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Cryptophane–xenon complexes for ^{129}Xe MRI applications

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The use of magnetic resonance imaging (MRI) and spectroscopy (MRS) in the clinical setting enables the acquisition of valuable anatomical information in a rapid, non-invasive fashion. However, MRI applications for identifying disease-related biomarkers are limited due to low sensitivity at clinical magnetic field strengths. The development of hyperpolarized (hp) ^{129}Xe MRI/MRS techniques as complements to traditional ^1H -based imaging has been a burgeoning area of research over the past two decades. Pioneering experiments have shown that hp ^{129}Xe can be encapsulated within host molecules to generate ultrasensitive biosensors. In particular, xenon has high affinity for cryptophanes, which are small organic cages that can be functionalized with affinity tags, fluorophores, solubilizing groups, and other moieties to identify biomedically relevant analytes. Cryptophane sensors designed for proteins, metal ions, nucleic acids, pH, and temperature have achieved nanomolar-to-femtomolar limits of detection via a combination of ^{129}Xe hyperpolarization and chemical exchange saturation transfer (CEST) techniques. This review aims to summarize the development of cryptophane biosensors for ^{129}Xe MRI applications, while highlighting innovative biosensor designs and the consequent enhancements in detection sensitivity, which will be invaluable in expanding the scope of ^{129}Xe MRI.

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1. Introduction

Proton magnetic resonance imaging (^1H MRI) is a commonly used imaging modality for the detection of disease pathology. Its high spatiotemporal resolution, penetration depth, and use of non-ionizing radiation make it an attractive technique for obtaining key anatomical and physiological information. However, due to the large background signal from water and fat molecules, it has limited applications in the identification of low-abundance molecular signatures of disease, which is often paramount in early disease detection. To address this sensitivity issue, MRI with heteronuclei lacking background signal has been investigated as a complement to standard ^1H MRI techniques. Here, we will focus on the spin-1/2 noble gas isotope ^{129}Xe , which has good solubility in aqueous and organic media,¹ a relatively high natural abundance of 26%,² low toxicity, and a large, polarizable electron cloud. Its high polarizability results in a ^{129}Xe NMR chemical shift window of over 300 ppm,^{3,4} which facilitates well-resolved detection in different chemical environments. This makes ^{129}Xe an attractive candidate for biomolecular imaging applications, including multiplexed detection. Additionally, ^{129}Xe can be readily hyperpolarized (hp) via spin-exchange optical pumping,⁵ whereby electron spin polarization from Rb or Cs atoms is transferred to the nuclear spin of ^{129}Xe . By this process, the signal-to-noise ratio of the

MRI measurement can be increased by several orders of magnitude over the normal Boltzmann distribution.^{6,7} Additionally, the T_1 of hp ^{129}Xe is sufficiently long (99 h at 14.1 T⁸ in the gas phase and 66 s when dissolved in saline at 9.4 T⁹) such that high polarization levels can be maintained during transport. With respect to clinically relevant conditions, the T_1 values of hp ^{129}Xe at 1.5 T in the lungs¹⁰ and oxygenated blood¹¹ are 20 and 8 s, respectively, which are acceptable for *in vivo* applications.

The compatibility of ^{129}Xe with *in vivo* imaging was first demonstrated by Albert and colleagues, who obtained a high-resolution image of mouse lungs after inhalation of hp ^{129}Xe .¹² This pioneering experiment has resulted in the development of hp ^{129}Xe MRI techniques for the imaging of human lungs^{10,13–17} and brain,^{18–20} of which there are now numerous examples. Efforts in ^{129}Xe MRI have also focused on using Xe host-guest chemistry, where the Xe guest interacts with a protein or small-molecule host, in biosensing applications. Genetically encoded Xe biosensors such as gas vesicles²¹ and monomeric proteins^{22–26} possessing hydrophobic cavities that allow for transient Xe interactions are being developed as MRI analogues of commonly used optical reporters such as green fluorescent protein. Additionally, small-molecule Xe hosts, which can be appended to an affinity tag for a biomarker of interest, can function as Xe biosensors via the modulation of the chemical shift or intensity of the ^{129}Xe signal by interactions between the Xe host and the target. For example, encapsulated Xe can generate a unique “bound” ^{129}Xe NMR chemical shift when the host binds to its target, well-resolved from the ^{129}Xe

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signal of unbound biosensor, as well as from free ^{129}Xe in solution. The diverse family of Xe biosensors has been thoroughly analyzed in recent reviews by our laboratory.^{27,28}

The most investigated Xe biosensors involve organic cage-like host molecules known as cryptophanes. As seen in Fig. 1, these cages provide a unique chemical environment for Xe, and highlight the considerable sensitivity of the ^{129}Xe NMR chemical shift to its environment: the chemical shifts of ^{129}Xe bound inside compounds 1–4 span a range of ca. 40 ppm with only an 8% change in cavity volume,^{29,30} while that of compound 5 is shifted to 308 ppm due to its six electron-withdrawing [$\eta^5\text{-C}_5\text{Me}_5$] Ru^{II} moieties.⁴ The prototypal cryptophane cage, cryptophane-A, was first synthesized in 1981, *via* the linkage of two cyclotrimeratrylene (CTV) caps by three ethylene linkers.³¹ Over the past four decades, considerable progress has been made in the synthesis of novel cryptophane derivatives, characterization of their structural aspects, and exploration of their unique host–guest chemistry, in both organic and aqueous solutions. We refer the reader to excellent in-depth reviews of these subjects.^{32,33} This review focuses on the biosensing features of Xe–cryptophane interactions, with applications in ^{129}Xe MRI. We seek to familiarize the reader with the fundamentals of cryptophane-based biosensing utilized in ^{129}Xe magnetic resonance spectroscopy (MRS), and highlight recent advances in biomarker detection and imaging. We conclude with several cutting-edge methods to advance the frontiers of Xe–cryptophane host–guest chemistry and offer our perspectives on future research.

2. Cryptophanes as biosensors in ^{129}Xe MRS

2.1 Direct detection of cryptophane-encapsulated ^{129}Xe

The Xe-binding properties of cryptophane-A were first discovered in 1998, with ^1H NMR experiments revealing good Xe affinity ($K_a \approx 3000 \text{ M}^{-1}$ in $\text{C}_2\text{D}_2\text{Cl}_4$ at 278 K), and well-resolved ^{129}Xe NMR signals from both solvated ^{129}Xe and cryptophane-bound ^{129}Xe .³⁴ In 2001, Pines, Schultz, and colleagues exploited these features of cryptophane-A in designing the first cryptophane-based biosensor.³⁵ After modifying the cryptophane-A molecule with a solubilizing peptide and a tether linking the cage to a biotin moiety, a ^{129}Xe NMR peak corresponding to bound ^{129}Xe was

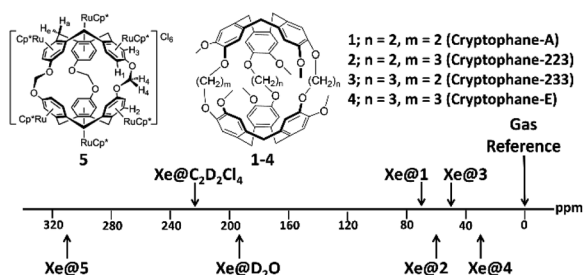


Fig. 1 Chemical structure and ^{129}Xe NMR chemical shifts of cryptophanes with varying alkoxy linker length in 1,1,2,2-tetrachloroethane- d_2 (1–4)³⁰ and of a cryptophane functionalized with six [$\eta^5\text{-C}_5\text{Me}_5$] Ru^{II} moieties (5)⁴ in D_2O .

observed at ca. 70.2 ppm, referenced to gaseous ^{129}Xe . Addition of 80 nmol avidin shifted this peak 2.3 ppm downfield, which was attributed to mechanical deformation of cryptophane and the corresponding distortion of xenon's highly polarizable electron cloud due to the biotin–avidin interaction (Fig. 2).

This innovative experiment demonstrated the feasibility of using functionalized cryptophanes and ^{129}Xe NMR to detect analytes with high sensitivity and has prompted the use of this technique in the detection of numerous biomarkers of interest. Generally, this methodology employs the use of a targeting moiety which can bind to or otherwise interact with the biomarker, resulting in a “bound” ^{129}Xe NMR signal that is distinct from “free” biosensor. Appending the targeting moiety to the cryptophane core can be accomplished *via* copper-catalyzed [3 + 2] azide–alkyne cycloaddition, whereby an affinity tag with an azido group can be appended onto a cryptophane-A derivative with propargyl moieties.³⁶ The connection of a targeting moiety with an amine group to a carboxylated cryptophane *via* an amide linkage is also commonly utilized. For hydrophobic cryptophane scaffolds, one or more solubilizing groups possessing such ionizable moieties as carboxylic acids^{37–39} or amines⁴⁰ can also be attached in order to increase the water solubility of the cage. Using this approach, the detection of such biomedically relevant targets as integrin receptor,⁴¹ carbonic anhydrase,^{42–44} matrix metalloproteinase-7 (MMP-7),⁴⁵ and major histocompatibility complex class II protein⁴⁶ was feasible with cryptophanes functionalized with affinity tags cyclic RGDyK peptide, benzenesulfonamide, MMP-7 specific peptide substrate, and hemagglutinin peptide, respectively.

