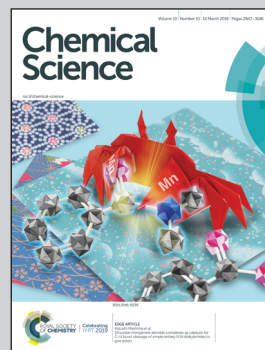


Showcasing research from Professor Michael Smietana's laboratory, Institut des Biomolécules Max Mousseron, University of Montpellier, France and Professor Stellios Arseniyadis' laboratory, School of Biological and Chemical Sciences, Queen Mary University of London, UK.

A rational quest for selectivity through precise ligand-positioning in tandem DNA-catalysed Friedel–Crafts alkylation/asymmetric protonation

We report here the development of a new DNA-based bio-hybrid catalyst in which the metallic co factor is precisely positioned within the DNA framework through a covalent anchoring strategy. Evaluated on the challenging copper(II)-catalysed asymmetric Friedel–Crafts alkylation/enantioselective protonation of α -substituted α,β -unsaturated enones, this approach showed unprecedented levels of enantioselectivity and broad scope, while unveiling specific structural features that account for an optimal chirality transfer from the duplex to the Friedel–Crafts adducts. Image courtesy of Bernard Günther.

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A rational quest for selectivity through precise ligand-positioning in tandem DNA-catalysed Friedel–Crafts alkylation/asymmetric protonation†

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Covalent anchorage of a metallic co-factor to a DNA-based architecture is merely the only way to ensure an accurate positioning of a catalytic site within the chiral micro-environment offered by the DNA double helix. Ultimately, it also allows a fine-tuning of the catalytic pocket through simple synthetic modifications of the DNA sequence. Here, we report highly selective copper(II)-catalysed asymmetric Friedel–Crafts conjugate addition/enantioselective protonation, which is due to a careful positioning of a bipyridine ligand within a DNA framework. Most importantly, this study unveils specific structural features that account for an optimal chirality transfer from the duplex to the Friedel–Crafts adducts.

Introduction

In the last decade, a large variety of artificial metalloenzymes based on various macromolecular architectures have been designed and engineered by chemists resulting in several particularly effective catalytic processes.¹ While the first protein-based artificial metalloenzymes date back to the 1970s,² it was only in 2005 that Roelfes and Feringa³ described the very first DNA-based artificial metalloenzyme, where the well-defined DNA structure provides a unique chiral microenvironment able to accommodate a transition metal complex which in turn catalysed Diels–Alder cycloaddition with high levels of enantiodiscrimination. Since then, DNA-based chiral amplifications have been applied to a wide range of Cu(II)-catalysed reactions including Friedel–Crafts alkylation,⁴ Michael addition,⁵ *syn*-hydration,⁶ and fluorination reactions.⁷ Among all the ligands tested, 2,2′-dimethyl-4,4′-bipyridine (dmbipy) was quickly identified as one of the most promising, which led multiple groups around the world including ours to evaluate the influence on the selectivity of various anchorage strategies by which the ligand can be incorporated in the DNA scaffold. While supramolecular interactions allow a straightforward self-assembly of the catalyst and thus a rapid screening/optimization of the reaction conditions, they unfortunately do

not enable a precise positioning of the metallic cofactor into the DNA duplex.¹ In contrast, the covalent attachment of a metallic cofactor allows the construction of finely tuned DNA-based catalysts with a clear knowledge of the surrounding environment, thus allowing a more straightforward rationalization of the catalytic efficacy of a given bio-hybrid system. Interestingly, while several groups have embarked on this route,⁸ none have taken into consideration the fundamental influence of the positioning of the metal-chelating ligand. This is all the more surprising that it is well known that DNA base modifications at the 5-position of the pyrimidines tend to place the substituents in the major groove of the double helix, while 2′-modifications place the substituents inside the minor groove.⁹ These fundamental structural differences prompted us to further investigate this matter with the aim of developing a strong mechanistic rational and eventually designing the ultimate bio-hybrid catalyst. The asymmetric Friedel–Crafts conjugate addition/enantioselective protonation of α -substituted α,β -unsaturated acyl imidazoles appeared to us as the perfect model reaction to study since the selectivities reported in the field were rather low and, most importantly, highly substrate dependent.^{4c} Indeed, the challenge in this reaction is that the chirality is not introduced during the conjugate addition step but rather during the protonation of the highly reactive pro-chiral enolate intermediate. As a general trend, the enantioselectivity in such reactions is usually obtained using a chiral enolate in conjunction with an achiral proton source or by using an achiral enolate in combination with a chiral protonating agent.¹⁰ In both cases however, exclusion of water is crucial in order to prevent any non-selective protonation. To the best of our knowledge only two examples of tandem Friedel–Crafts conjugate addition/asymmetric protonation in water have been reported in the literature so far. The first one was reported by Luo and Cheng in

