitant cation and anion binding affinities of the receptors in enabling ion-pair recognition.

Heteroditopic Receptors Utilising σ -Hole Interactions

Although hydrogen bonding motifs have traditionally dominated the field of anion recognition, recently the integration of σ -hole interactions into acyclic, macrocyclic and mechanically interlocked anion receptor systems has emerged as a promising alternative strategy. In particular, halogen bonding (XB)⁹⁷ interactions have grown in prominence by virtue of their often superior anion binding strength, more stringent linear interaction geometries and contrasting selectivity profiles relative to analogous HB receptors.^{50, 98–102} Encouraged by these studies, we have begun to explore the potential of

incorporating XB and related σ -hole binding motifs into heteroditopic ion-pair receptors as a means to further tune ion-pair binding affinity and selectivity.

Acyclic and Macrocyclic Heteroditopic Receptors Utilising σ -Hole Interactions

The potential efficacy of using halogen bonding and related σ -hole interactions to modulate the affinity and selectivity profile of ion-pair receptors was initially demonstrated in non-interlocked heteroditopic host molecules. The first example of using halogen bonding interactions in ion-pair binding was reported in 2005 by Resnati and co-workers, who synthesised a tripodal heteroditopic host **7** comprising a tris(olyoxyethylene)amine cation recognition site and 4-iodotetrafluorophenyl anion recognition sites (**Figure 8a**). ¹H NMR competition studies conducted in 60:35:5 CDCl₃/CD₃OD/D₂O (v/v/v) showed **7** was able to bind an NaI ion-pair with significantly higher affinity than the non-XB pentafluorophenyl analogue ($K_{\sigma}^{\text{XB}} = 2.6 \times 10^5 \text{ M}^{-1}$; $K_{\text{a}}^{\text{HB}} = 1.3 \times$

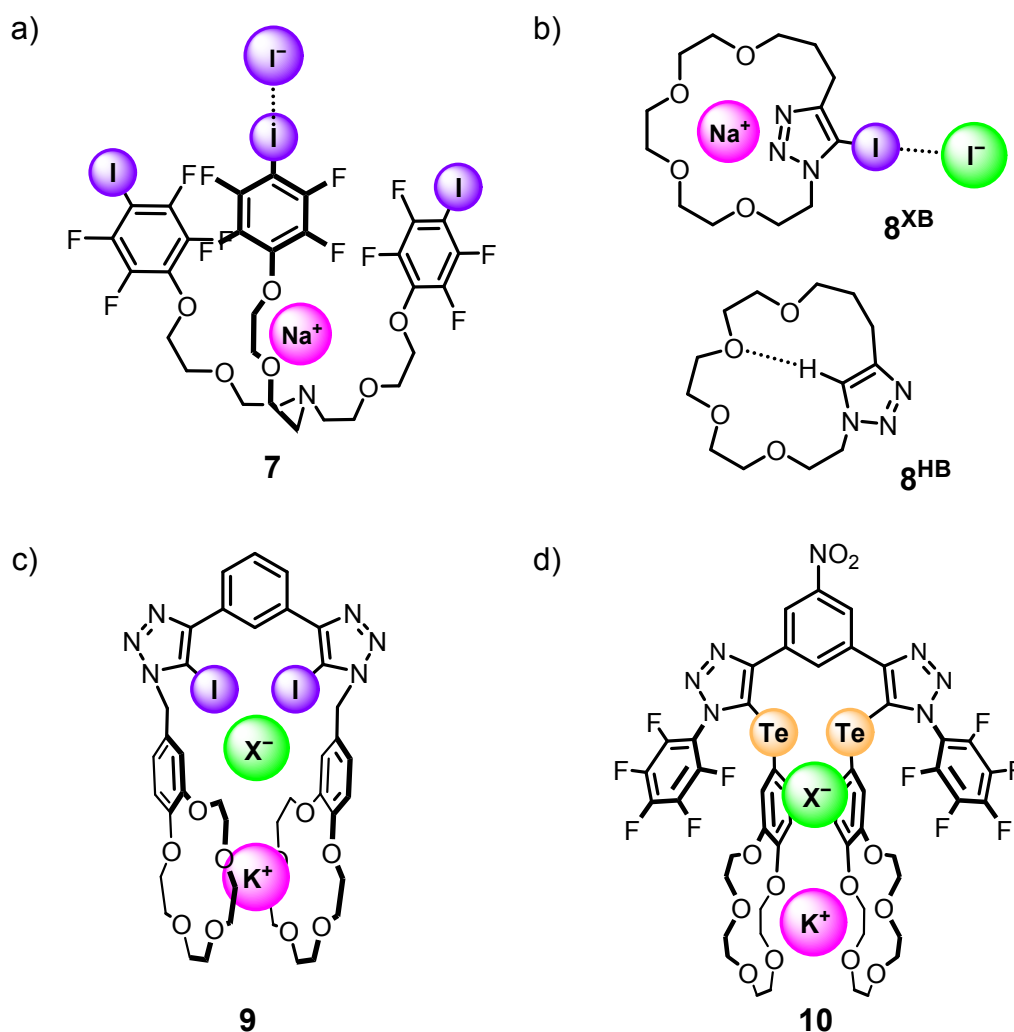


Figure 8. Heteroditopic ion-pair receptors utilising XB and ChB interactions: a) Resnati's tripodal receptor **7**; b) Schubert's triazole-incorporated cyclic ethylene glycol receptors **8**; c) Taylor's B15C5-functionalised XB receptors **9**; d) Docker's B15C15-functionalised ChB receptors **10**.



10^4 M^{-1}). However, single crystal X-ray analysis suggested that only one iodine donor in **7** participates in XB interactions with the iodide guest due to the divergent arrangement of the donor motifs.¹⁰³ Subsequently, Schubert and co-workers integrated a iodotriazole anion binding group into a tri(ethylene glycol)-containing macrocycle to generate a heteroditopic receptor **8^{XB}** capable of binding sodium iodide ion-pairs in 3:1 $\text{CD}_2\text{Cl}_2/\text{CD}_3\text{CN}$ solvent mixtures with significant positive cooperativity between the cation and anion binding events (**Figure 8b**). In contrast, the hydrogen bonding analogue **8^{HB}** showed no measurable ion-pair binding in the same solvent system, which was attributed to intramolecular hydrogen bond formation between the prototriazole and ether oxygen atoms outcompeting guest binding.¹⁰⁴