In addition, the detection of metal ions *via* functionalized cryptophanes has become a valuable tool to supplement

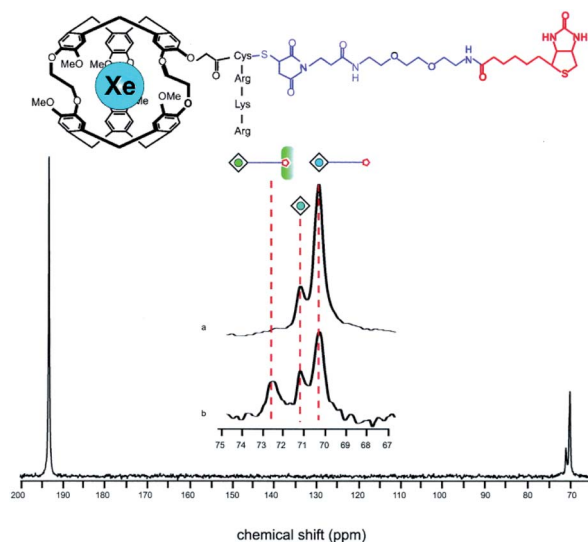


Fig. 2 Top: Chemical structure of a biosensor comprised of cryptophane-A, a solubilizing peptide and a tether linking the construct to biotin. Bottom: ^{129}Xe NMR spectrum showing the $^{129}\text{Xe}(\text{aq})$ signal at ca. 193 ppm and the bound ^{129}Xe signals at ca. 70 ppm. The peak corresponding to ^{129}Xe encapsulated by this biosensor shifts from ca. 70.2 to 72.5 ppm upon addition of avidin to solution. Reproduced with permission from ref. 35. Copyright 2001 National Academy of Sciences.

traditional analytical methods. In 2012, Kotera *et al.* demonstrated the first example of cryptophane-based metal ion detection by appending the zinc-chelating group nitrilotriacetic acid to cryptophane for the detection of Zn^{2+} .⁴⁷ A follow up study showed that this construct could also be used to detect Cd^{2+} and Pb^{2+} , with ^{129}Xe NMR chemical shifts migrating 1.5, 0.3, and 4.5 ppm downfield from free biosensor for Zn^{2+} , Cd^{2+} , and Pb^{2+} , respectively.⁴⁸ In recent years, efforts have focused on the design of cryptophane-based metal ion sensors with high selectivity and sensitivity. Zhang *et al.* designed a highly selective cryptophane functionalized with 2-(diphenylphosphino)benzenamine, which exhibited a sizable 6.4 ppm upfield ^{129}Xe NMR chemical shift upon Zn^{2+} binding, while being unresponsive to Ag^+ , Cu^+ , Pd^{2+} , Hg^{2+} , Mg^{2+} , and Ca^{2+} .⁴⁹ Additionally, Jeong *et al.* demonstrated how a cryptophane functionalized with the chelator 1,4,7,10-tetraazacyclododecane-1,4,7,10-tetraacetic acid was able to differentiate between Ca^{2+} , Cu^{2+} , Ce^{3+} , Zn^{2+} , Cd^{2+} , Ni^{2+} , Co^{2+} , Cr^{2+} , Fe^{3+} , and Hg^{2+} by exhibiting distinct ^{129}Xe NMR chemical shifts upon metal ion binding.⁵⁰

Cryptophane-based biosensors have also been designed to report on more complex systems. For example, Roy *et al.* observed a 1.5 ppm upfield ^{129}Xe NMR chemical shift change upon binding of a cryptophane functionalized with a 20-mer nucleotide to its complementary DNA strand, and no such change in the presence of a non-complementary strand.⁵¹ Berthault *et al.* demonstrated how the NMR chemical shift of ^{129}Xe encapsulated by a hexa-carboxylate-functionalized cryptophane is highly pH dependent,⁵² and later used the chemical shift difference of encapsulated ^{129}Xe in two different cryptophanes for more precise pH measurement.⁵³ Finally, Yang *et al.* designed a cryptophane functionalized with a thiol-reactive acrylate moiety, which allowed for the detection of, and differentiation between biothiols such as cysteine, homocysteine, and glutathione.⁵⁴ These innovative examples of cryptophane-based biosensor design highlight where recognition moieties can be tailored for highly specific and selective binding. Conversely, using a more universal approach to ^{129}Xe NMR biosensing, Kotera *et al.* demonstrated how Cys4-tagged proteins can be detected using a cryptophane linked to a CrAsH probe (a carboxy derivative of the bisarsenical FLAsH probe).^{55,56} This method allows for the detection of wide families of proteins that contain the small tetracysteine tag, which is commonly used for analyzing protein function and location.

Despite this recent progress, there is still much optimization necessary in the design of cryptophane-based biosensors for direct detection ^{129}Xe NMR. For example, there have been few examples of chemical shift changes >5 ppm upon biosensor-target binding, which can pose resolution issues at clinical magnetic field strengths. Additionally, due to chirality of the cryptophane (typically synthesized as a racemic mixture), and the chirality of many affinity tags, multiple peaks corresponding to different diastereomers can be observed in the ^{129}Xe NMR spectrum.⁵⁷ This phenomenon dilutes the signal-to-noise by the number of diastereomers, although this can be mitigated through enantiopure cryptophane synthesis.^{36,44,58} Xe access to the cryptophane interior can be limited by the energy necessary to displace confined water molecules inside the cage,⁵⁹ and precluded entirely

by collapsed conformations of the cryptophane cage in which one CTV unit folds inward into the cavity.⁶⁰ Spectral analysis can be further complicated by the presence of residual peaks corresponding to unbound biosensor. The use of 'turn-on' biosensors, where ^{129}Xe NMR signal is observed only in the presence of target, can simplify these analyses. Our laboratory recently reported the first turn-on cryptophane biosensor, which consisted of a FRRIAR peptide conjugated to cryptophane for the detection of calmodulin (CaM).⁶¹ Signal corresponding to bound ^{129}Xe was only observed in the presence of Ca^{2+} -bound CaM.

Additionally, we showed that cryptophane-based biosensors have a propensity to form water-soluble aggregates in solution,⁶² which can further complicate chemical shift assignments. Using a benzenesulfonamide-functionalized cryptophane (C8B) that was previously used⁴² for the detection of carbonic anhydrase (CA) as a model system, we observed that the biosensor formed aggregates that were hundreds of nanometers in diameter, but became monomeric upon binding to its protein target.⁶² We showed that as increasing amounts of CA were titrated into solutions of C8B, the chemical shift of encapsulated ^{129}Xe progressively shifted downfield, which we attributed to mechanical perturbation of the encapsulated ^{129}Xe electron cloud,^{35,57,63,64} and also the disaggregation phenomenon. Biosensor disaggregation can contribute significantly to the ^{129}Xe NMR chemical shift change observed upon target binding. Indeed, we hypothesize that only biosensor-target interactions that cause a significant change in the cryptophane aggregation state will produce substantial chemical shift changes. This hypothesis is supported by recent studies from our laboratory with an adamantyl-functionalized cryptophane-A sensor, which remains monomeric in aqueous solution, and does not exhibit a ^{129}Xe NMR chemical shift change upon binding the model receptor β -cyclodextrin.⁶⁵ The modulation of the ^{129}Xe NMR chemical shift as a function of such factors as stereochemistry and aggregation is a relatively unexplored area of cryptophane-based biosensing that is sure to receive additional attention as biosensing techniques are optimized.

2.2 Increasing ^{129}Xe NMR detection sensitivity with hyper-CEST

Hyperpolarization of ^{129}Xe offers a significant improvement in signal-to-noise relative to thermal polarization, but direct detection of encapsulated ^{129}Xe NMR signal nevertheless requires micromolar concentrations of biosensor to avoid long acquisition times.⁶⁶ Previous work showed that peptide-functionalized cryptophanes are relatively nontoxic to cells at concentrations of <100 μM ,⁶⁷ yet lower working concentrations of biosensor are still desired to maximize cell viability and to detect low-abundance molecular markers of disease. To increase detection sensitivity further, the hyper-CEST technique was developed, which combines ^{129}Xe hyperpolarization with chemical exchange saturation transfer (CEST).⁶⁸ Hyper-CEST takes advantage of the constant exchange of Xe into and out of the cryptophane, which typically occurs on the order of tens of Xe atoms per second in aqueous solution at rt.^{30,35} The encapsulated hp ^{129}Xe can be selectively depolarized by



applying radiofrequency (rf) pulses at the resonance frequency of the bound ^{129}Xe atom. The depolarized ^{129}Xe then exchanges into solution, causing depolarization of solvated ^{129}Xe nuclei (and rapid thermal equilibration). The subsequent loss of $^{129}\text{Xe}(\text{aq})$ signal acts as a reporter for the encapsulated ^{129}Xe pool, allowing for ultrasensitive detection of the cryptophane host (Fig. 3).