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2011 and featured the conjugate addition of various indoles on α -substituted acroleins through chiral enamine catalysis.¹¹ The second one was reported more recently by Roelfes and co-workers and involved the conjugate addition of a variety of indoles on 2-methyl-1-(thiazol-2-yl)prop-2-en-1-one catalysed by DNA in conjunction with a Cu-dmbipy metallic cofactor.^{4c} Interestingly, while the presence of DNA was found to accelerate the reaction at an exceptional rate (up to 990-fold), the enantioselectivities were nonetheless highly dependent on the nature of the indole. Moreover, this study unveiled a close connexion between the selectivity obtained and the binding affinity of the Cu(II)-dmbipy complex, the enone and the indole with DNA.^{4c} This indispensable affinity balance between all the species involved inspired us to evaluate the effect of a covalent anchorage of the metallic co-factor onto the DNA scaffold and ultimately ascertain the influence of the grooves on the enantioselectivity.

Results and discussion

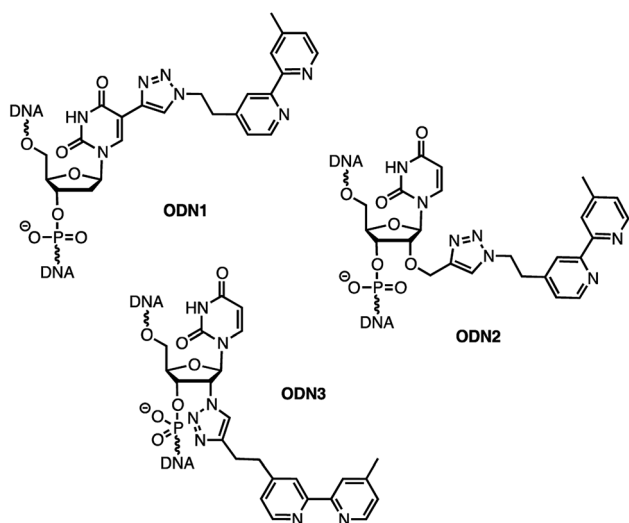
In order to analyse the influence of the positioning of the metallic co-factor on both the reactivity and the selectivity, we chose to synthesize **ODN1**, **ODN2** and **ODN3**, which incorporate a bipyridine ligand either in the major groove (**ODN1**) or in the minor groove (**ODN2** and **ODN3**) once hybridised with a common complementary strand **ODN4** (Scheme 1).

The usual methods to introduce modifications into oligonucleotides involve either the synthesis of appropriately modified phosphoramidite building blocks or the post-synthetic conjugation of a defined reactive group. This latter strategy is usually preferred as it is easier to handle, gives better yields and allows higher degrees of modularity.¹² As rate acceleration and improved enantioselectivities have been previously demonstrated with G-rich self-complementary dodecamers,¹³ we designed a duplex 5'-GCCAGCXGACCG-3'/5'-CGGTCAGCTGGC-3' which incorporates a unique modification (X) at residue 7 of the sense strand. Commercially available phosphoramidite

