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Conjugated Oligonucleotides for Biochemical Applications

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Abstract

In the last 30 years oligonucleotides i.e., relatively short polymers (usually 12-24 units long) based on DNA structure, have found a widespread use in biochemical studies and as biochemical probes and chemotherapeutic agents for the downregulation of genetic expression or for exon skipping. Here we present a short review of studies from our laboratories on the synthesis and applications of different kind of conjugates to address some of these techniques. Preparation of conjugates with small alkyl groups, intercalators, fluorescent oligothiophenes, and lipophilic bile-acids will be discussed.

Keywords: Conjugated oligonucleotides; Antisense; Anti-gene; Molecular probes

Introduction

Natural oligonucleotides, i.e., short fragments of single stranded DNA have been actively studied after the discovery of the significance of the double stranded DNA in living beings. Their chemical synthesis was attempted since the '60s after the pioneering works of Todd [1] and Khorana [2,3], based on a tedious and difficult repetition of chemical steps aimed at the addition of single nucleosides employing the methodology of active esters of P(V) [4,5]. Despite the difficulties of this approach, short sequences of DNA with a known sequence were employed for structural characterization X-ray crystallography and NMR [6,7], studies of drugs-binding (distamycins [8], daunomycins [9] etc.), and as antisense gene down-regulators [10]. In the '80s, following the development of the intuition of Beaucage and Carruthers [11], a new strategy of synthesis based on the high reactivity of P(III) was developed [12] and found a widespread application toward the automated synthesis starting from the 3'-terminal nucleoside attached to a solid support normally controlled pore glass (CPG). Since then little has changed, but as chemists, we were challenged with the poor performance of exogenous DNA. In fact, outside of the strict cellular control, exogenous oligonucleotides encounter difficulties to cross the cell membranes, and once inside cells are rapidly degraded by nucleases before reaching the desired target. A first response to the fast degradation challenge was the development of oligonucleotides with modified backbones (phosphorothioates, phosphoridithioates, etc.), modified sugars (arabino, 2'-O-Alkyl-RNA) or both (morpholino, PNA etc.). In this rush, we discovered, despite numerous successful applications, the uniqueness of natural sequences, for the fine regulation of biochemical processes; in fact modified oligo's simply have too many chiral centers, the wrong solubility, and a too high or too low affinity toward their complementary sequences. A more sensible approach for dealing with such precious molecules is considering oligonucleotides as strings of "biochemical information" and try to perturb the natural structure with the "minimal efficient modification" that sometime, can be obtained with the chemical conjugation of small molecules.

Antisense Stability

The antisense methodology is a way to impair the genetic expression of an unwanted gene interfering with its translation forming a duplex with the sense RNA using an exogenous oligonucleotide (usually of DNA type). There are many good reviews on this methodology [13,14] that can be carried out with several variations. In its basic form, the "antisense" oligonucleotide pairs according to Watson and Crick to the "sense" mRNA. This hetero-duplex can be degraded by RNase-H preventing the synthesis of the corresponding protein. Several factors complicate this nice picture; one of them is the degradation of the "antisense" especially by exonucleases, before it reaches its target. To overcome this problem, many researchers proposed the use of phosphorothioated oligonucleotides that have a good resistance against nuclease degradation. However, phosphorothioates have two serious drawbacks: they introduce one chiral phosphorous per substitution, so that an 18-mer PS oligo is a complex mixture of 217 different diasteroisomers; secondly they tend to bind a-specifically to cellular proteins, causing unwanted side effects [15,16]. A more sensible approach seems that of using terminal modification to the antisense oligonucleotide. One of the first attempts in this direction was that of Zamecnik who in 1978 published a work on an oligonucleotide to be used as antisense agents against the Ros sarcoma virus with both ends protected as phenylcarbamoyl derivatives [10]. According to the procedure of Agarwal [17] the protected tridecamer prepared with the phosphotriester method [4] was reacted, in its protected form but with 5' and 3' free hydroxyl groups, with phenylisocyanate to give after deprotection the end capped phenylcarbamoyl-oligonucleotide. The efficiency of the protection can be tested against the snake venom phosphodiesterase (SVPDE) a 3'-exonuclease or calf-spleen phosphodiesterase (CSPDE) the corresponding 5' enzyme.