In order to approximate the contrast that one would observe in a ^{129}Xe MRI experiment, a time-dependent saturation profile of the Xe host can be obtained. First demonstrated by Meldrum *et al.*,⁶⁹ this technique involves applying rf pulses at the bound ^{129}Xe resonance frequency and monitoring the decrease in $^{129}\text{Xe}(\text{aq})$ NMR signal as a function of saturation time. This on-resonance decay of the $^{129}\text{Xe}(\text{aq})$ signal can be compared to an off-resonance decay, where rf pulses are applied at a resonance frequency where no hyper-CEST effect is observed, and the difference between the two decay curves, which is weighted by the saturation time, can be quantified as saturation contrast.⁷⁰

Several recent examples highlight the increase in sensitivity afforded by the hyper-CEST technique. For instance, a peptide-functionalized cryptophane biosensor developed by Riggle *et al.* was utilized to label cells in acidic environments.⁷¹ A cryptophane-conjugated EALA-repeat peptide assumed a disordered form at neutral pH but became α -helical at pH 5.5, which resulted in insertion into HeLa cell membranes and a subsequent 13 ppm downfield chemical shift change of the encapsulated ^{129}Xe (Fig. 4). Importantly, significant hyper-CEST contrast was observed with 34 pM biosensor, which improved detection sensitivity by 6 orders of magnitude relative to direct detection of the “bound” ^{129}Xe NMR signal. In addition to

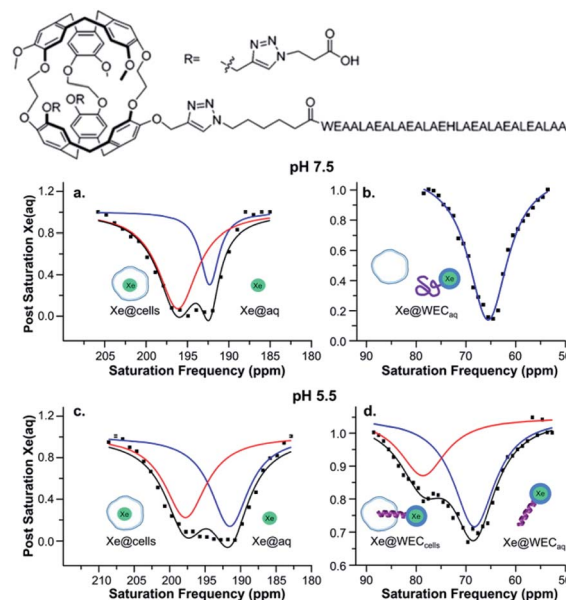


Fig. 4 Top: Chemical structure of water-soluble EALA-cryptophane (WEC). Bottom: Hyper-CEST ^{129}Xe NMR of 5–10 μM WEC with 1×10^7 HeLa cells per mL at pH 7.5 (a and b) and pH 5.5 (c and d). Reproduced with permission from ref. 71. Copyright 2015 American Chemical Society.

allowing for the highly sensitive detection of specific Xe host molecules, hyper-CEST can also facilitate the differentiation between biosensor in the absence and presence of target. Zeng *et al.* designed a mitochondria-targeted biosensor for the detection of biothiols, comprising cryptophane-A, a disulfide linker, a naphthalimide fluorescent moiety, and a triphenylphosphonium mitochondrial targeting tag.⁷² Hyper-CEST revealed a 200 pM detection limit for this biosensor, but also allowed for the monitoring of intracellular biothiol levels. It was observed that the CEST effect for this biosensor incubated with lung cancer cells was significantly larger than for biosensor incubated with lung cancer cells and *N*-ethylmaleimide, a thiol scavenger. This was due to the biosensor becoming more water-soluble and accessible to Xe after thiol-induced cleavage of the disulfide bond, which highlights the importance of understanding solution properties in biosensor design. Finally, Milanole *et al.* used hyper-CEST to show that their cryptophane-based biosensor, functionalized with a fluorescein moiety and an antibody targeted toward epidermal growth factor receptor (EGFR), was selective toward cells overexpressing EGFR.⁷³ Saturation contrast was only observed with biosensor incubated with the HCC827 cell line (high EGFR expression) and not with biosensor incubated with the A549 cell line (intermediate EGFR expression), with intracellular biosensor concentration in the low-nanomolar regime.

3. From spectroscopy to imaging

3.1 *In vitro* ^{129}Xe MRI studies

Heretofore, we have discussed the contributions made to cryptophane-based biosensing *via* spectroscopic studies. These

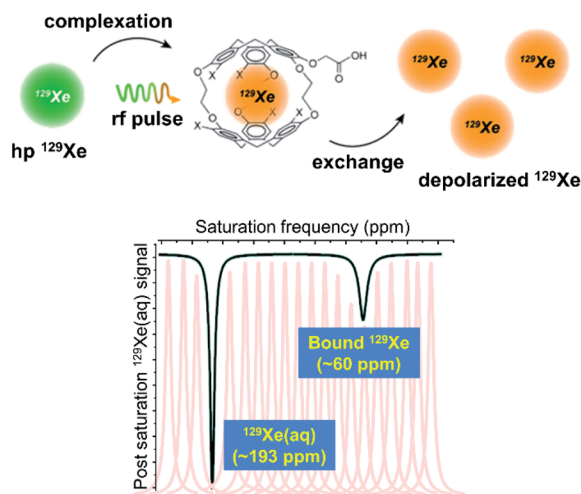


Fig. 3 Top: Mechanism of the hyper-CEST NMR experiment. Hp ^{129}Xe atoms (green) bind to the interior of the cryptophane cage, and are depolarized (orange) by a selective rf pulse. The exchange of bound ^{129}Xe out of the cage causes the depolarization of bulk ^{129}Xe nuclei. This scheme was modified with permission from ref. 66. Copyright 2014 Wiley-VCH. Bottom: Representative ^{129}Xe hyper-CEST NMR z-spectrum of a cryptophane host. The individual peaks represent the intensity of the ^{129}Xe signal as a function of the frequency at which rf pulses are applied and the black trace represents the collective z-spectrum.



experiments and analyses have not only advanced the growth of this field, but have also stimulated progression into the clinical realm, where obtaining high-contrast images of target is made feasible by designing cryptophane biosensors with optimal hyper-CEST properties. For more details on technical aspects related to the development of ^{129}Xe MRI techniques, we refer the reader to prior reviews of this subject.^{74–76} We note that several of the biosensors discussed previously, as well as several of those that will be discussed in this section, are classified as “bimodal” contrast agents, containing both cryptophane and a fluorophore, and thus detectable *via* both ^{129}Xe MRI and fluorescence imaging.^{55,56,73,77–82} The capacity for analysis *via* a secondary detection method such as fluorescence can be greatly beneficial in complementing the ^{129}Xe MRI readout. Additionally, it has been shown that differences in membrane fluidity can affect the depolarization time of hp ^{129}Xe . Studies by Schnurr *et al.* showed that high membrane fluidity resulted in short depolarization times and saturated hyper-CEST responses with short saturation pulses.⁸³ Additional work by Booker and Sum provided valuable insight with respect to Xe–membrane interaction, showing that Xe localizes in the hydrophobic core of membranes, and by doing so increases bilayer thickness, membrane fluidity, and lipid head group spacing.⁸⁴ This interaction of Xe with membranes has also been exploited in the design of liposome bilayers used for the delivery of the gas in rat models of stroke.^{85–87} Thus, having a secondary imaging modality unaffected by the dynamic interactions of Xe with the cellular environment becomes valuable when performing experiments.

Kotera *et al.* first demonstrated the enhanced sensitivity of ^{129}Xe MRI with their nitrilotriacetic acid-functionalized cryptophane for the detection of Zn^{2+} ions.⁴⁷ They observed a significant increase in MRI contrast when Ca^{2+} ions were replaced with Zn^{2+} in a solution of biosensor, and showed that the limit of Zn^{2+} detection was as low as 100 nM, which was at the time an improvement of two orders of magnitude over conventional MRI methods. Efforts from the Zhou laboratory and collaborators have also focused on the development of ^{129}Xe MRI contrast agents. For example, they designed a unique biosensor for Hg^{2+} detection comprised of two cryptophane-A cages linked by a dipyrrolylquinoxaline moiety.⁸⁸ Hg^{2+} binding resulted in a clamp-like conformational change of the biosensor, in which the two cryptophanes converge and their electron clouds overlap, causing a ^{129}Xe NMR chemical shift change. This change was enough to observe ^{129}Xe MRI contrast in the presence of Hg^{2+} , with no contrast observed with biosensor alone. They also demonstrated the use of ^{129}Xe MRI in the detection of biothiols using the previously mentioned acrylate-functionalized cryptophane,⁵⁴ where contrast was observed only when the biosensor was incubated with cysteine; and in the detection of H_2S .⁸² This latter study featured a biosensor comprised of cryptophane-A functionalized with 4-azide-1,8-naphthalic anhydride, which was able to detect HS^- due to the ^{129}Xe NMR chemical shift change that occurred after the HS^- -induced reduction of the azido group to an amine (Fig. 5). ^{129}Xe MRI contrast was observed only after the biosensor was incubated with HS^- . It should be noted that the design of this biosensor featured

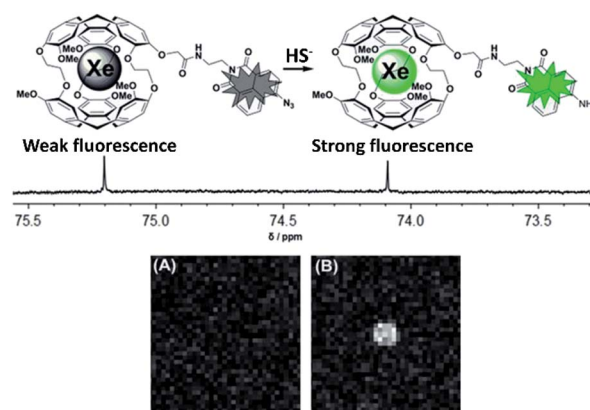


Fig. 5 Top: Scheme of biosensor for H_2S , activatable *via* both fluorescence and ^{129}Xe MRI modalities. Bottom: ^{129}Xe MRI images of an NMR tube containing 100 μM biosensor (A) and 100 μM biosensor with 10 equiv. HS^- (B). Reproduced with permission from ref. 82. Copyright 2017 Wiley-VCH.

a fluorescent tag that could simultaneously function as an analyte probe, as the HS^- -induced reduction of the azido group also caused an increase in the fluorescence intensity as well as a change in the ^{129}Xe NMR chemical shift. Integration of two detection modalities into one sensor should simplify synthetic efforts and facilitate biocompatibility in future designs.