derivatives of 2'-deoxy-5-ethynyluridine **7** and 2'-O-propargyluridine **10** were used to prepare CPG-bound oligonucleotides **8** and **11** respectively (DMT-off). The azido-dmbipy partner **4**, on the other hand, was synthesized in three steps starting from commercially available 4,4'-dimethyl-2,2'-bipyridine **1**. Hence, monolithiation of **1** using LDA followed by the addition of an excess of paraformaldehyde afforded the corresponding hydroxymethyl derivative **2**,¹⁴ which was eventually subjected to tosylation [TsCl, DIEA, DCM, rt, and 57%] and azidation [NaN₃, DMF, 0 °C, and 85%] to afford the desired azide **4** (Scheme 2A). A Cu(I)-catalysed azide-alkyne cycloaddition (CuAAC) reaction was then carried out on a solid support by mixing the representative **ODN** with azide **4** (2 equiv.) in a 1 : 1 H₂O/dioxane mixture in the presence of a freshly prepared aqueous solution of CuSO₄·5H₂O (1 equiv.), sodium ascorbate (5 equiv.) and THPTA (3 equiv.), and by heating the mixture for 75 min at 55 °C under microwave irradiation (Scheme 2B and C).¹⁵ The resulting CPG supports were then filtered and washed with a saturated solution of EDTA. After treatment with aqueous ammonia, **ODN1** and **ODN2** were purified by preparative HPLC and finally desalted.

In order to probe the influence of the flexibility of the ligand inside the minor groove, we also synthesized **ODN3**, which bears a triazole moiety directly attached onto the 2'-position at residue 7 of the sense strand. This modification required the use of 2'-azido uridine which is not compatible with standard phosphoramidite chemistry because of the inherent reactivity of the phosphorus(III) species in the presence of azides.¹⁶ Although the post-synthetic labelling of 2'-azido modified RNA has been described by Micura and co-workers by combining both phosphotriester and phosphoramidite chemistries,¹⁷ we decided to embark on the chemical synthesis of the appropriately modified phosphoramidite **15** (Scheme 2D). In this context, the monolithiation of **1** using LDA at -78 °C followed by the addition of bromo-1-(trimethylsilyl)-1-propyne (1 equiv.) afforded **5**, which was ultimately deprotected with potassium fluoride under microwave irradiation. The resulting terminal alkyne **6** was then reacted with 5'-dimethoxytrityl-2'-azido-2'-deoxyuridine **13** in the presence of CuSO₄·5H₂O and sodium ascorbate [H₂O/THF/pyridine (6 : 3 : 2), 30 min, and rt] to afford triazole derivative **14**, which was subsequently converted to the corresponding phosphoramidite **15** using 2-cyanoethyl-*N,N*-diisopropylchlorophosphoramidite (1.3 equiv.). The requisite oligonucleotide **ODN3** was then assembled through solid-phase synthesis, purified by HPLC and desalted. **ODN1**, **ODN2** and **ODN3** were eventually hybridised with complementary **ODN4** (5'-CGG TCA GCT GGC-3') and evaluated in the context of the tandem Friedel-Crafts conjugate addition/asymmetric protonation reaction.

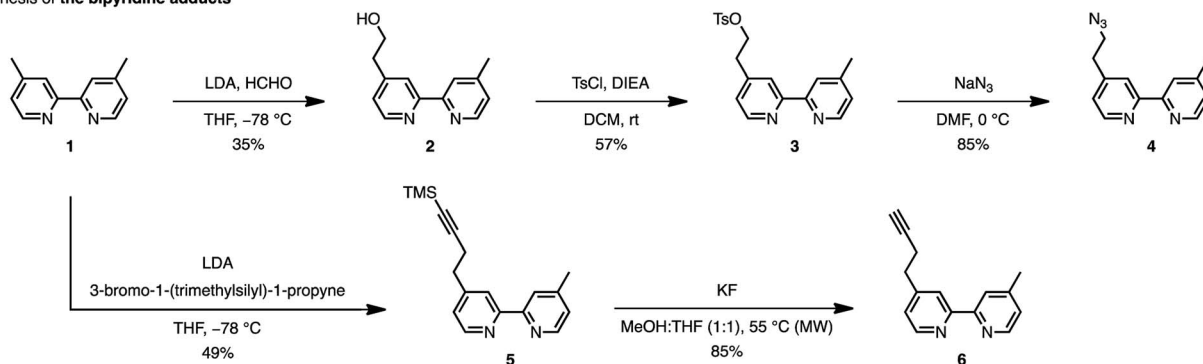
Based on previously optimized conditions, we decided to run the reactions using a 1 : 1 ratio of 2-methyl-1-(thiazol-2-yl)prop-2-en-1-one and indole, 30 mol% of Cu(NO₃)₂ and a 0.4 mM solution of **ODNx/ODN4** in a MES buffer (pH 5.0) at 4 °C for three days.¹⁸ The supramolecular approach using the non-covalently linked Cu-dmbipy complex in the presence of either st-DNA (2 mM base pair) or the non-modified duplex **ODN5/ODN4** (5'-GCCAGCTGACCG-3'/5'-CGGTCAGCTGGC-3')



