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Metal-catalyzed uncaging of DNA-binding agents in living cells

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Attachment of alloc protecting groups to the amidine units of fluorogenic DNA-binding bisbenzamidines or to the amino groups of ethidium bromide leads to a significant reduction of their DNA affinity. More importantly, the active DNA-binding species can be readily regenerated by treatment with ruthenium catalysts in aqueous conditions, even in cell cultures. The catalytic chemical uncaging can be easily monitored by fluorescence microscopy because the protected products display both different emission properties and cell distribution than the parent compounds.

Introduction

A major research goal in chemical biology is the efficient and selective targeting of double stranded DNA with small molecules. Towards this goal, a wide range of synthetic DNA binders, from small molecules to larger peptides have been developed over the years. In addition to the search for better sequence selectivity, there is also an increased focus in the external control of the DNA binding of these molecules, so that they could be activated at will, when and where required. In this context, we have recently described a photo-uncaging strategy for controlling the non-covalent DNA interaction of DNA-binding agents, such as those shown in the Figure 1.

Fig 1 Structure of selected DNA binders: pentamidine (1), propamidine (2), phenyl azapentamidine (3), DAPI (4) ethidium bromide (5).

Although irradiation with light represents a clean and simple way of activating molecules in a spatiotemporal controlled way, the potential biological application of the strategy is limited by the low penetrability the UV light typically used for photolysis and the likely secondary damage to the tissues. Therefore, other alternatives for the generation of bioactive agents in an externally controlled manner are highly desirable. Particularly appealing is the possibility of releasing the active molecules by using a catalytic reaction that could be performed, for instance, by specific enzymes. Although the use of naturally occurring biocatalysts is attractive, its scope is intrinsically limited to a relatively small number of transformations, and to compounds that could be recognized as substrates by the enzymes. A promising alternative to enzymes could rely on the use of transition metal catalysts. Curiously, despite the extensive use of organometallic catalysis in synthetic chemistry, metal-based catalytic reactions have been largely overlooked in biological settings. While applying organometallic catalysis in aqueous media is not straightforward, several relevant metal-catalyzed reactions have been successfully used in biocompatible solvents, and even in cells, and it is foreseeable that the next years will witness a surge on the use of metal catalysis in biological environments. Particularly relevant in this area is the seminal work by E. Meggers and coworkers, who have demonstrated that ruthenium catalysts, combined with thiophenol, can be used to uncage alloc-protected rhodamines in aqueous media (MeOH/H2O, 95:5), and even inside HeLa cells, without remarkably influencing the cell viability.

Owing to our recent demonstration that caging the amidinium groups of bisbenzamidines or the amino groups of ethidium with photolabile nitrobenzyl groups suppresses their DNA interaction, we wondered whether simple alloc-protecting groups might also prevent DNA binding, which would open the possibility of using metal-π-allyl chemistry for catalytic uncaging.

Herein we demonstrate the viability of this temporary allyl-carbamate protecting approach for controlling the DNA binding of representative minor groove binders, such as bisbenzamidines, and of classical intercalators like ethidium bromide, Importantly, we also show that the uncaging reaction can be efficiently performed in living cells, and that the redistribution of the released dyes can be monitored by fluorescence microscopy. This work represents the first demonstration of the use of metal catalysis to trigger DNA binding events in live cells.

Results and Discussion
Synthesis and \textit{in vitro} fluorescence studies of protected derivatives of 3

Bisbenzamidines like pentamidine (1) or propamidine (2) are structurally simple DNA binders that show good stability, sequence selectivity, and excellent internalization properties in a variety of cell lines. Although rather toxic, pentamidine and some of its derivatives are being used for the treatment of \textit{Pneumocystis carinii}, Leishmaniasis or early-phase African sleeping sickness, among other diseases. Their mode of action is attributed to their ability to interact with the A/T-rich mitochondrial DNA of the parasites. We have recently shown that substitution of the oxygen atom of classic bis-benzamidines by a nitrogen generates \textit{aza} derivatives, such as the phenyl \textit{aza}-pentamidine 3, with DNA-dependent fluorescent properties.