In addition to being used in the detection of biomedically relevant analytes, ^{129}Xe MRI with cryptophanes has also found use in thermometry applications. Schilling *et al.* showed that the chemical shift of ^{129}Xe encapsulated by cryptophane-A has a temperature dependence of $0.29 \text{ ppm } ^\circ\text{C}^{-1}$, which is *ca.* 30-fold greater than water.⁸⁹ This temperature dependence arises from the large electron cloud of Xe, which is able to interact with less shielded areas of the cryptophane interior, as well as from the greater conformational flexibility of the cryptophane at higher temperatures. Additionally, using chemical shift imaging with a cryptophane concentration of 150 μM , they were able to determine that the lower limit of the detectable temperature change was $0.1 ^\circ\text{C}$, which suggests that highly sensitive and accurate temperature mapping is feasible *via* ^{129}Xe hyper-CEST MRI.

3.2 *In cellulo* ^{129}Xe MRI studies

Even though considerable efforts are still needed to successfully bridge the gap to clinical *in vivo* studies, there have been numerous pioneering experiments that illustrate the potential of cryptophanes for *in vivo* ^{129}Xe MRI applications. In particular, the Schröder laboratory and their collaborators have made valuable contributions to developing ^{129}Xe MRI with cryptophane-based biosensors by performing numerous *in cellulo* studies. In 2012, they demonstrated that using cryptophane-A in imaging applications requires only *ca.* 100 μM ^{129}Xe and only 30 nM of NMR-active cryptophane with a CEST saturation pulse of 26 s, which can be reduced down to sub-second timescales if *ca.* 10 μM NMR-active cage is used.⁹⁰ Their group was also the first to use ^{129}Xe MRI with cryptophane



to image live cells,⁸⁰ which revealed valuable information about the *in cellulo* distribution of Xe. Using mouse fibroblasts, they observed the expected pools of Xe in bulk solution (192 ppm) and encapsulated by cryptophane-A (59 ppm), but also solvated Xe in cells (196 ppm) and Xe encapsulated by cell-internalized cryptophane (69 ppm). Due to the sizable chemical shift difference between Xe encapsulated in free and cell-internalized cryptophane, as well as the sensitivity of hyper-CEST, it was possible to perform selective ¹²⁹Xe MRI of this latter pool. By using a construct comprised of cryptophane-A functionalized with fluorescein, they were able to use ¹²⁹Xe MRI to clearly differentiate between labeled and non-labeled cells (Fig. 6), with an average intracellular cryptophane concentration of 15 μM, achieved after incubating cells with 50 μM cryptophane for 20 h. In a follow-up study, they showed that cryptophane-A functionalized with a different fluorophore, tetramethylrhodamine, was able to achieve similar ¹²⁹Xe MRI contrast to that observed with the former construct.⁹¹ This illustrated the flexibility of cryptophane-based ¹²⁹Xe imaging, and suggested the possibility of fluorescence multiplexing applications.

After having succeeded in obtaining ¹²⁹Xe MRI images of cells with nonspecific cryptophane uptake, they demonstrated the feasibility of performing cell-specific targeting using a functionalized cryptophane. In a key proof of principle study, they targeted cells expressing the cell surface protein and macrophage marker cluster of differentiation 14 (CD14).⁷⁸ By functionalizing cryptophane-A and fluorescein with biotin, conjugating an anti-CD14 antibody to avidin, and incubating these constructs with high-CD14-expressing macrophages and a control fibroblast cell line, they were able to distinguish between the two types of cells using ¹²⁹Xe MRI and fluorescence with only *ca.* 20 nM biosensor. Notably, these experiments demonstrated the versatility of their modular approach, as the biosensor can be adapted to detect any target for which there is

a specific antibody or other binding element available. For example, they later showed how this general targeting scheme could be used for the detection of members of the claudin (Cldn) protein family.⁹² In this study, they conjugated *Clostridium perfringens* enterotoxin, which has high affinity for specific claudins, to avidin, and used the previously mentioned cryptophane-biotin and fluorescein-biotin constructs as ¹²⁹Xe MRI and fluorescence readout modules, respectively. This biosensor was successfully used to image and distinguish Cldn4-expressing HEK cells (59% CEST effect) from control non-transfected HEK cells (11% CEST effect) *via* ¹²⁹Xe MRI.

In their next study, they demonstrated a technique for delivering cryptophane to human brain microvascular endothelial cells (HBMECs) by using large unilamellar vesicles (LUVs).⁹³ In their experimental design, they functionalized the lipid bilayer of cryptophane-containing LUVs (*ca.* 3800 cryptophane units per liposome) with arginine-rich peptides to facilitate uptake into HBMECs. The target cells and control human aortic endothelial cells (HAoECs) were then incubated with both functionalized and unfunctionalized LUVs, and cellular uptake was studied *via* fluorescence and ¹²⁹Xe MRI modalities. Indeed, uptake of the biosensor into HBMECs resulted in an MRI CEST effect *ca.* 3-fold larger than that observed with HAoECs, allowing for the two cell types to be spatially distinguished. Because of the high loading capacity of LUVs, these images were able to be obtained with only 1 nM of biosensor, illustrating the high sensitivity of this approach. This technique was later improved by using a lipopeptide comprised of a cationic peptide, a cryptophane-A unit, and a palmitoyl chain.⁹⁴ After micelle formation, it was estimated that the average cryptophane content per unit volume was *ca.* 7-fold higher in the micelles than in LUVs. This allowed for a significantly higher CEST effect to be observed with HBMECs relative to HAoECs, with the important difference being that the cells were incubated with a cryptophane concentration *ca.* 10-fold more dilute than that used in the LUV study. This was facilitated by the high local cryptophane concentration in this biosensor design, resulting in a high density of depolarization seed points. Finally, an additional study focused on targeting cell-surface glycans by functionalizing cryptophane-A with a fluorophore and bicyclo[6.1.0]nonyne for glycan labeling.⁷⁹ Cells treated with the synthetic sugar Ac₄ManNAz for the purpose of expressing an azido-functionalized sialic acid on their surface were able to be clearly imaged and differentiated from untreated cells using this biosensor. Notably, this study involved covalently linking the biosensor to its target *via* azide-alkyne cycloaddition, thus eliminating signal from unbound biosensor after washout.

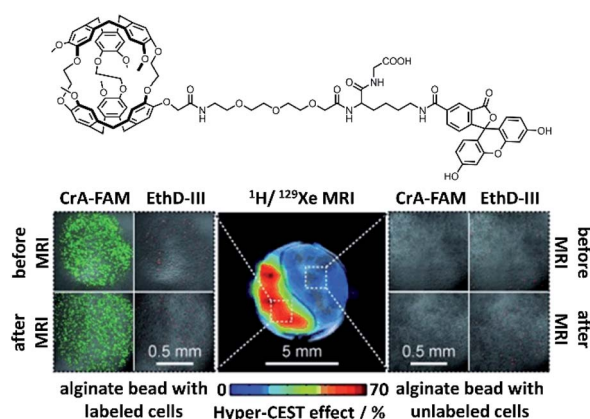


Fig. 6 Top: Chemical structure of fluorescein-conjugated cryptophane (CrA-FAM). Bottom: ¹²⁹Xe hyper-CEST MRI with underlying ¹H MRI images of mouse fibroblasts labeled with CrA-FAM (15 μM intracellular concentration) and control cells, encapsulated in alginate beads. Laser scanning microscopy images of each cell population are shown, with CrA-FAM accumulated in cells in green and dead cells stained with ethidium homodimer III (EthD-III) in red. Reproduced with permission from ref. 80. Copyright 2014 Wiley-VCH.

4. New frontiers in cryptophane-based biosensing

Over the past decade, there has been considerable progress in the creative utilization of cryptophanes beyond their traditional roles as single-unit hosts for ¹²⁹Xe biosensing. In this section, we highlight several inventive examples that push the boundaries of cryptophanes for ¹²⁹Xe MRI.