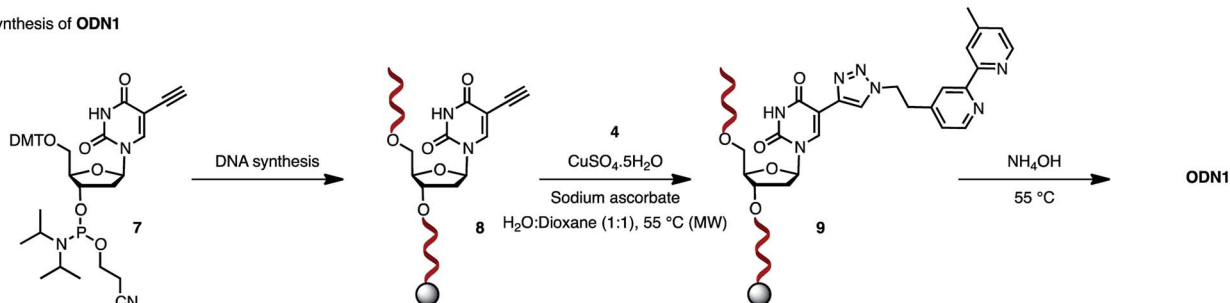
Scheme 1 Structures of bipyridine-functionalised ODNs.



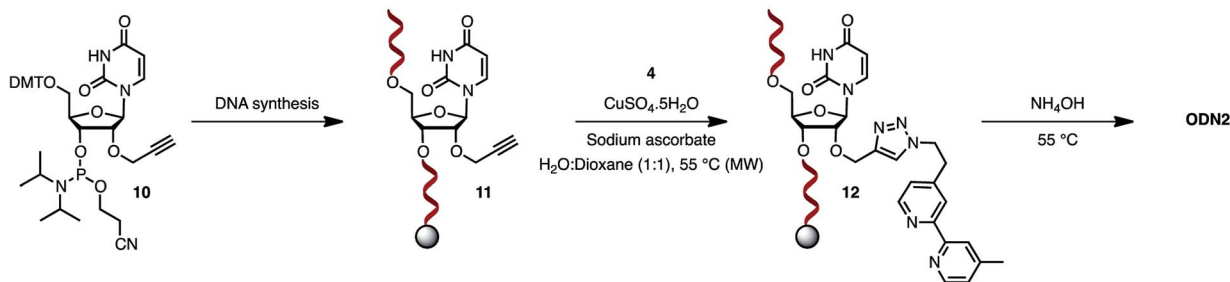
A Synthesis of the bipyridine adducts



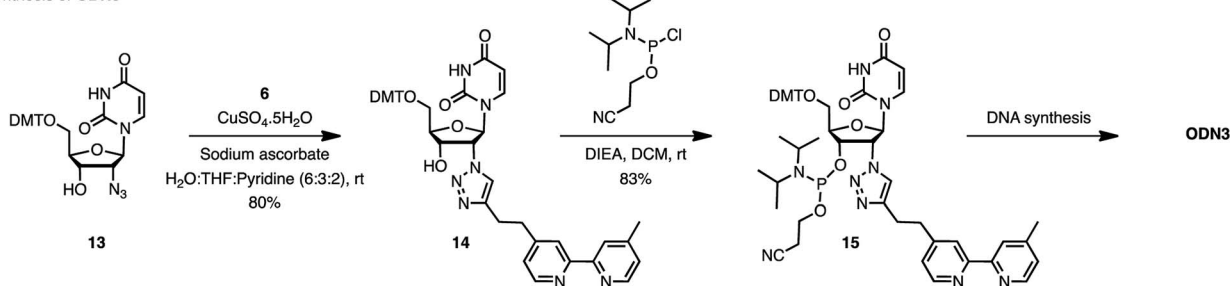
B Synthesis of ODN1



C Synthesis of ODN2



D Synthesis of ODN3



Scheme 2 Synthesis of oligonucleotides ODN1, ODN2 and ODN3.