In an elegant demonstration of utilising allosteric cooperativity in receptor design, a series of potassium halide-selective acyclic ion-pair receptors were designed to exploit the tendency of benzo-15-crown-5 ether (B15C5) to form 2:1 sandwich complexes with potassium ions.⁷⁶ One example reported by Taylor and co-workers features a 1,3-bis(iodotriazole)benzene-based XB ion-pair receptor **9** functionalised with B15C5 groups (**Figure 8c**). Pre-complexation of **9** to K^+ or Rb^+ facilitated the formation of an intramolecular sandwich complex which led to significant enhancements of up

to 700-fold in the apparent halide anion association constants of the receptor, whereas Na^+ -complexation resulted in only a 15-fold enhancement arising from electrostatic effects.⁷⁷ Recently, Docker and co-workers reported a chalcogen bonding (ChB)⁹⁸ heteroditopic receptor **10** based on a modified receptor design, which consists of an acyclic 3,5-bis-tellurotriazole nitro-benzene scaffold with B15C5 units directly appended to the tellurium-incorporated triazoles (**Figure 8d**). **10** showed marked selectivity for potassium chloride over other alkali metal chloride salts, with crystallographic studies confirming the formation of an intramolecular co-facial bis-B15C5 K^+ sandwich complex. In addition **10** was shown to solubilise KCl in CDCl_3 through a series of solid-liquid and liquid-liquid extraction studies, and preliminary U-tube membrane transport experiments further indicated its capability to selectively transport KCl across a membrane.⁶⁰

Tse and co-workers prepared a series of heteroditopic macrocycles comprising a phenanthroline-based cation binding site opposite a bidentate anion binding motif based on either halogen bond (**11^{XB}**), chalcogen bond (**11^{ChB}**) or hydrogen bond (**11^{HB}**) donor motifs (**Figure 9a**).⁵⁷ ^1H NMR titration studies in $\text{CDCl}_3/\text{CD}_3\text{CN}$ solvent mixtures indicated strong cooperative binding of lithium halide ion-pairs by **11^{XB}** and **11^{ChB}**, which exhibited enhanced binding affinities relative to **11^{HB}**.

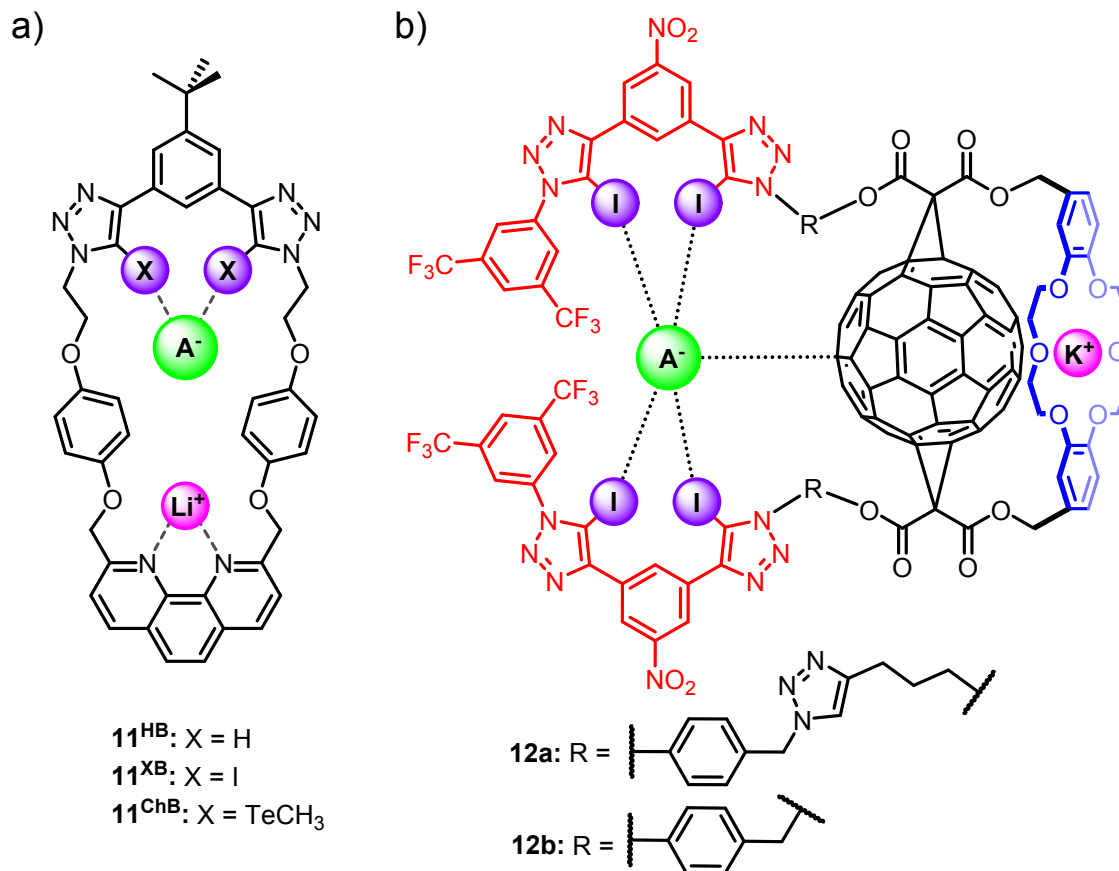


Figure 9. Non-interlocked heteroditopic incorporating XB anion binding motifs, including a) Tse's phenanthroline-functionalised macrocycles **11**, and b) Bak's fullerene-containing receptor **12a-b**.



Importantly, the ion-pair binding affinities and preferences of the macrocycles could be tuned by varying the nature of the anion binding interaction. **7^{XB}** showed overall strongest binding affinity for all LiX ion-pairs (X = Cl, Br, I) and is notably able to stabilise the 'hard' LiCl ion-pair. In contrast, **11^{ChB}** was unable to bind LiCl and instead demonstrates a marked preference for the 'soft' LiI ion-pair.

Another example of integrating XB...anion interactions into a macrocyclic scaffold for alkali metal halide ion-pair recognition was reported by Bak and co-workers,¹⁰⁵ who synthesised fullerene-containing heteroditopic receptors **12a-b** utilising iodotriazole XB donors as the anion binding motif and dibenzo-18-crown-6 ether as the cation binding group, wherein importantly the cation and anion binding sites are spatially separated by a C₆₀ unit (Figure 9b-c). The neutral receptors **12a-b** displayed enhanced XB-mediated anion binding relative to a non-fullerene analogue in 3:1 CD₃CN/CDCl₃, attributed to a combination of fullerene-mediated pre-organisation, solvent shielding and the formation of π -hole interactions with the polarisable C₆₀ surface. Pre-complexation of a potassium cation to the crown ether unit significantly enhanced the halide binding affinities of **12a-b** through positive electrostatic cooperativity associated with the increased polarisation of the C₆₀ surface by the co-bound cation. The important contribution from C₆₀ polarisation was reflected in the marked binding preference of the potassium bound receptor complex **12-K⁺** for the softer iodide ion ($K_a(\text{I}^-) = 36000 \text{ M}^{-1}$ (**12a**), 19800 M⁻¹ (**12b**)) over bromide ($K_a(\text{Br}^-) = 25600 \text{ M}^{-1}$ (**12a**), 8300 M⁻¹ (**12b**)) and chloride ($K_a(\text{Cl}^-) = 15100 \text{ M}^{-1}$ (**12a**), 9300 M⁻¹ (**12b**)). Interestingly, detailed analysis of the ¹H NMR spectral data suggests that a significant proportion (40%) of the chloride guest is directly coordinated to the potassium cation in a contact ion-pair binding fashion, and does not benefit from the polarisation-enhanced XB interactions, accounting for the halide's weaker binding to the receptor. These studies thereby highlight the potential of integrating XB interactions into heteroditopic receptor design to elicit nuanced selectivity effects in ion-pair binding.