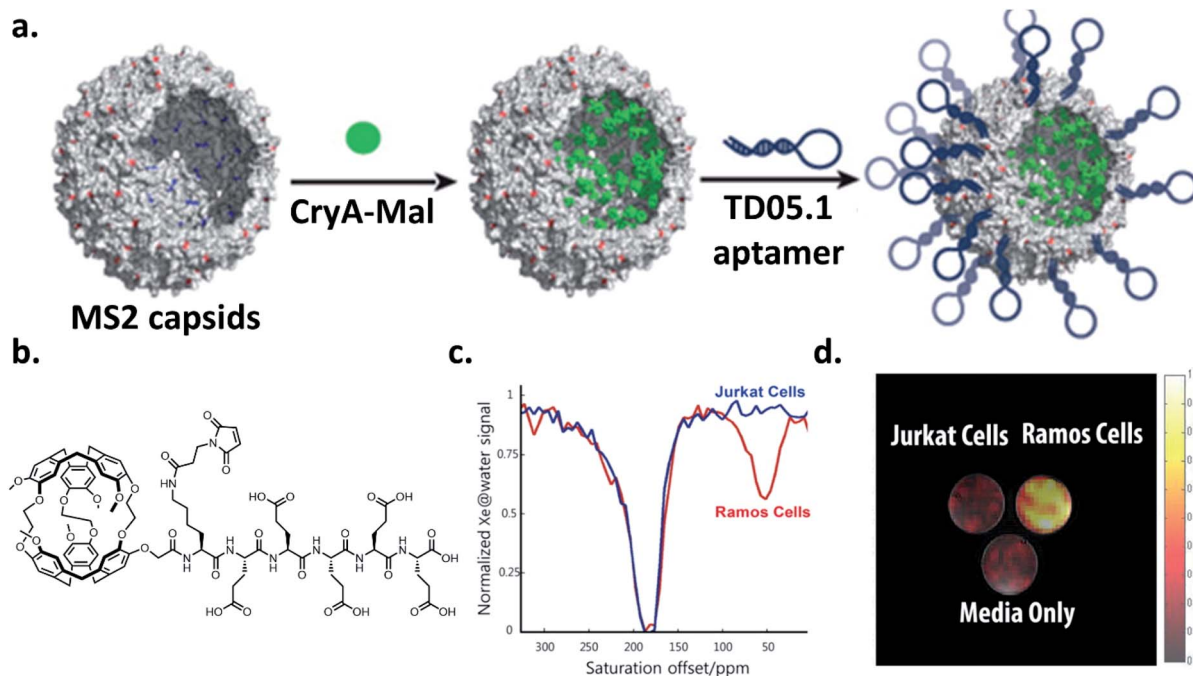


Fig. 7 (a) Scheme of bacteriophage–cryptophane biosensor assembly. The capsid interior of bacteriophage MS2 was modified with cryptophane-A conjugated to maleimide and five glutamic acids to increase solubility (CryA-Mal, green circle) at ca. 110 N87C positions. The capsid exterior was modified with 54 units of the TD05.1 DNA aptamer for targeting mIgM markers on lymphoma cells. (b) Chemical structure of CryA-Mal. (c) ^{129}Xe hyper-CEST NMR z-spectra of the biosensor (167 nM in capsids) incubated with 2×10^7 Ramos (mIgM $^+$) or Jurkat (mIgM $^-$) cells. (d) ^{129}Xe hyper-CEST MRI superimposed on ^1H MR image of tubes containing cell media only, and the two cell lines incubated with biosensor. Reproduced with permission from ref. 98. Copyright 2016 American Chemical Society.

One of the critical goals in this area has been to enhance the detection sensitivity of cryptophane hosts. In order to achieve this, significant efforts have focused on grafting multiple cryptophane units onto a larger scaffold, notably by the laboratories of Pines, Wemmer, and collaborators. In 2006, they demonstrated the first example of this strategy by attaching a biotin moiety to a polyamidoamine dendrimer, which was able to encapsulate two cryptophane-A units when bound to avidin, and increased signal-to-noise by a factor of 8 relative to their original biotinylated cryptophane biosensor.^{35,95} They improved upon this concept four years later by using bacteriophage MS2 as a scaffold for attaching multiple cryptophane-A units *via* linker molecules.⁶⁹ This allowed for the attachment of 125 cryptophane-A units per viral capsid, which resulted in a 700 fM detection limit for the cryptophane-capsid construct. Further improvements in detection sensitivity were achieved with the bacteriophage M13, which possesses 15 times as many coat proteins on its capsid as MS2, allowing for 1050 cryptophane-A host molecules per capsid to be attached, and a detection sensitivity of 230 fM.⁹⁶ Biosensing capability with a construct comprising aminoxy-functionalized cryptophane-A ligated to filamentous bacteriophage fd expressing single-chain antibody variable fragments (scFvs) was then established.⁹⁷ Significant hyper-CEST contrast in the presence of MDA-MB-231 cells expressing high levels of EGFR, recognized by the scFvs, and minimal contrast in the presence of Jurkat cells expressing low levels of EGFR was observed. Importantly, this demonstrated that constructs possessing multiple cryptophane units can

avoid the nonspecific cell uptake previously observed with singular cryptophane-A.⁸⁰ Finally, in the most recent demonstration of specific cell targeting, it was established that it was possible to use a cryptophane-bacteriophage construct modified on its exterior with DNA aptamers to target mIgM markers on the surfaces of lymphoma cells (Ramos line) (Fig. 7).⁹⁸ Taken together, these experiments illustrate the potential of greatly enhancing cryptophane detection sensitivity by grafting individual units onto a larger scaffold. We note that it should also be possible to attach other recognition elements, such as antibodies or peptides, onto the viral capsids for the detection of a wide range of targets.

The prospect of using multiplexed cryptophane-based biosensors with distinct chemical shifts to detect different molecular targets is also a burgeoning area of research. Berthault *et al.* first demonstrated the possibility of multiplexed cryptophane imaging by performing ^{129}Xe MRI with an NMR tube containing an aqueous solution of a carboxylated cryptophane layered on top of a solution of an organosoluble cryptophane in tetrachloroethane.⁹⁹ To facilitate cryptophane functionality, Tyagi *et al.* showed that cryptophane-A can be functionalized with polyglycerol dendrons and still preserve its optimal Xe binding affinity and exchange rate.¹⁰⁰ The multitude of hydroxyl groups at the outer surface of the cage allowed for high water solubility as well as for the potential of further functionalization with multiple targeting groups with high regioselectivity.¹⁰¹ Live cell multichannel ^{129}Xe MRI was demonstrated by Klippel *et al.*, who used cryptophane-A as well



as perfluorooctyl bromide (PFOB) nanodroplets as Xe hosts.¹⁰² Because the ^{129}Xe NMR chemical shift of cell-internalized cryptophane-A is approximately 40 ppm upfield of cell-internalized PFOB, it was possible to image mouse fibroblast cells *via* a frequency-selective saturation transfer approach, where observed contrast for the two Xe hosts was dependent upon the saturation frequency. Moreover, due to the ^{129}Xe NMR chemical shift of cell-internalized cryptophane being nearly independent of cell type,^{80,91,93} these experiments illustrate the possibility of imaging varying cell types using this multichannel detection approach. Importantly, the use of cryptophanes in tandem with established contrast media such as PFOB nanodroplets is a valuable example of how the unique chemical environment, high detection sensitivity, and multifunctionality inherent to cryptophane contrast agents can drive their integration into modern biomedical applications.

5. Concluding remarks

The field of cryptophane biosensing has greatly evolved since the conception of the first xenon biosensor two decades ago. We have reviewed examples of the ultrasensitive detection and monitoring of proteins,^{35,41,42,45,46,55,56,61,73,78,79,92,97} ions,^{47–50,82,88} nucleic acids,⁵¹ biothiols,^{54,72} and even environmental conditions such as pH^{52,53,71} and temperature⁸⁹ using direct and indirect detection methods. We have highlighted instances of innovative biosensor design, such as the use of a singular moiety for both fluorescence and ^{129}Xe MRI readout,⁸² modulation of solubility after target binding,⁷² and loading up to thousands of individual cryptophane host molecules onto a single carrier.^{69,93–98} Finally, we have shown examples of selective and sensitive ^{129}Xe MRI contrast attainable with these biosensors. This past decade of expanding the scope of ^{129}Xe MRI using cryptophane-based biosensing has been extremely productive, and the future of this field remains bright.

Considerable progress is still necessary for cryptophane–xenon biosensors and ^{129}Xe MRI to achieve widespread use as imaging tools. Synthetic challenges including low overall yields from multistep syntheses and difficulties in performing large scale cryptophane syntheses due to the propensity of polymer formation at high concentrations are notable limiting factors.³² Also, translation of the end products into bioimaging applications can be challenging due to poor biosensor solubility and propensity for self-aggregation. We note that these limitations can be overcome with other small-molecule Xe hosts, such as members of the cucurbit[*n*]uril (CB[*n*]) family, which are commercially available and relatively inexpensive macrocycles with excellent Xe exchange kinetics¹⁰³ and versatile host–guest chemistry.¹⁰⁴ Indeed, detection of cucurbit[6]uril *via* ^{129}Xe hyper-CEST MRI in the vasculature of a living rat was recently demonstrated by Albert and colleagues.¹⁰⁵ However, the targeted biosensing capabilities of CB[*n*] are limited, as these compounds are difficult to functionalize and have high affinity for naturally occurring small molecules in living organisms, such as polyamines,¹⁰³ necessitating the use of relatively high (mM) concentrations for *in vivo* imaging studies. Self-assembled metal–organic nanocapsules have recently emerged as

promising candidates for ^{129}Xe MRI applications due to their highly tunable characteristics,^{106,107} but further work is needed with respect to their functionalization in order to enable targeted imaging. Thus, at the current time, we believe that cryptophanes are the gold standard for ultrasensitive, targeted detection. The key experiment to validate their status as the definitive, biocompatible Xe host must be a demonstration of their detection *via* ^{129}Xe MRI in a living organism, and we believe that the next several years of research should view this as a critical endeavor. For example, a study examining the bio-distribution of unfunctionalized cryptophane-A in a mouse model, followed by a study with functionalized cryptophane targeting an overexpressed biomarker in transgenic mice would be essential in establishing cryptophane biocompatibility and targetability. In order to reach and progress past this milestone, efforts must be focused on optimizing syntheses of functionalized cryptophane. Recent progress in improving synthetic techniques to achieve shorter, scalable cryptophane syntheses with improved yields, milder reaction conditions, and easier purification procedures is promising,^{36,108} though efficient gram-scale syntheses are yet to be developed.