was also evaluated for comparison purposes. A large variety of indoles differing in their substitution pattern were assessed (Table 1). Three major trends could be identified from these reactions: (1) good to excellent conversions were observed albeit lower than the ones obtained using the supramolecular approach,^{4c} (2) the non-covalent approach using st-DNA or ODN5/ODN4 in the presence of dmbipy afforded very similar ee values, and (3) the best selectivities were obtained with the covalently modified sequences, particularly ODN2/ODN4, which clearly outperformed the supramolecular approach in terms of

enantioselectivity. In contrast, ODN1/ODN4 (major groove) and ODN3/ODN4 (minor groove) could not clearly differentiate between the two faces of the pro-chiral enolate, thus suggesting that the positioning of the ligand in either groove did not seem to have a significant effect on the selectivity of the reactions. Moreover, the catalytic efficacy of our lead duplex appeared to be far less dependent on the nature of the indole than st-DNA as showcased by the good to excellent ee values obtained with the various indoles tested. Finally, an interesting inversion of selectivity was observed when using ODN2/ODN4 compared to



Table 1 Scope of the a tandem Friedel–Crafts conjugate addition/asymmetric protonation reaction^{a,b}

| 16 | + | 17a-o |
|---|---|--|
| st-DNA or ODNx/ODN4 Cu(NO ₃) ₂ (30 mol%) MES (pH 5.0), 4 °C | | |
| 18a-o | | |
| 18a | 18b | 18c |
| st-DNA 95% +21% ee ODN5/ODN4 87% +40% ee ODN1/ODN4 30% -20% ee ODN2/ODN4 76% -80% ee ODN3/ODN4 30% +7% ee | st-DNA 96% +17% ee ODN5/ODN4 80% +38% ee ODN1/ODN4 52% -12% ee ODN2/ODN4 78% -86% ee ODN3/ODN4 32% -29% ee | st-DNA 88% +29% ee ODN5/ODN4 65% +5% ee ODN1/ODN4 44% -16% ee ODN2/ODN4 64% -86% ee ODN3/ODN4 21% -19% ee |
| 18d | 18e | 18f |
| st-DNA 83% +43% ee ODN5/ODN4 99% +41% ee ODN1/ODN4 78% -6% ee ODN2/ODN4 85% -74% ee ODN3/ODN4 65% 0% ee | st-DNA 92% +11% ee ODN5/ODN4 92% +37% ee ODN1/ODN4 25% -14% ee ODN2/ODN4 88% -82% ee ODN3/ODN4 26% +8% ee | st-DNA 95% +36% ee ODN5/ODN4 99% +36% ee ODN1/ODN4 44% -14% ee ODN2/ODN4 99% -73% ee ODN3/ODN4 42% +13% ee |
| 18g | 18h | 18i |
| st-DNA 99% +42% ee ODN5/ODN4 90% +43% ee ODN1/ODN4 46% -4% ee ODN2/ODN4 81% -67% ee ODN3/ODN4 50% +20% ee | st-DNA 83% +37% ee ODN5/ODN4 >99% +29% ee ODN1/ODN4 71% -20% ee ODN2/ODN4 88% -70% ee ODN3/ODN4 53% -15% ee | st-DNA 87% +27% ee ODN5/ODN4 68% +42% ee ODN1/ODN4 15% -2% ee ODN2/ODN4 39% -80% ee ODN3/ODN4 13% +8% ee |
| 18j | 18k | 18l |
| st-DNA 69% +19% ee ODN5/ODN4 47% +4% ee ODN1/ODN4 11% +27% ee ODN2/ODN4 32% -63% ee ODN3/ODN4 5% +7% ee | st-DNA 70% +16% ee ODN5/ODN4 37% +20% ee ODN1/ODN4 10% +11% ee ODN2/ODN4 18% -54% ee ODN3/ODN4 4% +19% ee | st-DNA 43% +23% ee ODN5/ODN4 24% +24% ee ODN1/ODN4 <5% nd ODN2/ODN4 9% -49% ee ODN3/ODN4 <5% nd |
| 18m | 18n | 18o |
| st-DNA 95% +72% ee ODN5/ODN4 84% +72% ee ODN1/ODN4 <5% nd ODN2/ODN4 47% -67% ee ODN3/ODN4 <5% +4% ee | st-DNA 76% +79% ee ODN5/ODN4 77% +71% ee ODN1/ODN4 6% +8% ee ODN2/ODN4 50% -65% ee ODN3/ODN4 13% +8% ee | st-DNA >99% +3% ee ODN5/ODN4 >99% -36% ee ODN1/ODN4 <5% nd ODN2/ODN4 >99% +61% ee ODN3/ODN4 <5% nd |