MIM Heteroditopic Receptors Utilising σ -Hole Interactions

To the best of our knowledge, the first instance of a heteroditopic [2]rotaxane-based ion-pair receptor **13** employing XB donor groups for anion binding was reported by our group in 2022.¹⁰⁶ The rotaxane design consists of a bis(iodotriazole)-functionalised macrocycle as well as an iodotriazole-functionalised axle component (Figure 10a), which was prepared *via* a CuAAC active metal template reaction. Quantitative ¹H NMR studies showed that **13** was capable of binding lithium bromide and iodide ion-pairs in CDCl₃/CD₃CN mixtures, an impressive feat considering the high lattice enthalpies of lithium halide salts. Crystallographic and DFT modelling studies indicated that the interlocked nature of **13** was pivotal in creating the requisite cation and anion binding sites. Cation binding occurs *via* the 2,6-dialkoxypyridyl motif of the macrocycle and the triazole *N*-donors of the axle, which forms a binding cavity complementary in size to the small Li⁺ cation. Li⁺ complexation within this cavity pre-organises the iodotriazole XB motifs of the macrocycle and axle to participate in convergent anion binding, giving rise to a positive allosteric effect that contributes to favourable cooperativity in ion-pair binding. Importantly, the non-interlocked axle and macrocycle components were unable to bind lithium halide ion-pairs. The selectivity of the rotaxanes for binding lithium halide ion-pairs is particularly interesting given the widespread adoption of lithium-based battery technologies,¹⁰⁷ which has in turn driven efforts to develop efficient methods for the recovery and remediation of lithium salts.

Subsequently, related rotaxanes **14^{HB}** and **14^{XB}** employing a similar design concept, with an isophthalamide-based macrocycle and an axle containing a prototriazole (**14^{HB}**) or iodotriazole (**14^{XB}**) motif, displayed remarkably selective binding of lithium chloride over other alkali metal halides (Figure 10b). Interestingly, ¹H NMR, crystallographic and DFT computational studies revealed the free rotaxanes adopted a co-conformation which allows the axle triazole nitrogen atoms to participate in intramolecular hydrogen bonding interactions

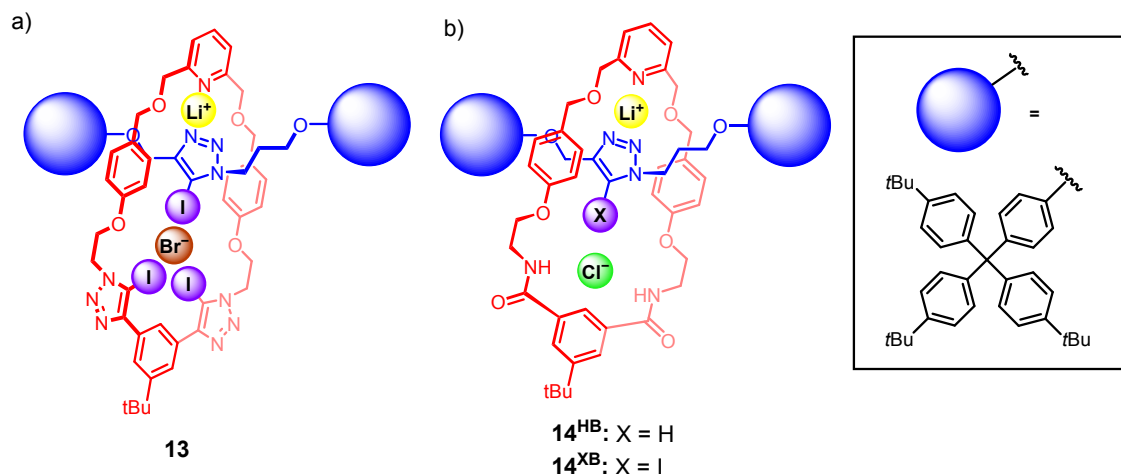


Figure 102. Beer's heteroditopic XB [2]rotaxanes capable of cooperatively binding LiX ion-pairs, including: a) rotaxane **13** which preferentially bind LiBr and LiI ion-pairs; b) rotaxanes **14^{HB}** and **14^{XB}** which are capable of stabilising LiCl ion-pairs.



ARTICLE

Journal Name

with the isophthalamide motifs. Binding of Li^+ induces a re-orientation of the triazole to create pre-organised cation and anion binding cavities, thereby exploiting the co-conformational dynamism of the MIM architecture to achieve a positive cooperative allosteric enhancement of anion binding upon Li^+ pre-complexation.

Integrating XB donor motifs into pre-organised catenane-based host architectures proved to be a promising strategy for modulating the binding affinities and selectivity profiles of ion-pair binding (**Figure 11**). A series of neutral all-XB homo[2]catenanes **15**^{DEG} and **15**^{TEG} were prepared *via* an elegant template-directed synthetic strategy, in which a sodium