Following the Zamecknic's example, we prepared a set of oligonucleotides of identical sequence, but having the extremities conjugated with propandiol, hexandiol, or *seco*-uridine (Figure 1) and compared them with phosphorothioated analogs toward resistance against 3'- and 5'- exonucleases. The efficiency of the conjugation was tested by HPLC against SVPDE and CSPDE finding an increase resistance toward both degradations of all the end-modified oligonucleotides, with a more than ten times increased resistance of the *seco*-uridine conjugates respect to the natural unproteted oligonucleotide [18].

The 5'-end *seco*-uridine modification, and similarly the propandiol and hexandiol derivatives, were obtained preparing the corresponding

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phosphoramidites, suitable for the automated synthesis, while for the 3'-end modification we had to derivatize the used solid support reacting the aminoalkyl-CPG with the dimethoxytrityl protected uridine with succinic anhydride and a condenser. The logic behind the preparation of those compounds is sketched in Scheme 1. Briefly: uridine 1 was protected selectively at the 5' position with 4,4'-dimethoxytrityl chloride in pyridine to give 2 that was oxidized to the dialdehyde 3 with sodium periodate in water/acetone; 3 was reduced with sodium borohydride in ethanol to 4 that was converted to the monobenzoyl ester mixture 5+6 with one equivalent of benzoyl chloride in pyridine. The mixture was separated on silica gel and compound 5 was used for the preparation of amidite 7 by use of 2-cyanoethyl-N,N-diisopropylchlorophosphorami dite in dichloromethane and 5 eq. of diisopropylethylamine (DIPEA). The mixture 5+6 was used to prepare the succinic ester by reaction with the corresponding anhydride, and dicyclohexylcarbodiimide (DCC) as a condenser, in pyridine, then the crude filtrate was reacted with long chain amine-CPG (n=6) or amine-CPG (n=3) by gentle shaking in a sealed vial to the modified CPG 8, finally capped with acetic anhydride.

Exon Skipping

A variant of the antisense approach is that of exon skipping. Antisense-mediated exon skipping [19] is a promising approach for the treatment of Duchenne Muscular Dystrophy DMD, a rare genetic disease due to a mutation of the dystrophin gene that cause the production of a truncated not functional protein. Targeting the antisense oligonucleotide to a selected region of the pre-mRNA, it is possible to modify the normal introne-hexone reading frame in a way to exclude the sequence that during the translation codes for a stop signal, in favor of a shorter but still partly functional variant of dystrophin. For this application, that do not require the intervention of RNase-H, 2'OMe-RNA oligonucleotides are normally used, in the form of phosphorothioates (PS). Among other factors, the efficient delivery of the oligonucleotide remains a significant challenge for its use as therapeutic drugs, and we demonstrated that a possible solution was to convey the oligonucleotides across the cell membrane through the use of cationic core-shell nanoparticles [20,21].

A new possibility raised some years ago when Regen and coll. conjugated a 5'-thiol modified T18 with two molecules of bile

acids joined by a flexible scaffold [22,23]. They found that this assembly allowed the bile acids hydrophilic faces to wrap around the oligonucleotide presenting the opposite lipophilic surfaces toward an artificial liposomal membrane, simulating a cellular one helping the conjugated oligonucleotide to cross the bilayer membrane, with the ingenious mechanism (dubbed molecular umbrella) depitched in Figure 2.