A successful demonstration of *in vivo* detection of a cryptophane-based biosensor will surely give insight on many additional ways to optimize biosensor design, as we currently have no knowledge of how cryptophanes behave in multicellular organisms with respect to biodistribution. For example, the recently discovered cryptophane aggregation phenomenon⁶² suggests that cryptophane *in vivo* circulation time and stability may be modulated by the size of the nanoscale clusters. The facile nature of cryptophane functionalization should allow for a rapid response with respect to optimizing biosensor design to enhancing biocompatibility. Indeed, there are currently many innovative ways to tune cryptophane solution properties, including appending cryptophane with dendrons carrying numerous hydroxyl groups¹⁰⁰ and functionalizing cryptophane with PEG moieties,¹⁰⁹ and we expect additional techniques to emerge in the near future.

Continued research highlighting the use of ^{129}Xe MRI and cryptophane-based biosensors for the *in vivo* detection of molecular markers of disease will facilitate the transition to the clinical realm. Based on numerous recent examples of the use of ^{129}Xe MRI in assessing physiological processes in human lungs and kidneys,^{110–113} we believe that approval for hp ^{129}Xe to be used in the clinical setting will progressively become increasingly widespread. At the time of this review, the use of hp ^{129}Xe has not yet attained universal approval for use in general clinical settings, nor have cryptophane-based biosensors been studied in multicellular organisms. However, hp ^{129}Xe is currently being used as an investigational medical product in humans at many institutions globally, and has received approval for routine clinical lung imaging in the United Kingdom.¹¹⁴ Xemed and Polarean, two prominent U.S.-based industry members in the field of hyperpolarized gas MRI, have filed requests with the United States Food and Drug Administration for the use of hp ^{129}Xe in the clinical setting. Emerging new vendors, such as XeUS Technologies, are also contributing to the growth of the field by providing affordable



instrumentation for clinical ^{129}Xe production. In summary, it is clear that once clearance by regulatory agencies is granted, the infrastructure will be readily available for a rapid transition into the clinical realm.

Additionally, modern technological advancements, such as progress in generating larger quantities of isotopically enriched ^{129}Xe and the development of ^{129}Xe polarizers capable of achieving near-unity polarization levels^{115,116} will likely allow for significantly lower working concentrations of Xe and biosensor, thereby facilitating biocompatibility and lowering cost. The integration of cryptophane-based biosensors with established imaging tools such as PFOB nanodroplets¹⁰² and LUVs,⁹³ and with common fluorescence imaging modalities, should also aid in the transition of cryptophanes into the clinical domain. Lastly, the continued development and optimization of Xe delivery systems, MRI scanners, and pulse sequences will undoubtedly expand the potential of ^{129}Xe for *in vivo* imaging applications. With these advances on the horizon, we believe that the next decade of research into cryptophane-based biosensors with applications in ^{129}Xe MRI promises to be even more rewarding than the last one.

Conflicts of interest

There are no conflicts to declare.

Acknowledgements

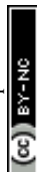
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References

- H. L. Clever, *Krypton, Xenon and Radon - Gas Solubilities*, Pergamon Press, New York, 1979.
- W. M. Haynes, *CRC Handbook of Chemistry and Physics*, 79th edn, CRC Press, Boca Raton, 2012.
- D. Rafferty, *Annu. Rep. NMR Spectrosc.*, 2006, **57**, 205–270.
- R. M. Fairchild, A. I. Joseph, K. T. Holman, H. A. Fogarty, T. Brotin, J.-P. Dutasta, C. Boutin, G. Huber and P. Berthault, *J. Am. Chem. Soc.*, 2010, **132**, 15505–15507.
- T. G. Walker and W. Happer, *Rev. Mod. Phys.*, 1997, **69**, 629.
- B. M. Goodson, *J. Magn. Reson.*, 2002, **155**, 157–216.
- D. A. Barskiy, A. M. Coffey, P. Nikolaou, D. M. Mikhaylov, B. M. Goodson, R. T. Branca, G. J. Lu, M. G. Shapiro, V. V. Telkki, V. V. Zhivonitko, I. V. Koptug, O. G. Salnikov, K. V. Kovtunov, V. I. Bukhtiyarov, M. S. Rosen, M. J. Barlow, S. Safavi, I. P. Hall, L. Schröder and E. Y. Chekmenev, *Chem.–Eur. J.*, 2017, **23**, 725–751.
- B. C. Anger, G. Schrank, A. Schoeck, K. A. Butler, M. S. Solum, R. J. Pugmire and B. Saam, *Phys. Rev. A: At., Mol., Opt. Phys.*, 2008, **78**, 43406.
- A. Bifone, Y. Q. Song, R. Seydoux, R. E. Taylor, B. M. Goodson, T. Pietrass, T. F. Budinger, G. Navon and A. Pines, *Proc. Natl. Acad. Sci. U. S. A.*, 1996, **93**, 12932–12936.
- J. P. Mugler III and T. A. Altes, *J. Magn. Reson. Imaging*, 2013, **37**, 313–331.
- G. Norquay, G. Leung, N. J. Stewart, G. M. Tozer, J. Wolber and J. M. Wild, *Magn. Reson. Med.*, 2015, **74**, 303–311.
- M. S. Albert, G. D. Cates, B. Driehuys, W. Happer, B. Saam, C. S. Springer and A. Wishnia, *Nature*, 1994, **370**, 199–201.
- W. Ruan, J. Zhong, K. Wang, G. Wu, Y. Han, X. Sun, C. Ye and X. Zhou, *J. Magn. Reson. Imaging*, 2017, **45**, 879–888.
- T. N. Matin, N. Rahman, A. H. Nickol, M. Chen, X. Xu, N. J. Stewart, T. Doel, V. Grau, J. M. Wild and F. V. Gleeson, *Radiology*, 2017, **282**, 857–868.
- B. Driehuys, S. Martinez-Jimenez, Z. I. Cleveland, G. M. Metz, D. M. Beaver, J. C. Noulis, S. S. Kaushik, R. Firszt, C. Willis, K. T. Kelly, J. Wolber, M. Kraft and P. H. McAdams, *Radiology*, 2012, **262**, 279–289.
- Y. Shukla, A. Wheatley, M. Kirby, S. Svenningsen, A. Farag, G. E. Santyr, N. A. M. Paterson, D. G. McCormack and G. Parraga, *Acad. Radiol.*, 2012, **19**, 941–951.
- J. P. Mugler III, B. Driehuys, J. R. Brookeman, G. D. Cates, S. S. Berr, R. G. Bryant, T. M. Daniel, E. E. De Lange, J. H. Downs, C. J. Erickson, W. Happer, D. P. Hinton, N. F. Kassel, T. Maier, C. D. Phillips, B. T. Saam, K. L. Sauer and M. E. Wagshul, *Magn. Reson. Med.*, 1997, **37**, 809–815.
- M. R. Rao, N. J. Stewart, P. D. Griffiths, G. Norquay and J. M. Wild, *Radiology*, 2018, **286**, 659–665.
- M. R. Rao, N. J. Stewart, G. Norquay, P. D. Griffiths and J. M. Wild, *Magn. Reson. Med.*, 2016, **75**, 2227–2234.
- J. P. Mugler III, B. Driehuys, J. R. Brookeman, G. D. Cates, S. S. Berr, R. G. Bryant, T. M. Daniel, E. E. De Lange, J. H. Downs, C. J. Erickson, W. Happer, D. P. Hinton, N. F. Kassel, T. Maier, C. D. Phillips, B. T. Saam, K. L. Sauer and M. E. Wagshul, *Magn. Reson. Med.*, 1997, **37**, 809–815.
- M. G. Shapiro, R. M. Ramirez, L. J. Sperling, G. Sun, J. Sun, A. Pines, D. V. Schaffer and V. S. Bajaj, *Nat. Chem.*, 2014, **6**, 629–634.
- Y. Wang, B. W. Roose, E. J. Palovcak, V. Carnevale and I. J. Dmochowski, *Angew. Chem., Int. Ed.*, 2016, **55**, 8984–8987.
- B. W. Roose, S. D. Zemerov and I. J. Dmochowski, *Chem. Sci.*, 2017, **8**, 7631–7636.
- B. W. Roose, S. D. Zemerov, Y. Wang, M. Kasimova, V. Carnevale and I. J. Dmochowski, *ChemPhysChem*, 2019, **20**, 260–267.
- S. D. Zemerov, B. W. Roose, K. L. Farenhem, Z. Zhao, M. A. Stringer, A. R. Goldman, D. W. Speicher and I. J. Dmochowski, *Anal. Chem.*, 2020, **92**, 12817–12824.
- Z. Zhao, B. W. Roose, S. D. Zemerov, M. A. Stringer and I. J. Dmochowski, *Chem. Commun.*, 2020, **56**, 11122–11125.
- O. Taratula and I. J. Dmochowski, *Curr. Opin. Chem. Biol.*, 2010, **14**, 97–104.
- Y. Wang and I. J. Dmochowski, *Acc. Chem. Res.*, 2016, **49**, 2179–2187.
- L. Garel, J.-P. Dutasta and A. Collet, *Angew. Chem., Int. Ed.*, 1993, **32**, 1169–1171.
- G. Huber, T. Brotin and L. Dubois, *J. Am. Chem. Soc.*, 2006, **128**, 6239–6246.
- J. Gabard and A. Collet, *J. Chem. Soc., Chem. Commun.*, 1981, 1137.