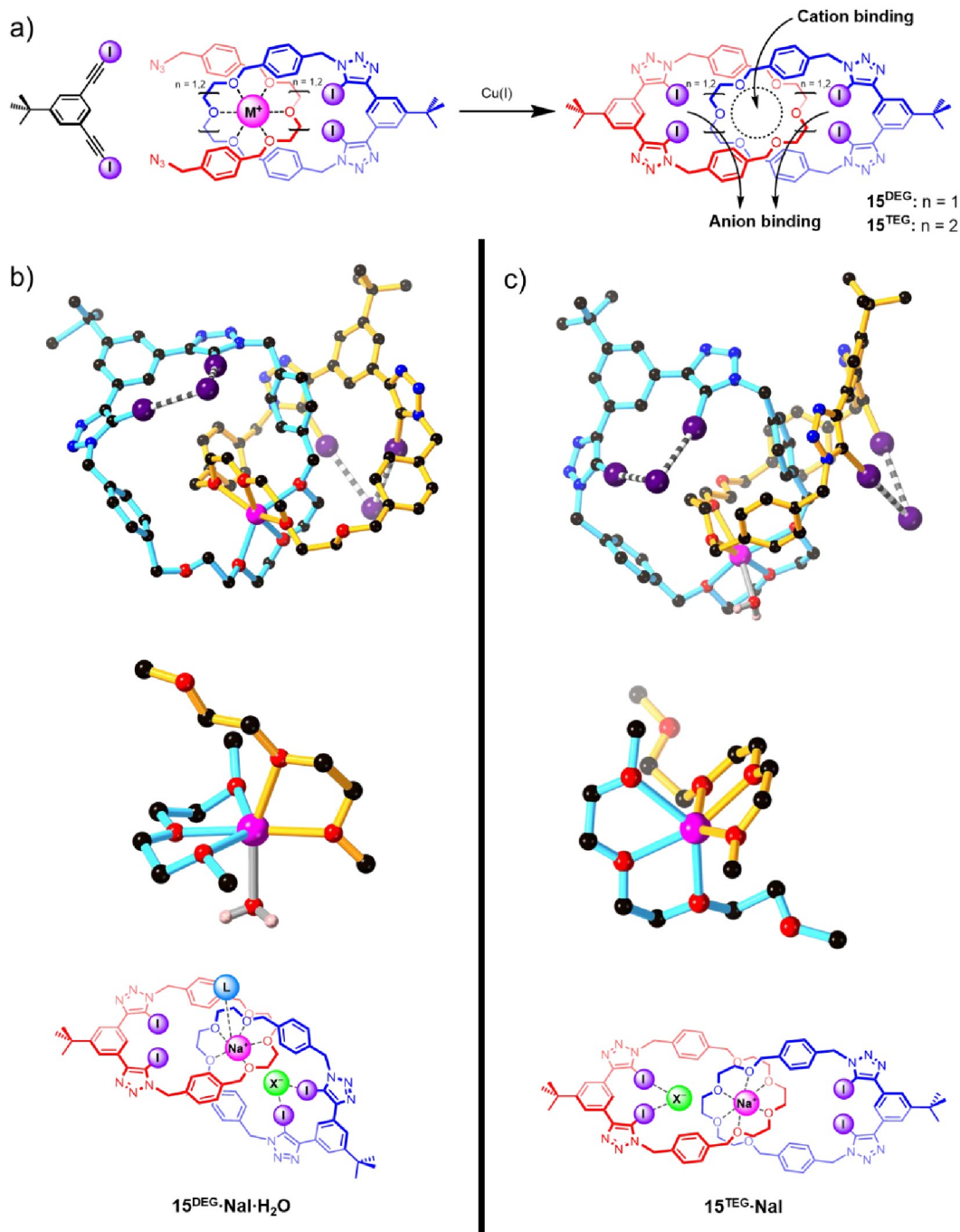


Figure 11. a) Sodium cation template-directed synthesis of heteroditopic all-XB [2]catenanes **15** for alkali metal halide ion-pair binding; Solid-state structures of (b) **15**^{DEG}·NaI·H₂O and (c) **15**^{TEG}·NaI, showing a single [2]catenane unit bound to a sodium cation and two iodide anions (middle) and a truncated representation of the ethylene glycol region (bottom) highlighting the coordination environment around the sodium ion. Atom colours are as follows: black (carbon), blue (nitrogen), red (oxygen), purple (iodine), light pink (hydrogen), bright pink (sodium). The chemical structures of **15**^{DEG} and **15**^{TEG} are shown on the left for reference.



cation directs the formation of pseudo-[2]rotaxane complex between an oligo(ethylene glycol)-functionalised XB macrocycle and a bis-azide. The acyclic bis-azide precursor subsequently underwent a CuAAC-mediated macrocyclisation reaction with a bis-iodoalkyne to form a second interlocked macrocycle containing a bidentate XB donor motif. The mechanical bond formation step thereby simultaneously generated both cation and anion recognition sites (**Figure 11a**).

The resulting homo[2]catenanes exhibited moderate affinity for sodium and potassium cations, with 1:1 host-guest stoichiometric binding constants (K_a) in the range 700-1800 M^{-1} . Notably, the smaller di(ethylene glycol)-based catenane **15^{DEG}** preferentially binds sodium over potassium and vice versa for the larger tri(ethylene glycol)-based catenane **15^{TEG}**, reflecting the size complementarity between the interlocked cation binding site and the cation guests. No metal cation binding was observed for the non-interlocked macrocycles, attesting to the importance of the mechanical bond effect (MBE) in switching on the cation affinity of the host. The modest halide affinities of the free catenanes ($K_a = 100-200 M^{-1}$) were found to be significantly enhanced by pre-complexation of a sodium or potassium cation, reflecting the positive electrostatic cooperativity in ion-pair binding. Interestingly, a detailed analysis of the 1H NMR titration isotherms showed that depending on the identity of the catenane and the pre-complexed metal ion, the halides were bound in either a 1:1 or 1:2 host-guest stoichiometry. XRD crystallographic analysis revealed that, in addition to the anticipated XB-mediated anion binding, the smaller [2]catenane **15^{DEG}** can also adopt a direct contact ion-pair binding mode in which an anion guest coordinates to the unsaturated Na^+ centre (**Figure 11b**), giving rise to an overall 1:1:2 host-cation-anion stoichiometry. In contrast, the larger [2]catenane **15^{TEG}** exhibits a 1:1:1 host-cation-anion binding stoichiometry as its coordinatively saturated Na^+ cation guest precludes direct contact ion-pair binding (**Figure 11c**).¹⁰⁸

While the homo[2]catenane systems highlighted the potential of exploiting the highly pre-organised binding cavities of catenanes to elicit ion-pair binding, the relatively moderate individual cation and anion affinities of **15^{DEG}** and **15^{TEG}** precluded strong ion-pair binding. Building on this work, a heteroditopic [2]catenane ion-pair host system **16** containing a more potent bis-di(ethylene glycol) cation binding site was synthesised (**Figure 12**), which demonstrated a dramatic

increase in affinity for sodium and potassium cations compared to homo[2]catenane systems, as well as marked sodium over potassium selectivity arising from the size complementarity between the di(ethylene glycol)-based cation binding site and the smaller sodium cation. Similar to the homo[2]catenanes, ion-pair titrations conducted on **16** showed that its anion binding affinity could be 'switched on' by pre-complexation with a metal cation. Notably, **16** displayed significantly stronger binding of alkali metal halide salts compared to the homo[2]catenanes, indicating that the modified host design was effective in enhancing the ion-pair affinity of the interlocked receptor. In addition, the binding preference for sodium halide over potassium halide ion-pairs was maintained, with **16** displaying 3 to 5-fold stronger binding to NaX ion-pairs ($X = Br, I$) relative to the corresponding KX salts. This binding preference was impressively translated to the solid-liquid extraction properties of the catenane, with competitive SLE experiments indicating preferential extraction of sodium halide salts from a solid mixture of NaX/KX salts ($X = Br, I$).¹⁰⁹

To the best of our knowledge, comparative studies of 'mechanical bond isomers' for ion-pair binding and recognition remain unprecedented to date. This is somewhat surprising considering the different co-conformational behaviours of rotaxanes and catenanes and the potential effects on their recognition properties. As such, encouraged by the promising ion-pair binding properties of the catenane-based heteroditopic receptors, attention was turned to investigating their [2]rotaxane analogues (**Figure 13a**). To this end, a series of heteroditopic [2]rotaxanes **17a-d** were prepared *via* a modified sodium cation template-directed methodology, in which a stoppering-type reaction was performed on the intermediate pseudo[2]rotaxane assembly using sterically bulky iodoalkyne-functionalised terphenyl groups. Notably, 1H NMR binding studies revealed that metal cation binding to the rotaxanes induces a translocation of the macrocycle component towards the central ethylene glycol stations to facilitate convergent metal coordination by the polyether binding motifs of the macrocycle and axle. The need for a large-amplitude co-conformational change led to reduced cation binding affinities of the [2]rotaxanes relative to the analogous homo[2]catenanes, which in turn gave rise to complex parallel and competing equilibria in the ion-pair binding studies that complicated the quantitative determination of ion-pair binding affinities. Importantly, comparing the apparent alkali metal