These derivatives can be easily synthesized in a single step by reductive amination of commercial dialdehydes. The fluorogenic properties of these molecules resemble those of other bisbenzamidine bis-\textit{aza} derivatives, such as the phenyl \textit{aza}-pentamidine 3, with DNA-dependent fluorescent properties. The fluorigenic properties of these molecules resemble those of other DNA binders that are widely used as fluorescent stains, such as the blue nuclear stain DAPI (4',6-diamidino-2-phenylindole, 4; Fig 1), or ethidium bromide (5, Fig 1).

Installation of the alloc protecting groups in the \textit{aza}-pentamidine 3 was readily accomplished by reaction with allyl chloroformiate in the presence of NaOH, in H2O/THF (85% yield). The resulting dialloc protected derivative was readily accomplished by reaction with allyl chloroformiate in the presence of NaOH, in H2O/THF (85% yield). The resulting dialloc protected derivative in the presence of NaOH, in H2O/THF (85% yield). The resulting dialloc protected derivative in the presence of NaOH, in H2O/THF (85% yield). The resulting dialloc protected derivative in the presence of NaOH, in H2O/THF (85% yield). The resulting dialloc protected derivative.

As expected, while incubation of 3 with allyl derivative displays a relatively weak fluorescence intensity and slightly red-shifted emission in comparison to 3. As expected, while incubation of 3 with a hairpin oligonucleotide featuring an A/T-rich site (h-A3•T3) leads to a large increase in its fluorescence emission upon irradiation at 320 nm, 3 displays only a very modest emission intensity enhancement under the same conditions (Fig 2).

Titration of 3 with the oligonucleotide h-A3•T3 allowed us to calculate an apparent dissociation constant of \( K_d \approx 21 \mu M \), which is over 30 times weaker than that exhibited by 3 for this same oligo (\( K_d \approx 0.7 \mu M \)). The lower DNA affinity of the alloc-protected 3 with respect to 3 can be attributed to the loss of the electrostatic and hydrogen bonding interactions of the amidinium groups in the caged derivative. This hypothesis is consistent with the results obtained with the analogous bisbenzamidine bis-\textit{allyl} derivative 6 that, in contrast to 3, displays comparable binding affinity to that of the unprotected amidinium derivative 3 (\( K_d \approx 0.5 \mu M \), see ESI). As expected, 3 also shows a very small increase in its emission intensity upon incubation with a G/C-rich oligonucleotide (see the ESI).

\textbf{Catalytic uncaging}

The deprotection reaction was first carried out by treatment of 3 with thiophenol (5 equiv) and 10% of ruthenium catalyst \textit{RuCp*(COD)Cl} ([Ru]), at rt in MeOH/H2O (95:5). HPLC analysis of the reaction mixture after 20 min barely shows any starting material, and the major peak corresponds to the parent bisbenzamidine 3. Importantly, the reaction can also be efficiently carried out in more biologically relevant media (phosphate buffer 100 mM, pH 7.5, and 10% DMSO), although under these conditions, after 20 min we still observe a minor amount of the mono-protected product m-3 (Fig 3). Keeping the reaction for a longer time did not produce a significant change in the proportion of the products (85:15).

The deprotection reaction could also be carried out with Pd catalysts using standard conditions for alloc removal. Thus, treatment of 3 with Pd(OAc)2, N-methyl morpholine, PPh3 and PhSiH3 in 100 mM phosphate buffer pH 7.5, and 10% DMSO, led after 5 min to an almost quantitative conversion to 3; as expected, this deprotection did not proceed in absence of the

\begin{figure}[h]
\centering
\includegraphics[width=\textwidth]{fig2.png}
\caption{Fig 2 Top: Structures of the alloc (©3), and allyl (©6) protected derivatives of bisbenzamidine 3. Bottom: Fluorescence emission of 5 \( \mu M \) solutions of 3 and ©3 in Tris-HCl 20 mM, 100 mM NaCl, pH 7.5 (dashed lines), and in the presence of 1 equiv of h-A3•T3 (solid lines). h-A3•T3: 5'−GCC-AAATTTCAGTTTTCCTG-AAATTTCGC−3'.}
\end{figure}