^a Conditions with st-DNA: st-DNA [2 mM in Milli-Q H₂O (29 μL)], 200 mM MES buffer solution (10 μL, pH 5.0), 1.0 mM of Cu(NO₃)₂-dmbipy in Milli-Q H₂O solution (33 μL, 30 mol%), 0.05 M solution of freshly prepared enone in DMSO (2.0 μL, 1 equiv.), 0.05 M solution of indole in DMSO (2.0 μL, 1 equiv.), 3 d, and 4 °C. Conditions with ODNx/ODN4: ODNx/ODN4 (40 mol%), 200 mM MES buffer solution (10 μL, pH 5.0), 1.0 mM of Cu(NO₃)₂ in Milli-Q H₂O solution (30 μL, 30 mol%), 0.05 M solution of freshly prepared enone in DMSO (2.0 μL, 1 equiv.), 0.05 M solution of indole in DMSO (2.0 μL, 1 equiv.), 3 d, and 4 °C. ^b Conversion and ee values were determined by High Pressure Liquid Chromatography (HPLC) analysis. ODNx/ODN4: 5'-GCCAGCXGACCG-3'/5'-CGGTCAGCTGGC-3'.

any of the non-covalent approaches (ODN5/ODN4 or st-DNA) and this trend was confirmed for pretty much all the indoles tested independently of their substitution pattern. Hence, the use of 1*H*-indole led to the corresponding saturated ketone in -80% ee with ODN2/ODN4 while only +21% ee was obtained with st-DNA (Table 1, **18a**). Likewise, the 5-, 6- and 7-methoxy-substituted indoles afforded ee values ranging from 74 to 86% (Table 1, **18b–d**) with ODN2/ODN4 while the same reactions run in the presence of st-DNA afforded only moderate selectivities (up to +43% ee). A similar trend was also observed with the 5-, 6- and 7-methyl-substituted indoles as well as the 5-hydroxy indole, which afforded ee values ranging between -67 and -82%, while the enantioselectivities did not exceed +43% ee when using the supramolecular approach (Table 1, **18e–h**).

ODN2/ODN4 also proved to be superior to st-DNA when indoles bearing an electron-withdrawing group at the 5 position were used. This was the case with 5-fluoro indole (-80% ee vs. +27% ee, **18i**), 5-chloro indole (-63% ee vs. +19% ee, **18j**), 5-bromo indole (-54% ee vs. +16% ee, **18k**) and methyl indole-5-carboxylate (-49% ee vs. +23% ee, **18l**). ODN2/ODN4 displayed a similar selectivity to st-DNA when indoles bearing a tertiary amine at the 5 position such as a piperidine (-67% ee vs. +72% ee, **18m**) or a morpholine (-65% ee vs. +79% ee, **18n**) were used. Protonation of the amine in the reaction media was advanced to explain the higher ee values obtained with st-DNA;^{4c} however in view of our results, this does not seem to be the case in our covalent approach. Finally, the use of *N*-methyl indole led to the corresponding product in only +3% ee with st-



DNA, while **ODN2/ODN4** afforded up to +61% ee under otherwise identical conditions (Table 1, **18o**). It is worth pointing out as well that this was actually the only example that did not lead to an inversion of the selectivity thus confirming the importance of specific interactions between the substrate and the oligonucleotides.