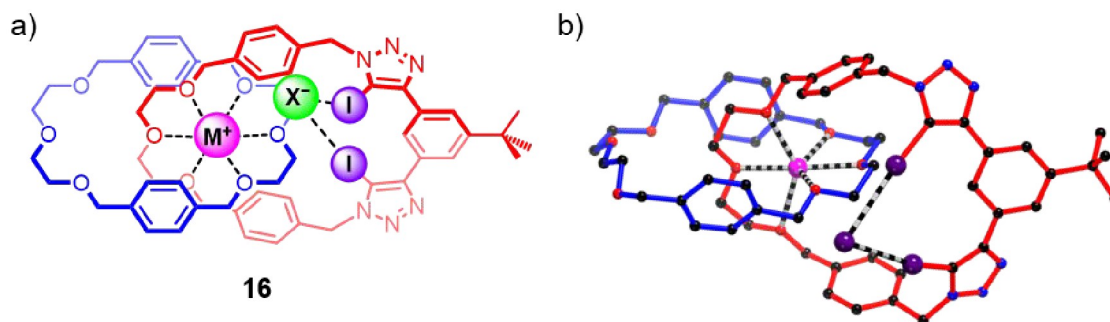


Figure 12. A heteroditopic XB [2]catenane **16** showing enhanced alkali metal halide ion-pair binding and extraction properties.



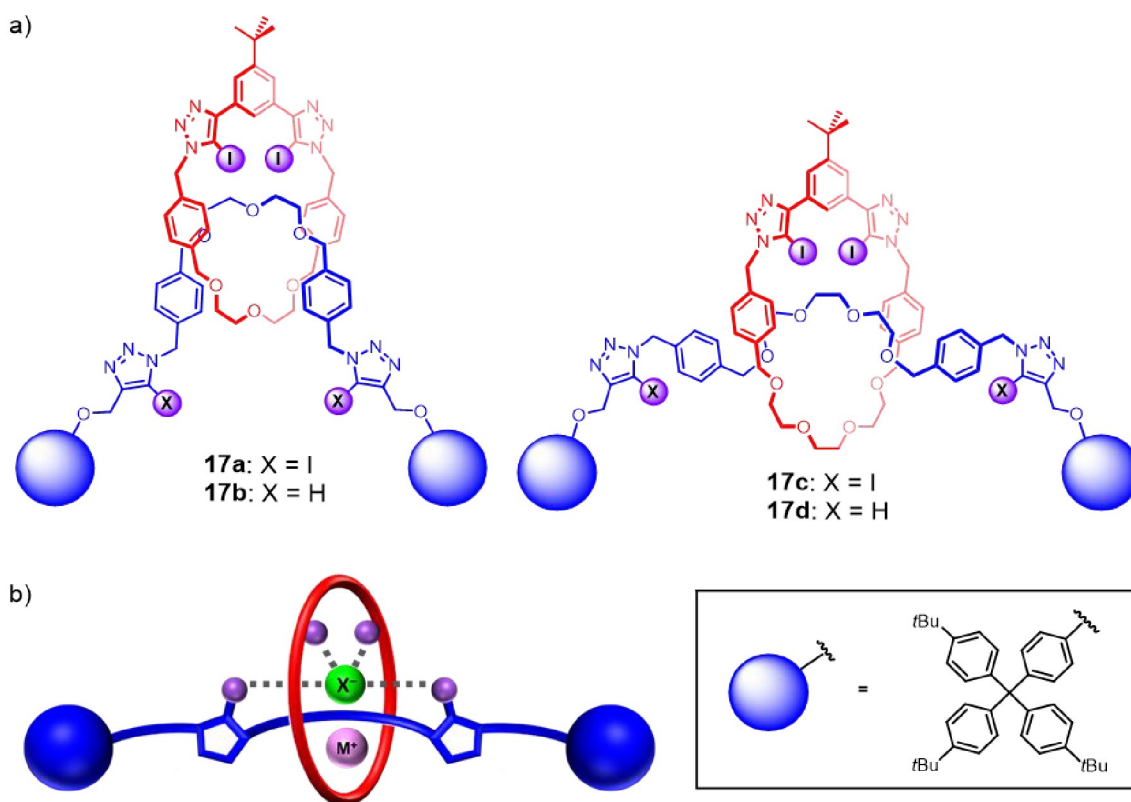


Figure 13. a) Heteroditopic HB or XB rotaxanes **17a-d** demonstrating cooperative binding of alkali metal halide ion-pairs; b) Proposed ion-pair binding mode of heteroditopic rotaxanes.

cation iodide ion-pair binding constants of the [2]rotaxanes and the [2]catenane analogues revealed that the rotaxanes exhibited a higher degree of positive cooperativity in ion-pair binding despite weaker cation binding (**Figure 13b**), suggesting the advantages of incorporating mechanical bond co-conformational flexibility into the design of future heteroditopic interlocked receptors.¹¹⁰

Conclusions

In the pursuit of novel heteroditopic receptors capable of improved ion-pair binding affinity and selectivity, the use of mechanically interlocked host topologies is emerging as a promising avenue of research. Although efforts to incorporate the mechanical bond into heteroditopic host design has thus far remained limited, the examples discussed in this review importantly highlight significant binding affinity enhancements and positive cooperativity in ion-pair recognition by MIM-based heteroditopic receptors relative to non-interlocked acyclic and macrocyclic analogues. Crucially, in many cases, MIM-based heteroditopic receptors leverage not only favourable proximal cation-anion bound electrostatic effects but also their inherent co-conformational dynamism to achieve allosteric cooperativity, wherein the binding of one guest ion induces a co-conformational rearrangement of the mechanically interlocked components, including macrocycle translocation, pirouetting or inter-ring circum-rotation, to create a pre-

organised binding cavity suited for binding to the second guest counter-ion of the ion-pair. Impressively, chemically similar alkali metal cations or halide anions can often be distinguished by varying the nature of the cation/anion binding motifs and the size of the interlocked cavity, which serves as a powerful means to achieve highly selective ion-pair binding. Notably, recent advancements in the incorporation of halogen bonding (XB) anion recognition motifs into MIM receptor scaffolds have enabled further modulation of ion-pair binding affinity and selectivity profiles.

Given the relatively small number of heteroditopic MIM receptors reported to date, future work in this area is expected to initially focus on laying the groundwork for this expanding field. From a synthetic perspective, this would entail exploiting and combining emerging classes of non-covalent interactions, such as chalcogen bonding and anion- π interactions in innovative host designs. Equally important is the development of robust analytical methods to quantify the strength and cooperativity of ion-pair binding, which would enable systematic comparisons to be made between various receptor systems. Combining these synthetic and analytical tools to evaluate the cation, anion and electrostatic/allosteric cooperative ion-pair binding properties would elucidate guiding principles to achieve selective ion-pair recognition in future strategic MIM heteroditopic host design.