\begin{figure}[h]
\centering
\includegraphics[width=\textwidth]{fig3.png}
\caption{Fig 3 Top: Catalytic deprotection in phosphate buffer and 10% DMSO. Bottom: HPLC trace of the uncaging reaction in phosphate buffer 100 mM pH 7.5, and 10% DMSO. a) ©3 (75 mM); b) ©3 + PhSH (325 mM); c) ©3 + PhSH + [Ru] (10%), t = 1 min; d) ©3 + PhSH + [Ru], t = 20 min. The deprotection reaction could also be carried out with Pd catalysts using standard conditions for alloc removal. Thus, treatment of ©3 with Pd(OAc)2, N-methyl morpholine, PPh3 and PhSiH3 in 100 mM phosphate buffer pH 7.5, and 10% DMSO, led after 5 min to an almost quantitative conversion to ©3; as expected, this deprotection did not proceed in absence of the}
\end{figure}
additives (see the ESI).

The Ru-catalyzed deprotection also takes place in the presence of calf thymus DNA (ct-DNA), although in this case after 20 min at rt we observed the formation of a 15:45:40 mixture of the free (3), monoprotected (m-©3) and fully protected (©©3) products, respectively (see the ESI). Given that the deprotection reaction generates different amounts of the monoprotected derivative m-©3, we studied its DNA binding properties. Not surprisingly titration of m-©3 with h-A2•T2 revealed that this compound binds to the DNA with an intermediate affinity to that of the bis-alloc-protected product ©©3 and the parent unprotected molecule 3 (Kd [m-©3] ≈ 6.0 µM, see the ESI).

Deprotection and DNA binding studies of DAPI and Ethidium derivatives

As with ©3, the bis-alloc DAPI derivative ©4 was readily obtained in good yield by treatment of 4 with allyl chloroformiate and Et3N in DMSO. In contrast to DAPI, ©4 displays a weak intrinsic fluorescence, but experienced a comparatively lower increase in emission when mixed with the h-A2•T2 hairpin oligonucleotide (Fig 4). Steady state fluorescence titration of ©4 with this hairpin oligo revealed a Kd [©4] ≈ 0.41 µM, which represents more than 40 times weaker affinity than that of DAPI (Kd [©4] ≤ 10 nM).17 Likewise, as observed previously with 3, inhibition of the DNA binding required the protection of both amidinium groups, since the single protected DAPI still exhibited significant DNA binding affinity (Kd [m-©4] ≈ 49 nM, see the ESI).

The Ru-catalyzed uncaging of ©4 was faster and more efficient than that of ©3. Thus, treatment of ©4 (25 mM) with thiophenol (5 equiv) and 10% of the ruthenium catalyst [Ru], in 100 mM phosphate buffer pH 7.5 and 10% DMSO in the presence of ct-DNA led, after only 5 min at rt, to the desired parent compound 4 (clean conversion by HPLC).

Finally, the catalytic uncaging strategy was also tested with ethidium bromide (©5), a classic intercalator that is widely used for DNA and RNA staining.14 Commercially available ethidium bromide (©5) reacted efficiently with allyl chloroformiate in DIEA/DMF to yield the desired dialloc derivative ©6 as a bright yellow solid (yield ≈ 58%). Remarkably, in contrast to the parent ethidium, the biscarbamate ©6 is highly fluorescent in aqueous solution in the absence of DNA, displaying a maximum emission wavelength at 510 nm. While the addition of ct-DNA to ethidium induces a significant increase in its fluorescence emission at 595 nm, addition of ct-DNA to the bis-caged ethidium ©5 leads to a decrease in its emission.15 As in the case of the minor groove binders discussed before, the caged derivative displayed weaker DNA binding than the parent ethidium; near a 20-fold decrease in affinity for h-AGA (see the ESI).