Mimicking this approach, we are now testing a 2'OMe-RNA-PS oligonucleotide targeting exon 23 in mdx mouse, provided with a tri-valent bile acid scaffold (Scheme 2). To this aim, a post-synthetic conjugation approach was assessed, i.e., during the solid phase synthesis, the oligonucleotide was equipped at the 5'-end with the appropriate functionality to which the lipophilic moiety was coupled in solution after release and purification of the antisense. Firstly, we have prepared the hydrophobic moiety. Thus, the known 3a-N₂-ursodeoxycholic methyl ester (3aN₃UDC) [24] and the tris-propargyloxyamino derivative 9 obtained from TRIS according to a literature procedure [25] were chosen as building blocks for its synthesis. For assembling the desired tri-valent bile acid scaffold 10 we used the copper-catalyzed azide-alkyne cycloaddition (CuAAC) under microwave irradiations. In these conditions we isolated 10 in 24% overall yield after flash chromatography. Compound 10, was subjected to N-succinylation with succinic anhydride, and then transformed into the reactive ester 11 by using N-hydroxysuccinimide (NHS) in presence of DCC, with an overall yield of 83%. Finally, the conjugation of the 5'-amino-modified antisense (5'-NH2-AON) with the lipophilic N-hydroxysuccinimide ester 11 was accomplished in solution, by carrying out the reaction in bicarbonate buffer at pH 8.2 for 20 hours to achieve the desired lipophilic oligonucleotide 12 with 30% yield after reverse phase purification. The ability of the new synthesized lipophilic oligonucleotide 12 to cross the phospholipid membrane is currently underway in our laboratories.

Anti-gene Studies with Oligonucleotide-Daunomycin Conjugates

The anti-gene approach, i.e., the use of an oligonucleotide to interfere with the transcription of the DNA and ultimately with the expression of the encoded protein, arises from the discovery of the possibility of binding an oligonucleotide to some regions of a regular duplex. In 1957,



in fact, Felsenfeld found evidences of the existence of a three stranded structure of composition 2PolyU:polyA [26]; several years later Lyamichev published the "Structures of homopurine-homopyrimidine tract in superhelical DNA" [27] then called H-DNA. When it was clear that the resulting "triplex" complex was formed by an oligonucleotide located into the major groove of the DNA, binding the homopurinic strand, through Hoogsteen or reverse Hoogsteen hydrogen bond patterns, the gene itself became a target for several kind of biochemical applications. Pioneers in this field were Dervan [28] and Hélène [29]. At difference with what happen by using oligonucleotides as antisense

agents, in the triplex approach the oligonucleotide is designed to bind directly to the gene of interest, inside the nucleus of eukaryotic cells. The theoretical tremendous advantage of such a technique over the antisense methodology is evident from the count of the target molecules, in principle a single molecule could shut down a gene in a single cell. Moreover it was demonstrated that the approach is feasible despite the presence of chromatine [30]. The success of this methodology depends among other factors (cellular uptake, biodistribution, stability etc.) on the strength of binding of the triplex forming oligonucleotide (TFO) to its target, and several groups used to conjugate the TFO with DNA

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Figure 2: Scheme of synthesis and representation of the crossing of a bilayer phospholipid membrane aided by the "umbrella effect". Reprinted adapted with permission from: [22,23].



intercalators to enhance the stability of the triplex aggregate [31]. We decided to follow this strategy employing daunomycin. Daunomycin is a powerful antitumor molecule that interact with DNA, intercalating its anthraquinone planar moiety between two couples of bases (preferentially GpC) laying the aminosugar into the minor groove [32]. Following this intercalation, its natural fluorescence is quenched, (this can be monitored to demonstrate the binding), and the complex DNA-daunomycin prevents the topoisomerase II resealing after the cleavage of one of the strands required by the unwinding, eventually leading to cellular death.