Further progress in the field would encourage the construction of functional MIM-based ion-pair receptors for a



diverse range of applications. For example, MIM-based heteroditopic architectures are particularly suited for use as ion-pair sensors, as their high binding affinity and selectivity for target guests allows them to function at low concentrations. Initially, sensors can be prepared in a relatively facile manner via the integration of fluorophoric or redox-active reporter groups into established heteroditopic MIM receptor scaffolds. Additionally, the inherent co-conformational dynamism of MIMs is a powerful tool for the development of novel sensing mechanisms for ion-pairs, as has been demonstrated in a series of reports featuring chromophore-functionalised rotaxane shuttles which utilise anion recognition induced macrocycle translocation to elicit a selective optical response to specific anionic guest species.^{48, 111-115} There is also potential for utilising MIM-based heteroditopic receptors in selective ion-pair transport or extraction. Finally, the scope for guest binding may further expand from simple inorganic ion-pairs to the recognition of complex guest species such as biological zwitterions including amino acids, where chiral heteroditopic MIM structural host frameworks can be employed.¹¹⁶⁻¹¹⁸

Data Availability Statement

No primary research results, software or code have been included and no new data were generated or analysed as part of this review.

Conflicts of interest

There are no conflicts to declare.

Acknowledgements

H.M.T. acknowledges the Clarendon Fund and the Oxford Australia Scholarships Fund for a postgraduate research scholarship.

Notes and references

**Dedicated to Professor Stephen J. Loeb on the occasion of his 70th birthday

- 1 D. Banerjee, D. Kim, M. J. Schweiger, A. A. Kruger and P. K. Thallapally, *Chem. Soc. Rev.*, 2016, **45**, 2724-2739.
- 2 M. D. Rao, K. K. Singh, C. A. Morrison and J. B. Love, *RSC Adv.*, 2020, **10**, 4300-4309.
- 3 M. P. Anderson, R. J. Gregory, S. Thompson, D. W. Souza, S. Paul, R. C. Mulligan, A. E. Smith and M. J. Welsh, *Science*, 1991, **253**, 202.
- 4 F. Delange, *Thyroid*, 1994, **4**, 107-128.
- 5 J. L. Way, *Annu. Rev. Pharmacol. Toxicol.*, 1984, **24**, 451-481.
- 6 V. H. Smith and D. W. Schindler, *Trends. Ecol. Evol.*, 2009, **24**, 201-207.
- 7 J. H. P. Watson and D. C. Ellwood, *Nucl. Eng. Des.*, 2003, **226**, 375-385.
- 8 P. K. Dasgupta, J. V. Dyke, A. B. Kirk and W. A. Jackson, *Environ. Sci. Technol.*, 2006, **40**, 6608-6614.

- 9 N. H. Evans and P. D. Beer, *Angew. Chem. Int. Ed.*, 2014, **53**, 11716-11754.
- 10 F. P. Schmidtchen, *Angew. Chem. Int. Ed. Engl.*, 1977, **16**, 720-721.
- 11 E. Graf and J. M. Lehn, *J. Am. Chem. Soc.*, 1976, **98**, 6403-6405.
- 12 J. M. Lehn, E. Sonveaux and A. K. Willard, *J. Am. Chem. Soc.*, 1978, **100**, 4914-4916.
- 13 G. Müller, J. Riede and F. P. Schmidtchen, *Angew. Chem. Int. Ed. Engl.*, 1988, **27**, 1516-1518.
- 14 K. Chellappan, N. J. Singh, I.-C. Hwang, J. W. Lee and K. S. Kim, *Angew. Chem. Int. Ed.*, 2005, **44**, 2899-2903.
- 15 A. Kumar and P. S. Pandey, *Org. Lett.*, 2008, **10**, 165-168.
- 16 J. T. Davis, P. A. Gale and R. Quesada, *Chem. Soc. Rev.*, 2020, **49**, 6056-6086.
- 17 N. Akhtar, O. Biswas and D. Manna, *Chem. Commun.*, 2020, **56**, 14137-14153.
- 18 K. M. Båk, K. Porfyrakis, J. J. Davis and P. D. Beer, *Mater. Chem. Front.*, 2020, **4**, 1052-1073.
- 19 J. F. Stoddart, *Chem. Soc. Rev.*, 2009, **38**, 1802-1820.
- 20 E. A. Neal and S. M. Goldup, *Chem. Commun.*, 2014, **50**, 5128-5142.
- 21 M. Xue, Y. Yang, X. Chi, X. Yan and F. Huang, *Chem. Rev.*, 2015, **115**, 7398-7501.
- 22 G. Gil-Ramírez, D. A. Leigh and A. J. Stephens, *Angew. Chem. Int. Ed.*, 2015, **54**, 6110-6150.
- 23 J. E. M. Lewis, M. Galli and S. M. Goldup, *Chem. Commun.*, 2017, **53**, 298-312.
- 24 W.-X. Gao, H.-J. Feng, B.-B. Guo, Y. Lu and G.-X. Jin, *Chem. Rev.*, 2020, **120**, 6288-6325.
- 25 H.-N. Zhang, Y.-J. Lin and G.-X. Jin, *J. Am. Chem. Soc.*, 2021, **143**, 1119-1125.
- 26 J. T. Wilmore and P. D. Beer, *Adv. Mater.*, 2024, **36**, 2309098.
- 27 G. Baggi and S. J. Loeb, *Chem. Eur. J.*, 2017, **23**, 14163-14166.
- 28 M. Nandi, S. Bej, T. K. Ghosh and P. Ghosh, *Chem. Commun.*, 2019, **55**, 3085-3088.
- 29 S.-M. Chan, F.-K. Tang, C.-S. Kwan, C.-Y. Lam, S. C. K. Hau and K. C.-F. Leung, *Mater. Chem. Front.*, 2019, **3**, 2388-2396.
- 30 T. Shukla, A. K. Dwivedi, R. Arumugaperumal, C.-M. Lin, S.-Y. Chen and H.-C. Lin, *Dyes Pigm.*, 2016, **131**, 49-59.
- 31 M. Denis, J. Pancholi, K. Jobe, M. Watkinson and S. M. Goldup, *Angew. Chem. Int. Ed.*, 2018, **57**, 5310-5314.
- 32 A. Caballero, F. Zapata and P. D. Beer, *Coord. Chem. Rev.*, 2013, **257**, 2434-2455.
- 33 R. J. Goodwin, A. Docker, H. I. MacDermott-Opeskin, H. M. Aitken, M. L. O'Mara, P. D. Beer and N. G. White, *Chem. Eur. J.*, 2022, **28**, e202200389.
- 34 J. M. Mercurio, F. Tyrrell, J. Cookson and P. D. Beer, *Chem. Commun.*, 2013, **49**, 10793-10795.
- 35 M. J. Langton, O. A. Blackburn, T. Lang, S. Faulkner and P. D. Beer, *Angew. Chem. Int. Ed.*, 2014, **53**, 11463-11466.
- 36 N. G. White and P. D. Beer, *Org. Biomol. Chem.*, 2013, **11**, 1326-1333.
- 37 Y. Cheong Tse, R. Hein, E. J. Mitchell, Z. Zhang and P. D. Beer, *Chem. Eur. J.*, 2021, **27**, 14550-14559.
- 38 J.-P. Sauvage, *Angew. Chem. Int. Ed.*, 2017, **56**, 11080-11093.
- 39 F. Bitsch, C. O. Dietrich-Buchecker, A. K. Khemiss, J. P. Sauvage and A. Van Dorsselaer, *J. Am. Chem. Soc.*, 1991, **113**, 4023-4025.
- 40 C. O. Dietrich-Buchecker, C. Hemmert, A. K. Khemiss and J. P. Sauvage, *J. Am. Chem. Soc.*, 1990, **112**, 8002-8008.
- 41 C. O. Dietrich-Buchecker and J.-P. Sauvage, *Angew. Chem. Int. Ed. Engl.*, 1989, **28**, 189-192.
- 42 J. Guilhem, C. Pascard, J. P. Sauvage and J. Weiss, *J. Am. Chem. Soc.*, 1988, **110**, 8711-8713.
- 43 J. P. Sauvage and J. Weiss, *J. Am. Chem. Soc.*, 1985, **107**, 6108-6110.
- 44 C. O. Dietrich-Buchecker, J. P. Sauvage and J. M. Kern, *J. Am. Chem. Soc.*, 1984, **106**, 3043-3045.