With regard to the catalytic uncaging: treatment of ©6 (50 mM) with thiophenol (5 equiv) and 10% of [Ru] in phosphate buffer 100 mM pH 7.5 and 10% DMSO at rt in presence of ct-DNA, led to the expected uncaged products, although the reaction is slower than in the case of ©4. HPLC of the reaction mixture after 20 min showed the formation of mono-uncaged derivatives; however, after 1 h most of the starting material had been transformed into the desired fully deprotected product (©5). Interestingly, since the absorption of the protected derivative (©6) is different to that of the parent ethidium, the reaction progress can be monitored by observing the changes in the solution color (inset in Fig 6). Competitive titrations with ©5 showed that the monoprotected ethidium derivatives bind DNA also worst than the parent ethidium (see the ESI).
The above results validate the attachment of allo derivatives as an efficient strategy to considerably inhibit the DNA binding of three different types of small DNA-binding agents and confirm the viability of a metal-catalyzed removal of the caging group under aqueous (and hypothetically biocompatible) conditions. We explored the possibility of performing the catalytic de-allylation process to induce their DNA binding in cellular settings. We focused our efforts in the study of DAPI and ethidium derivatives, because their cellular distribution might be easily monitored by fluorescence microscopy thanks to the intrinsic fluorescence of the protected dyes and the characteristic blue staining of the whole cell indicating that the alloc-protected DAPI is internalized very efficiently, and it was required the use of higher concentrations of DAPI (up to 15 µM) to see a clear staining (Fig 7h). This enhanced staining with the dye evenly distributed between the cytoplasm and the cell nuclei (Fig 8a); addition of the reagents (2.5 µM Ru-catalyzed deprotection of 2.5 µM DAPI and 100 µM PhSH) led to a more or less homogeneous display a phenotype viability (see the ESI). Moreover, comparative staining with propidium iodide (20 µM) showed that treatment of CEF untreated cultures (see the ESI). As expected, the integrity of cell membranes (see the ESI). As expected, the integrity of cell membranes was maintained after 20 min of treatment with 2.5 µM Ru(catalyzed deprotection of 2.5 µM DAPI and 100 µM PhSH) and 100 µM PhSH, the cell model (Fig 8a), addition of the reagents (2.5 µM [Ru] and 100 µM PhSH) led to a clear redistribution of the emission suggesting that this method does not significantly leak out of the cell (see the ESI). While further studies are needed, these results provide support for an intracellular uncaging reaction.

Fig 6 HPLC trace of the deprotection reaction of C5 at t= 1 h. 0.60 g (50 mM) c-DNA + Ru(10%) t= 0; d) c-DNA + Ru(10%) t= 20 min; e) c-DNA + Ru(10%) t= 30 min; f) c-DNA + Ru(10%) t= 40 min; g) c-DNA + Ru(10%) t= 50 min; h) c-DNA + Ru(10%) t= 60 min; i) c-DNA + Ru(10%) t= 70 min; j) c-DNA + Ru(10%) t= 80 min; k) c-DNA + Ru(10%) t= 90 min; l) c-DNA + Ru(10%) t= 100 min; m) c-DNA + Ru(10%) t= 110 min; n) c-DNA + Ru(10%) t= 120 min; o) c-DNA + Ru(10%) t= 130 min; p) c-DNA + Ru(10%) t= 140 min; q) c-DNA + Ru(10%) t= 150 min; r) c-DNA + Ru(10%) t= 160 min; s) c-DNA + Ru(10%) t= 170 min; t) c-DNA + Ru(10%) t= 180 min; u) c-DNA + Ru(10%) t= 190 min; v) c-DNA + Ru(10%) t= 200 min; w) c-DNA + Ru(10%) t= 210 min; x) c-DNA + Ru(10%) t= 220 min; y) c-DNA + Ru(10%) t= 230 min; z) c-DNA + Ru(10%) t= 240 min.

In consonance with previous findings by Meggers et al., we found that both the metal catalyst and the thiophenol additive are required to obtain an efficient cellular uncaging reaction. As expected, the integrity of cell membranes was maintained after 20 min of treatment with 2.5 µM Ru(catalyzed deprotection of 2.5 µM DAPI and 100 µM PhSH) and 100 µM PhSH, the cell model (Fig 8a), addition of the reagents (2.5 µM [Ru] and 100 µM PhSH) led to a clear redistribution of the emission.
into the cell nuclei (Fig 8c) as expected for the release of the uncaged DAPI and its sticking to nuclear DNA.