Discussion

The catalytic performance of **ODN2/ODN4** compared to either **ODN1/ODN4** or **ODN3/ODN4** or the non-covalent approach is quite remarkable and leads to unprecedented levels of enantioselectivities for tandem Friedel–Crafts conjugate addition/asymmetric protonation in water. In order to rationalize the differences observed, especially when the bipyridine ligand is positioned in the major groove or in the minor groove through a 2'-triazolyl linker, we recorded circular dichroism (CD) spectra of the different modified duplexes in the absence and presence of Cu(II) (Fig. 1). For all three duplexes, the CD spectra were very similar and exhibited a typical B-type CD spectrum characterized by a negative band around 240 nm and a positive band around 260 nm. The attachment of the dmbipy ligand through a triazolyl linker either at the 5-position of the thymidine or at the 2'-position of the 2'-deoxyuridine did not seem to induce any significant change in the spectrum compared to that of the non-modified duplex **ODN4/ODN5**. Similarly, addition of Cu(II) ions did not induce substantial ICD (Fig. 1). According to the CD spectra, these observations indicate that the covalent attachment of the bipyridine ligand has no major influence on the folding of these duplexes. **ODN2/ODN4** is the only duplex that shows a small decrease in the intensity of the positive band after addition of Cu(II) ions thus revealing a small change in the interaction of the ligand with the DNA helix (Fig. 1b). This small structural change may be the key factor that accounts for the higher enantioselectivities observed with **ODN2/ODN4** compared to all the other modified duplexes. To further investigate these systems, the influence of the covalent modification was analysed by thermal denaturation studies.

A single sigmoidal transition was obtained in all cases. Compared with the non-modified duplex **ODN4/ODN5**, the presence of the bipyridine linker destabilised the duplex ranging from $\Delta T_m = -5^\circ\text{C}$ for **ODN1/ODN4** and **ODN2/ODN4** to $\Delta T_m = -11^\circ\text{C}$ for **ODN3/ODN4**. This strong destabilization

might also affect the catalytic efficiency observed with **ODN3/ODN4**. Interestingly, no significant effect was observed upon addition of copper(II) ions (Table 2). The steric restrictions imposed by the ligand and the substrate/DNA-binding interactions might be the main parameter affecting the stereochemical outcome of the reaction (Table 1, **18g–j**, comparing **ODN2/ODN4** with **ODN3/ODN4**). In terms of the catalytic activity, the only difference between **ODN2/ODN4** and **ODN3/ODN4** is the attachment of the triazolyl linker. Interestingly however, we observed no influence of the groove on the selectivity outcome as only one duplex which happened to position the ligand in the minor groove gave successful results. The structural difference was related to the attachment of the triazolyl linker on **ODN2/ODN4**, which was longer and more flexible. Structural changes caused by modification of the sugar at the 2'-position have been extensively investigated. In particular, it was shown that 2'-*O*-modifications, like the ones found in **ODN2**, deviate the conformational equilibrium of the sugar toward the C3'-*endo* (North) pucker in order to prevent steric clash between the phosphate backbone and the neighboring nucleobases.^{19–22} This puckering generates locally an A-form geometry into the helix, causing the distance between two adjacent bases to be reduced, a characteristic of RNA duplexes. Interestingly, 2'-azido groups have also been shown to induce a 3'-*endo* sugar puckering conformation.²³ However, examination of these modifications revealed that the 2'-azido group mainly interacts with the adjacent 3'-phosphate group,²³ whereas in the case of 2'-*O*-modifications, the 2'- and 3'-oxygen atoms as well as the 2'-

Table 2 Melting temperatures (T_m) of covalently modified duplexes in the absence (–) and presence (+) of Cu(II) ions

| Entry | Duplex | Cu(NO ₃) ₂ | T_m^a (°C) |
|-------|------------------|-----------------------------------|--------------|
| 1 | ODN4/ODN5 | – | 56.9 |
| 2 | ODN1/ODN4 | – | 52.1 |
| 3 | ODN1/ODN4 | + | 53.2 |
| 4 | ODN2/ODN4 | – | 52.1 |
| 5 | ODN2/ODN4 | + | 49.8 |
| 6 | ODN3/ODN4 | – | 46.1 |
| 7 | ODN3/ODN4 | + | 44.9 |

^a Melting temperatures are obtained from the maxima of the first derivatives of the melting curve (A260 vs. temperature) recorded in a buffer containing 1 M NaCl and 10 mM sodium cacodylate. Curve fits data were averaged from fits of three denaturation curves.