ARTICLE

Journal Name

- 45 M. Cirulli, A. Kaur, J. E. M. Lewis, Z. Zhang, J. A. Kitchen, S. M. Goldup and M. M. Roessler, *J. Am. Chem. Soc.*, 2019, **141**, 879-889.
- 46 A. Docker, Y. C. Tse, H. M. Tay, A. J. Taylor, Z. Zhang and P. D. Beer, *Angew. Chem. Int. Ed.*, 2022, **61**, e202214523.
- 47 H. M. Tay, T. G. Johnson, A. Docker, M. J. Langton and P. D. Beer, *Angew. Chem. Int. Ed.*, 2023, **62**, e202312745.
- 48 H. M. Tay, A. Docker, A. Taylor and P. D. Beer, *Chem. Eur. J.*, 2024, **30**, e202400952.
- 49 K. M. Bąk, B. Trzaskowski and M. J. Chmielewski, *Chem. Sci.*, 2024, **15**, 1796-1809.
- 50 S. K. Kim and J. L. Sessler, *Chem. Soc. Rev.*, 2010, **39**, 3784-3809.
- 51 A. J. McConnell, A. Docker and P. D. Beer, *ChemPlusChem*, 2020, **85**, 1824-1841.
- 52 P. D. Beer, P. K. Hopkins and J. D. McKinney, *Chem. Commun.*, 1999, 1253-1254.
- 53 M. Zakrzewski, D. Załubiniak and P. Piątek, *Dalton Trans.*, 2018, **47**, 323-330.
- 54 Q. He, G. M. Peters, V. M. Lynch and J. L. Sessler, *Angew. Chem. Int. Ed.*, 2017, **56**, 13396-13400.
- 55 S. G. Galbraith, P. G. Plieger and P. A. Tasker, *Chem. Commun.*, 2002, 2662-2663.
- 56 T. Bunchuay, A. Docker, U. Eiamprasert, P. Surawatanawong, A. Brown and P. D. Beer, *Angew. Chem. Int. Ed.*, 2020, **59**, 12007-12012.
- 57 Y. C. Tse, A. Docker, Z. Zhang and P. D. Beer, *Chem. Commun.*, 2021, **57**, 4950-4953.
- 58 A. Docker, J. G. Stevens and P. D. Beer, *Chem. Eur. J.*, 2021, **27**, 14600-14604.
- 59 A. Docker, T. Bunchuay, M. Ahrens, A. J. Martinez-Martinez and P. D. Beer, *Chem. Eur. J.*, 2021, **27**, 7837-7841.
- 60 A. Docker, I. Marques, H. Kuhn, Z. Zhang, V. Félix and P. D. Beer, *J. Am. Chem. Soc.*, 2022, **144**, 14778-14789.
- 61 Q. He, N. J. Williams, J. H. Oh, V. M. Lynch, S. K. Kim, B. A. Moyer and J. L. Sessler, *Angew. Chem. Int. Ed.*, 2018, **57**, 11924-11928.
- 62 Q. He, Z. Zhang, J. T. Brewster, V. M. Lynch, S. K. Kim and J. L. Sessler, *J. Am. Chem. Soc.*, 2016, **138**, 9779-9782.
- 63 A. V. Koulov, J. M. Mahoney and B. D. Smith, *Org. Biomol. Chem.*, 2003, **1**, 27-29.
- 64 J. M. Mahoney, G. U. Nawaratna, A. M. Beatty, P. J. Duggan and B. D. Smith, *Inorg. Chem.*, 2004, **43**, 5902-5907.
- 65 A. Galan, D. Andreu, A. M. Echavarren, P. Prados and J. De Mendoza, *J. Am. Chem. Soc.*, 1992, **114**, 1511-1512.
- 66 W. Walczak, M. Zakrzewski, G. Cichowicz and P. Piątek, *Org. Biomol. Chem.*, 2020, **18**, 694-699.
- 67 R. C. Knighton and P. D. Beer, *Chem. Commun.*, 2014, **50**, 1540-1542.
- 68 A. Brown, K. M. Mennie, O. Mason, N. G. White and P. D. Beer, *Dalton Trans.*, 2017, **46**, 13376-13385.
- 69 M. J. Deetz, R. Shukla and B. D. Smith, *Tetrahedron*, 2002, **58**, 799-805.
- 70 J. R. Romero, G. Aragay and P. Ballester, *Chem. Sci.*, 2017, **8**, 491-498.
- 71 M. Denis, L. Qin, P. Turner, K. A. Jolliffe and S. M. Goldup, *Angew. Chem. Int. Ed.*, 2018, **57**, 5315-5319.
- 72 D.-H. Li and B. D. Smith, *J. Org. Chem.*, 2019, **84**, 2808-2816.
- 73 A. J. McConnell and P. D. Beer, *Angew. Chem. Int. Ed.*, 2012, **51**, 5052-5061.
- 74 J. L. Sessler, S. K. Kim, D. E. Gross, C.-H. Lee, J. S. Kim and V. M. Lynch, *J. Am. Chem. Soc.*, 2008, **130**, 13162-13166.
- 75 J. Scheerder, J. P. M. van Duynhoven, J. F. J. Engbersen and D. N. Reinhoudt, *Angew. Chem. Int. Ed. Engl.*, 1996, **35**, 1090-1093.
- 76 A. J. Evans and P. D. Beer, *Dalton Trans.*, 2003, 4451-4456.
- 77 A. J. Taylor, A. Docker and P. D. Beer, *Chem. Asian J.*, 2023, **18**, e202201170.
- 78 M. J. Deetz, M. Shang and B. D. Smith, *J. Am. Chem. Soc.*, 2000, **122**, 6201-6207.
- 79 J. M. Mahoney, A. M. Beatty and B. D. Smith, *J. Am. Chem. Soc.*, 2001, **123**, 5847-5848.
- 80 Y. Yeon, S. Leem, C. Wagen, V. M. Lynch, S. K. Kim and J. L. Sessler, *Org. Lett.*, 2016, **18**, 4396-4399.
- 81 B. Akhuli and P. Ghosh, *Chem. Commun.*, 2015, **51**, 16514-16517.
- 82 D. Jaglenieć, S. Siennicka, Ł. Dobrzycki, M. Karbarz and J. Romański, *Inorg. Chem.*, 2018, **57**, 12941-12952.
- 83 P. R. A. Webber and P. D. Beer, *Dalton Trans.*, 2003, 2249-2252.
- 84 Z.-H. Sun, F.-F. Pan, Triyanti, M. Albrecht and G. Raabe, *Eur. J. Org. Chem.*, 2013, **2013**, 7922-7932.
- 85 J. Romański and P. Piątek, *J. Org. Chem.*, 2013, **78**, 4341-4347.
- 86 J. H. Lee, J. H. Lee, Y. R. Choi, P. Kang, M.-G. Choi and K.-S. Jeong, *J. Org. Chem.*, 2014, **79**, 6403-6409.
- 87 D.-y. Duan, L. L. H. Liu, N. Bozeat, Z. M. Huang, S. Y. Xiang, G.-I. Wang, L. Ye and J. R. Hume, *Acta Pharmacol. Sin.*, 2005, **26**, 265-278.
- 88 L. Puljak and G. Kilic, *Biochimica et Biophysica Acta (BBA) - Molecular Basis of Disease*, 2006, **1762**, 404-413.
- 89 N. Carrasco, *Biochimica et Biophysica Acta (BBA) - Reviews on Biomembranes*, 1993, **1154**, 65-82.
- 90 X. Zheng, Z. Zhu, X. Lin, Y. Zhang, Y. He, H. Cao and Z. Sun, *Engineering*, 2018, **4**, 361-370.
- 91 M. Kamenica, R. R. Kothur, A. Willows, B. A. Patel and P. J. Cragg, *Journal*, 2017, **17**.
- 92 R. C. Knighton and P. D. Beer, *Org. Chem. Front.*, 2021, **8**, 2468-2472.
- 93 S. K. Kim and J. L. Sessler, *Acc. Chem. Res.*, 2014, **47**, 2525-2536.
- 94 V. Valderrey, E. C. Escudero-Adán and P. Ballester, *J. Am. Chem. Soc.*, 2012, **134**, 10733-10736.
- 95 A. Arun, A. Docker, H. Min Tay and P. D. Beer, *Chem. Eur. J.*, 2023, **29**, e202301446.
- 96 Y.-J. Lee, T.-H. Ho, C.-C. Lai and S.-H. Chiu, *Org. Biomol. Chem.*, 2016, **14**, 1153-1160.
- 97 L. C. Gilday, S. W. Robinson, T. A. Barendt, M. J. Langton, B. R. Mullaney and P. D. Beer, *Chem. Rev.*, 2015, **115**, 7118-7195.
- 98 J. Y. C. Lim and P. D. Beer, *Chem*, 2018, **4**, 731-783.
- 99 J. Pancholi and P. D. Beer, *Coord. Chem. Rev.*, 2020, **416**, 213281.
- 100 A. Docker and P. D. Beer, in *Halogen Bonding in Solution*, 2021, pp. 83-120.
- 101 T. Bunchuay, A. Docker, A. J. Martinez-Martinez and P. D. Beer, *Angew. Chem. Int. Ed.*, 2019, **58**, 13823-13827.
- 102 G. Turner, A. Docker and P. D. Beer, *Dalton Trans.*, 2021, **50**, 12800-12805.
- 103 A. Mele, P. Metrangolo, H. Neukirch, T. Pilati and G. Resnati, *J. Am. Chem. Soc.*, 2005, **127**, 14972-14973.
- 104 R. Tepper, B. Schulze, P. Bellstedt, J. Heidler, H. Görls, M. Jäger and U. S. Schubert, *Chem. Commun.*, 2017, **53**, 2260-2263.
- 105 K. M. Bąk, I. Marques, H. Kuhn, K. E. Christensen, V. Félix and P. D. Beer, *J. Am. Chem. Soc.*, 2023, **145**, 27367-27379.
- 106 V. K. Munasinghe, J. Pancholi, D. Manawadu, Z. Zhang and P. D. Beer, *Chem. Eur. J.*, 2022, **28**, e202201209.
- 107 G. Martin, L. Rentsch, M. Höck and M. Bertau, *Energy Storage Mater.*, 2017, **6**, 171-179.
- 108 H. M. Tay, Y. C. Tse, A. Docker, C. Gateley, A. L. Thompson, H. Kuhn, Z. Zhang and P. D. Beer, *Angew. Chem. Int. Ed.*, 2023, **62**, e202214785.
- 109 H. M. Tay, A. Docker, C. Hua and P. D. Beer, *Chem. Sci.*, 2024, **15**, 13074-13081.
- 110 H. M. Tay, A. Docker, Y. Cheong Tse and P. D. Beer, *Chem. Eur. J.*, 2023, **29**, e202301316.