Fig 8 Ru-catalyzed deprotection of 2.5 µM of C4 in Vero cells. Top row: C4, 30 min at 37 ºC; a) blue channel; b) brightfield; c) red channel. Bottom row: C4 with 2.5 µM [Ru] and 100 µM PhSH, 20 min; c) blue channel; d) brightfield.

Fig 9 Ru-catalyzed deprotection of 10 µM C5 in CEF cells. Left column: C5, 30 min at 37 ºC; a) green channel; b) brightfield; c) red channel. Right column: C5 incubated with 20 µM [Ru] and 100 µM PhSH, 20 min; d) green channel; e) brightfield; f) red channel. Fluorescence microscopy settings for green channel: filter 530–550 nm, emission filter 590 nm and dichromatic mirror 570 nm; and for red channel: filter 530–550 nm, emission filter 590 nm and dichromatic mirror 570 nm.

With regard to the uncaging of ethidium, the distinctive emission pattern of the protected and unprotected derivatives allowed a detailed monitoring of the uncaging process. Thus, C5 displays a clear green staining of the cells (excitation filter 460–490 nm/emission filter 520 nm) with a preferential accumulation of the caged dye in cytoplasmatic structures (Fig 9a); as expected from the fluorescence spectrum (Fig 5) no emission was observed in the red channel (excitation filter 530–550 nm/emission filter 590 nm). Treatment of the cells with [Ru] (20 µM) and PhSH (100 µM) for 20 min led to the characteristic staining of ethidium bromide in the red channel, with the emission mainly concentrated in the nuclei and nucleoli (Fig. 7f), while almost no signal was observed in the green channel (Fig. 7c), suggesting that the uncaging has been very efficient.

Biological activity

The above results demonstrate that caging key functional groups of DNA binding agents, even with simple alloc moieties, results in a significant decrease on their DNA affinity. A potential biomedical application of this chemistry demanded the demonstration that such decrease in DNA binding is also associated with a decreased biological effect. Therefore, we checked the activity of the protected (C3) and unprotected bis-benzamidines (3) against cancer cell lines. Interestingly, in the cisplatin-resistant A2780 cell line (Human ovarian carcinoma) the inhibitory effect of the caged derivative was more than ten times lower than that of the parent aza pentamidine (IC50[C3] ≈ 5.0 µM, IC50[3] ≈ 0.4 µM), which in turn had a stronger inhibitory effect than cisplatin (IC50[cisplatin] ≈ 7.4 µM). Similar inhibitory effects of the alloc caging were also observed with the related A2780 EDP cell line (IC50[C3] ≈ 5.1 µM, IC50[3] ≈ 0.9, IC50[cisplatin] ≈ 0.8). The ruthenium catalyst had negligible inhibitory effect by itself with an IC50 ≥ 45 µM. These results open the door for future pro-drug strategies that rely in the combination of a caged precursors and an appropriate metal complex for controlled catalytic uncaging.

Conclusions

In conclusion, we have demonstrated that installing simple allylcarbamate groups in key positions of several DNA binders not only prevents their interaction with DNA but also modulates their spectroscopic properties and cellular location. In particular, we have demonstrated that the bis alloc-protected derivatives of DAPI or ethidium bromide are weaker DNA binders than the parent compounds, which can be efficiently regenerated in aqueous buffers, and even in intracellular environments, by reaction with a ruthenium catalytic system. Our results set the stage for future developments on metal-catalyzed activation of DNA-binding compounds in biological media.

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Notes and references

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† Electronic Supplementary Information (ESI) available: Synthesis and characterization of the studied molecules and required precursors, NMR, UV, and Fluorescence spectra, titrations, control experiments, and
detailed procedures for cell uptake and co-staining experiments. See DOI: 10.1039/b000000x


20 The incubation time required for uncaging (30 min) is much shorter than that used in the growth inhibition studies (96 h).