- 32 T. Brotin and J.-P. Dutasta, *Chem. Rev.*, 2009, **109**, 88–130.
- 33 T. Brotin and J.-P. Dutasta, *Compr. Supramol. Chem. II*, 2017, **1**, 317–335.
- 34 K. Bartik, M. Luhmer, J.-P. Dutasta, A. Collet and J. Reisse, *J. Am. Chem. Soc.*, 1998, **120**, 784–791.
- 35 M. M. Spence, S. M. Rubin, I. E. Dimitrov, E. J. Ruiz, D. E. Wemmer, A. Pines, S. Q. Yao, F. Tian and P. G. Schultz, *Proc. Natl. Acad. Sci. U. S. A.*, 2001, **98**, 10654–10657.
- 36 O. Taratula, P. A. Hill, Y. Bai, N. S. Khan and I. J. Dmochowski, *Org. Lett.*, 2011, **13**, 1414–1417.
- 37 P. A. Hill, Q. Wei, R. G. Eckenhoff and I. J. Dmochowski, *J. Am. Chem. Soc.*, 2007, **129**, 9262–9263.
- 38 P. A. Hill, Q. Wei, T. Troxler and I. J. Dmochowski, *J. Am. Chem. Soc.*, 2009, **131**, 3069–3077.
- 39 Y. Bai, P. A. Hill and I. J. Dmochowski, *Anal. Chem.*, 2012, **84**, 9935–9941.
- 40 D. R. Jacobson, N. S. Khan, R. Colle, R. Fitzgerald, L. Laureano-Perez, Y. Bai and I. J. Dmochowski, *Proc. Natl. Acad. Sci. U. S. A.*, 2011, **108**, 10969–10973.
- 41 G. K. Seward, Y. Bai, N. S. Khan and I. J. Dmochowski, *Chem. Sci.*, 2011, **2**, 1103–1110.
- 42 J. M. Chambers, P. A. Hill, J. A. Aaron, Z. Han, D. W. Christianson, N. N. Kuzma and I. J. Dmochowski, *J. Am. Chem. Soc.*, 2009, **131**, 563–569.
- 43 J. A. Aaron, J. M. Chambers, K. M. Jude, L. Di Costanzo, I. J. Dmochowski and D. W. Christianson, *J. Am. Chem. Soc.*, 2008, **130**, 6942–6943.
- 44 O. Taratula, Y. Bai, E. L. D'Antonio and I. J. Dmochowski, *Supramol. Chem.*, 2015, **27**, 65–71.
- 45 Q. Wei, G. K. Seward, P. A. Hill, B. Patton, I. E. Dimitrov, N. N. Kuzma and I. J. Dmochowski, *J. Am. Chem. Soc.*, 2006, **128**, 13274–13283.
- 46 A. Schlundt, W. Kilian, M. Beyermann, J. Sticht, S. Günther, S. Höpner, K. Falk, O. Roetzschke, L. Mitschang and C. Freund, *Angew. Chem., Int. Ed.*, 2009, **48**, 4142–4145.
- 47 N. Kotera, N. Tassali, E. Léonce, C. Boutin, P. Berthault, T. Brotin, J.-P. Dutasta, L. Delacour, T. Traoré, D. A. Buisson, F. Taran, S. Coudert and B. Rousseau, *Angew. Chem., Int. Ed.*, 2012, **51**, 4100–4103.
- 48 N. Tassali, N. Kotera, C. Boutin, E. Léonce, Y. Boulard, B. Rousseau, E. Dubost, F. Taran, T. Brotin, J.-P. Dutasta and P. Berthault, *Anal. Chem.*, 2014, **86**, 1783–1788.
- 49 J. Zhang, W. Jiang, Q. Luo, X. Zhang, Q. Guo, M. Liu and X. Zhou, *Talanta*, 2014, **122**, 101–105.
- 50 K. Jeong, C. C. Slack, C. C. Vassiliou, P. Dao, M. D. Gomes, D. J. Kennedy, A. E. Truxal, L. J. Sperling, M. B. Francis, D. E. Wemmer and A. Pines, *ChemPhysChem*, 2015, **16**, 3573–3577.
- 51 V. Roy, T. Brotin, J.-P. Dutasta, M. H. Charles, T. Delair, F. Mallet, G. Huber, H. Desvaux, Y. Boulard and P. Berthault, *ChemPhysChem*, 2007, **8**, 2082–2085.
- 52 P. Berthault, H. Desvaux, T. Wendlinger, M. Gyejacquot, A. Stopin, T. Brotin, J.-P. Dutasta and Y. Boulard, *Chem.–Eur. J.*, 2010, **16**, 12941–12946.
- 53 E. Léonce, J. P. Dognon, D. Pitrat, J. C. Mulatier, T. Brotin and P. Berthault, *Chem.–Eur. J.*, 2018, **24**, 6534–6537.
- 54 S. Yang, W. Jiang, L. Ren, Y. Yuan, B. Zhang, Q. Luo, Q. Guo, L.-S. Bouchard, M. Liu and X. Zhou, *Anal. Chem.*, 2016, **88**, 5835–5840.
- 55 N. Kotera, E. Dubost, G. Milanole, E. Doris, E. Gravel, N. Arhel, T. Brotin, J.-P. Dutasta, J. Cochrane, E. Mari, C. Boutin, E. Leonce, P. Berthault and B. Rousseau, *Chem. Commun.*, 2015, **51**, 11482–11484.
- 56 E. Mari, Y. Bousmah, C. Boutin, E. Léonce, G. Milanole, T. Brotin, P. Berthault and M. Erard, *ChemBioChem*, 2019, **20**, 1450–1457.
- 57 M. M. Spence, E. J. Ruiz, S. M. Rubin, T. J. Lowery, N. Winssinger, P. G. Schultz, D. E. Wemmer and A. Pines, *J. Am. Chem. Soc.*, 2004, **126**, 15287–15294.
- 58 T. Brotin, R. Barbe, M. Darzac and J.-P. Dutasta, *Chem.–Eur. J.*, 2003, **9**, 5784–5792.
- 59 L. Gao, W. Liu, O.-S. Lee, I. J. Dmochowski and J. G. Saven, *Chem. Sci.*, 2015, **6**, 7238–7248.
- 60 O. Taratula, P. A. Hill, N. S. Khan, P. J. Carroll and I. J. Dmochowski, *Nat. Commun.*, 2010, **1**, 148.
- 61 B. A. Riggall, M. L. Greenberg, Y. Wang, R. F. Wissner, S. D. Zemerov, J. Petersson and I. J. Dmochowski, *Org. Biomol. Chem.*, 2017, **15**, 8883–8887.
- 62 S. D. Zemerov, B. W. Roose, M. L. Greenberg, Y. Wang and I. J. Dmochowski, *Anal. Chem.*, 2018, **90**, 7730–7738.
- 63 T. J. Lowery, S. Garcia, L. Chavez, E. J. Ruiz, T. Wu, T. Brotin, J.-P. Dutasta, D. S. King, P. G. Schultz, A. Pines and D. E. Wemmer, *ChemBioChem*, 2006, **7**, 65–73.
- 64 D. N. Sears and C. J. Jameson, *J. Chem. Phys.*, 2003, **119**, 12231–12244.
- 65 S. D. Zemerov, Y. Lin and I. J. Dmochowski, *Anal. Chem.*, 2021, **93**, 1507–1514.
- 66 K. K. Palaniappan, M. B. Francis, A. Pines and D. E. Wemmer, *Isr. J. Chem.*, 2014, **54**, 104–112.
- 67 G. K. Seward, Q. Wei and I. J. Dmochowski, *Bioconjugate Chem.*, 2008, **19**, 2129–2135.
- 68 L. Schröder, T. J. Lowery, C. Hilty, D. E. Wemmer and A. Pines, *Science*, 2006, **314**, 446–449.
- 69 T. Meldrum, K. L. Seim, V. S. Bajaj, K. K. Palaniappan, W. Wu, M. B. Francis, D. E. Wemmer and A. Pines, *J. Am. Chem. Soc.*, 2010, **132**, 5936–5937.
- 70 Y. Bai, Y. Wang, M. Goulian, A. Driks and I. J. Dmochowski, *Chem. Sci.*, 2014, **5**, 3197–3203.
- 71 B. A. Riggall, Y. Wang and I. J. Dmochowski, *J. Am. Chem. Soc.*, 2015, **137**, 5542–5548.
- 72 Q. Zeng, Q. Guo, Y. Yuan, Y. Yang, B. Zhang, L. Ren, X. Zhang, Q. Luo, M. Liu, L.-S. Bouchard and X. Zhou, *Anal. Chem.*, 2017, **89**, 2288–2295.
- 73 G. Milanole, B. Gao, A. Paoletti, G. Pieters, C. Dugave, E. Deutsch, S. Rivera, F. Law, J. L. Perfettini, E. Mari, E. Léonce, C. Boutin, P. Berthault, H. Volland, F. Fenaille, T. Brotin and B. Rousseau, *Bioorg. Med. Chem.*, 2017, **25**, 6653–6660.
- 74 A. M. Oros and N. J. Shah, *Phys. Med. Biol.*, 2004, **49**, R105–R153.
- 75 M. S. Albert and D. Balamore, *Nucl. Instrum. Methods Phys. Res., Sect. A*, 1998, **402**, 441–453.
- 76 P. Berthault, G. Huber and H. Desvaux, *Prog. Nucl. Magn. Reson. Spectrosc.*, 2009, **55**, 35–60.