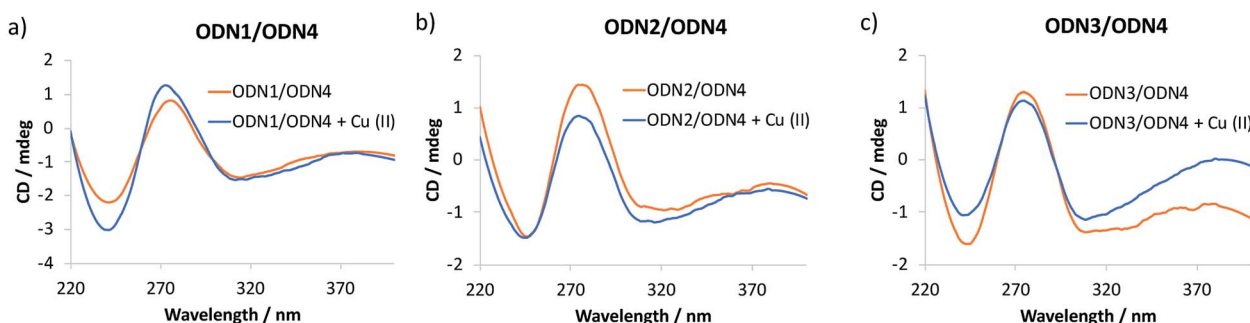
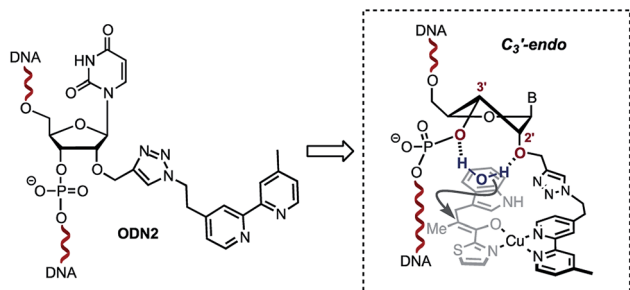


Fig. 1 CD spectra of (a) **ODN1/ODN4**, (b) **ODN2/ODN4** and (c) **ODN3/ODN4** duplexes folding in the absence and in the presence of Cu(II) ions.





Scheme 3 Plausible sugar pucker conformation in ODN2. The 2'-modification is likely to favour the C3'-endo conformation positioning a water molecule between the 2'-O and the 3'-phosphate.

O-substituents provide a stable cavity which can potentially coordinate to a molecule of water.^{22,24} We believe that this structural feature combined with the local A-form geometry adopted by the duplex accounts for the high enantioselectivities observed with ODN2/ODN4. Indeed, the ability of the 2' and 3' oxygen atoms in the 2'-O-substituted RNA residues to coordinate to a molecule of water, which was demonstrated by Egli and co-workers,^{22,24} prompted us to propose the hypothesis that the configuration adopted by the ODN2/ODN4 duplex is likely to sequester a molecule of H₂O inside the cavity through H-bond interactions with the 2'-O present in ODN2 and the 3'-O of the 3'-phosphate group (Scheme 3). In contrast, this organized coordination of H₂O is most probably lacking in both ODN1/ODN4 and ODN3/ODN4. This plausible mechanism by which a molecule of water sequestered in an organized cavity can readily protonate a highly reactive pro-chiral enolate intermediate is likely to differ from the mechanism taking place in the non-covalent approach reported by Roelfes and co-workers.^{4c} This may actually explain why the selectivities obtained in our case are not dependent on the electronics of the indoles.

Conclusions

Through this study, we were able to devise a particularly effective bio-hybrid catalyst capable of achieving unprecedented levels of enantioselectivity in the challenging copper(II)-catalysed asymmetric Friedel-Crafts conjugate addition/enantioselective protonation of α -substituted α,β -unsaturated enones. Interestingly, our 2'-modified bipyridine-containing duplex ODN2/ODN4 proved not only to be highly effective but also highly versatile affording excellent levels of enantioselectivity with minimum substrate dependency thus setting a new benchmark in the field. Most importantly, by closely examining the structure of the duplex and that of the catalytic pocket, we suggested that the high selectivities could be induced by both the formation of a local A-type helix around the metallic co-factor and the sequestration of a molecule of water between the hydroxymethyltriazole arm and the 3'-phosphate which can readily protonate the highly reactive enolate intermediate. Finally, we believe that this study demonstrates the value of the covalent anchorage approach, which allows a fine tuning of the chiral micro-environment around the metallic co-

factor and thus allows designing highly selective catalytic systems for specific transformations.

Conflicts of interest

There are no conflicts to declare.

Acknowledgements

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