Journal Name

ARTICLE

- 111 J. J. Gassensmith, S. Matthys, J.-J. Lee, A. Wojcik, P. V. Kamat and B. D. Smith, *Chem. Eur. J.*, 2010, **16**, 2916-2921.
- 112 T. A. Barendt, S. W. Robinson and P. D. Beer, *Chem. Sci.*, 2016, **7**, 5171-5180.
- 113 T. A. Barendt, A. Docker, I. Marques, V. Félix and P. D. Beer, *Angew. Chem. Int. Ed.*, 2016, **55**, 11069-11076.
- 114 T. A. Barendt, L. Ferreira, I. Marques, V. Félix and P. D. Beer, *J. Am. Chem. Soc.*, 2017, **139**, 9026-9037.
- 115 T. A. Barendt, I. Rašović, M. A. Lebedeva, G. A. Farrow, A. Auty, D. Chekulaev, I. V. Sazanovich, J. A. Weinstein, K. Porfyrakis and P. D. Beer, *J. Am. Chem. Soc.*, 2018, **140**, 1924-1936.
- 116 J. Y. C. Lim, I. Marques, V. Félix and P. D. Beer, *Angew. Chem. Int. Ed.*, 2018, **57**, 584-588.
- 117 R. L. Spicer, C. C. Shearman and N. H. Evans, *Chem. Eur. J.*, 2023, **29**, e202203502.
- 118 J. Y. C. Lim, I. Marques, V. Félix and P. D. Beer, *J. Am. Chem. Soc.*, 2017, **139**, 12228-12239.



Data Availability Statement for: Mechanically Interlocked Host Systems for Ion-Pair Recognition

View Article Online
DOI: 10.1039/D4CC03916E

No primary research results, software or code have been included and no new data were generated or analysed as part of this review.