- 77 C. Boutin, A. Stopin, F. Lenda, T. Brotin, J.-P. Dutasta, N. Jamin, A. Sanson, Y. Boulard, F. Leteurtre, G. Huber, A. Bogaert-Buchmann, N. Tassali, H. Desvaux, M. Carrière and P. Berthault, *Bioorg. Med. Chem.*, 2011, **19**, 4135–4143.
- 78 H. M. Rose, C. Witte, F. Rossella, S. Klippel, C. Freund and L. Schröder, *Proc. Natl. Acad. Sci. U. S. A.*, 2014, **111**, 11697–11702.
- 79 C. Witte, V. Martos, H. M. Rose, S. Reinke, S. Klippel, L. Schröder and C. P. R. Hackenberger, *Angew. Chem., Int. Ed.*, 2015, **54**, 2806–2810.
- 80 S. Klippel, J. Döpfert, J. Jayapaul, M. Kunth, F. Rossella, M. Schnurr, C. Witte, C. Freund and L. Schröder, *Angew. Chem., Int. Ed.*, 2014, **53**, 493–496.
- 81 N. S. Khan, B. A. Riggle, G. K. Seward, Y. Bai and I. J. Dmochowski, *Bioconjugate Chem.*, 2015, **26**, 101–109.
- 82 S. Yang, Y. Yuan, W. Jiang, L. Ren, H. Deng, L. S. Bouchard, X. Zhou and M. Liu, *Chem.-Eur. J.*, 2017, **23**, 7648–7652.
- 83 M. Schnurr, C. Witte and L. Schröder, *Phys. Chem. Chem. Phys.*, 2013, **15**, 14178–14181.
- 84 R. D. Booker and A. K. Sum, *Biochim. Biophys. Acta, Biomembr.*, 2013, **1828**, 1347–1356.
- 85 G. L. Britton, H. Kim, P. H. Kee, J. Aronowski, C. K. Holland, D. D. McPherson and S. L. Huang, *Circulation*, 2010, **122**, 1578–1587.
- 86 M. P. Dandekar, T. Peng, D. D. McPherson, J. Quevedo, J. C. Soares and S. L. Huang, *Prog. Neuro-Psychopharmacol. Biol. Psychiatry*, 2018, **86**, 140–149.
- 87 Y. F. Miao, T. Peng, M. R. Moody, M. E. Klegerman, J. Aronowski, J. Grotta, D. D. McPherson, H. Kim and S. L. Huang, *Sci. Rep.*, 2018, **8**, 1–12.
- 88 Q. Guo, Q. Zeng, W. Jiang, X. Zhang, Q. Luo, X. Zhang, L.-S. Bouchard, M. Liu and X. Zhou, *Chem.-Eur. J.*, 2016, **22**, 3967–3970.
- 89 F. Schilling, L. Schröder, K. K. Palaniappan, S. Zapf, D. E. Wemmer and A. Pines, *ChemPhysChem*, 2010, **11**, 3529–3533.
- 90 M. Kunth, J. Döpfert, C. Witte, F. Rossella and L. Schröder, *Angew. Chem., Int. Ed.*, 2012, **51**, 8217–8220.
- 91 F. Rossella, H. M. Rose, C. Witte, J. Jayapaul and L. Schröder, *ChemPlusChem*, 2014, **79**, 1463–1471.
- 92 A. Piontek, C. Witte, H. M. Rose, M. Eichner, J. Protze, G. Krause, J. Piontek and L. Schröder, *Ann. N. Y. Acad. Sci.*, 2017, **1397**, 195–208.
- 93 M. Schnurr, K. Sydow, H. M. Rose, M. Dathe and L. Schröder, *Adv. Healthcare Mater.*, 2015, **4**, 40–45.
- 94 M. Schnurr, I. Volk, H. Nikolenko, L. Winkler, M. Dathe and L. Schröder, *Adv. Biosyst.*, 2020, **4**, 1900251.
- 95 J. L. Mynar, T. J. Lowery, D. E. Wemmer, A. Pines and J. M. J. Fréchet, *J. Am. Chem. Soc.*, 2006, **128**, 6334–6335.
- 96 T. K. Stevens, K. K. Palaniappan, R. M. Ramirez, M. B. Francis, D. E. Wemmer and A. Pines, *Magn. Reson. Med.*, 2013, **69**, 1245–1252.
- 97 K. K. Palaniappan, R. M. Ramirez, V. S. Bajaj, D. E. Wemmer, A. Pines and M. B. Francis, *Angew. Chem., Int. Ed.*, 2013, **52**, 4849–4853.
- 98 K. Jeong, C. Netirojjanakul, H. K. Munch, J. Sun, J. A. Finbloom, D. E. Wemmer, A. Pines and M. B. Francis, *Bioconjugate Chem.*, 2016, **27**, 1796–1801.
- 99 P. Berthault, A. Bogaert-Buchmann, H. Desvaux, G. Huber and Y. Boulard, *J. Am. Chem. Soc.*, 2008, **130**, 16456–16457.
- 100 R. Tyagi, C. Witte, R. Haag and L. Schröder, *Org. Lett.*, 2014, **16**, 4436–4439.
- 101 S. Malhotra, M. Calderón, A. K. Prasad, V. S. Parmar and R. Haag, *Org. Biomol. Chem.*, 2010, **8**, 2228–2237.
- 102 S. Klippel, C. Freund and L. Schröder, *Nano Lett.*, 2014, **14**, 5721–5726.
- 103 Y. Wang and I. J. Dmochowski, *Chem. Commun.*, 2015, **51**, 8982–8985.
- 104 Y. Wang, B. W. Roose, J. P. Philbin, J. L. Doman and I. J. Dmochowski, *Angew. Chem., Int. Ed.*, 2016, **55**, 1733–1736.
- 105 F. T. Hane, T. Li, P. Smylie, R. M. Pellizzari, J. A. Plata, B. DeBoef and M. S. Albert, *Sci. Rep.*, 2017, **7**, 41027.
- 106 K. Du, S. D. Zemerov, S.-H. Parra, J. M. Kikkawa and I. J. Dmochowski, *Inorg. Chem.*, 2020, **59**, 13831–13844.
- 107 K. Du, S. D. Zemerov, P. J. Carroll and I. J. Dmochowski, *Inorg. Chem.*, 2020, **59**, 12758–12767.
- 108 T. Traoré, L. Delacour, S. Garcia-Argote, P. Berthault, J.-C. Cintrat and B. Rousseau, *Org. Lett.*, 2010, **12**, 960–962.
- 109 L. Delacour, N. Kotera, T. Traoré, S. Garcia-Argote, C. Puente, F. Leteurtre, E. Gravel, N. Tassali, C. Boutin, E. Léonce, Y. Boulard, P. Berthault and B. Rousseau, *Chem.-Eur. J.*, 2013, **19**, 6089–6093.
- 110 M. Chen, O. Doganay, T. Martin, A. McIntyre, N. Rahman, D. Bulte and F. Gleeson, *Eur. Radiol.*, 2020, **30**, 1145–1155.
- 111 J. Chacon-Caldera, A. Maunder, M. R. Rao, G. Norquay, O. I. Rodgers, M. Clemence, C. Puddu, L. R. Schad and J. M. Wild, *Magn. Reson. Med.*, 2020, **83**, 262–270.
- 112 E. J. Song, C. R. Kelsey, B. Driehuys and L. Rankine, *J. Med. Imaging Radiat. Oncol.*, 2018, **62**, 91–93.
- 113 L. Ebner, M. He, R. S. Virgincar, T. Heacock, S. S. Kaushik, M. S. Freemann, H. P. McAdams, M. Kraft and B. Driehuys, *Invest. Radiol.*, 2017, **52**, 120–127.
- 114 J. M. Wild, G. Collier, H. Marshall, L. Smith, G. Norquay, A. J. Swift, F. C. Horn, F. Chan, N. J. Stewart, L. C. Hutchison, M. R. Rao, I. Sabbroe, R. Niven, A. Horsley, S. Siddiqui, K. Ugonna and R. Lawson, *Thorax*, 2015, **70**, A220–A221.
- 115 J. R. Birchall, P. Nikolaou, A. M. Coffey, B. E. Kidd, M. Murphy, M. Molway, L. B. Bales, B. M. Goodson, R. K. Irwin, M. J. Barlow and E. Y. Chekmenev, *Anal. Chem.*, 2020, **92**, 4309–4316.
- 116 J. R. Birchall, R. K. Irwin, P. Nikolaou, A. M. Coffey, B. E. Kidd, M. Murphy, M. Molway, L. B. Bales, K. Ranta, M. J. Barlow, B. M. Goodson, M. S. Rosen and E. Y. Chekmenev, *J. Magn. Reson.*, 2020, **319**, 106813.