Some groups had already performed this conjugation binding the TFO with the aminogroup of the glycone of the daunomycin [33], but with little gain in terms of triplex stabilization and biological effect. We thought that better results could be achieved binding the daunomycin to the TFO at the opposite end, to preserve the natural way of intercalation of daunomycin with DNA, since the TFO binds the homopurinic strand displaying itself into the major groove of the duplex. Following this idea, we made a partnership with researchers of Menarini Laboratories, who helped us to modify the daunomycin. Ultimately we settled on a procedure based on the attachment of a iodohexametylene chain on the O-4 of the daunomycin and a post synthetic conjugation of the oligonucleotide via a chemoselective binding of a terminal thiophosphate group of the TFO [33] as depitched in Scheme 3 [34,35] the ⁵PS-TFO 14 was prepared using a 0.1 M acetonitrile solution of bis-2-cyanoethyl(N,N-diisopropyl)-phosphoramidite [36] with a longer coupling time on the automated synthesizer to ensure the complete coupling, followed by thio-oxidation, we get the PSunprotected oligo after usual ammonia deblocking. The ³¹P-NMR allowed us to judge if the oligonucleotide had the right PS/PO ratio. The conjugation with the protected daunomycin 13 was performed in water/DMF. After water/dichloromethane extraction only the conjugate 15 remains in water. The protective groups on the amino sugar were removed by reaction with an aqueous diluted soda solution; finally, the unprotected oligo 16 was purified by preparative reversed phase HPLC. The progress of the reaction could easily be monitored by HPLC, moreover the ³¹P-NMR signal of the phosphorothioate changed from 46 ppm of the monoester derivative 14 to 23 ppm of compound 16 (and 15). Later on we were able to prepare daunomicyn conjugates also in the 3' position (and in both 5' and 3' position) preparing a modified CPG according to a published procedure [37]. We utilized this modified CPG to prepare 3'-PS oligonucleotides [38] and from this a 3'-conjugate. Daunomiycin-TFO conjugates were applied with some interesting results against the poly purine tract of HIV-1 and against the promoter P2 of c-myc oncogene [35]. In the end, the efforts spent in this research leaded, as hoped, to a demonstration of an active role of the conjugated daunomycin after intercalation, mediated by topoisomerase. Indeed, targeting the same site MDR1 in resistant cells lines (MCF7-R and NIH-3T3) with a 5' or a 3' daunomycin-TFO conjugate, we observed the higher biological response with the less stable of the pair [39].

Fluorescent Probes

We had demonstrated for the first time the compatibility of oligothiophenes as fluorescent markers for oligonucleotides Scheme 4 [40].

The ready available terthiophene alcohol **17** can be reacted with the 2-cyanoethyl-*N*,*N*-diisopropylchlorophosphoramidite to give the corresponding amidite **18** that was found suitable for the conventional solid phase synthesis of oligonucleotides. We prepared the showed tetrathymidine conjugate **19** characterized by NMR, UV and fluorescence. Unlikely the conventional "old-fashionable" fluorescein labeled conjugates, these derivatives are stable under UV light without showing any bleaching.

The developments of those fluorescent labels made possible the design of biological probes in the form of molecular beacons (MBs). MBs are oligonucleotides bearing a fluorophore at one end and a quencher on the other. They are designed with short self-complementary regions flanking a central recognition sequence. In absence of a target the flanking regions pair, putting the fluorophore and the quencher in close proximity, in a conformation that graphically resemble to a hairpin; in that form the fluoresce is reduced by fluorescence resonance energy transfer (FRET). Conversely in presence of a target complementary to the central sequence, the MB forms a duplex with the target, the quencher is distanced from the fluorophore (more than the Förster distance) and the fluorescence is restored [41] (Figure 3).

We tried to prepare MBs employing oligothiophenes, thus we synthesized a 5'-dabcyl conjugate oligonucleotide with a free 3'-amino group. From this, we prepared several different oligothiophene derivatives using the corresponding succinimide-oligothiophene (Figure 4 left panel). Surprisingly we found that to detect the complementary target the presence of a quencher was not strictly necessary [42,43] (Figure 4 right panel), probably because this kind of







Figure 3: Sketch of the functioning of a molecular beacon (MB). The hairpin structure puts the fluorophore and the quencher in close proximity so that the fluorescence is strongly reduced. Fluorescence is restored after thermal denaturation or hybridization of the MB with a complementary target.



Figure 4: Left panel: sequences, methodology of synthesis and structure of the fluorophores. Right panel: emitted fluorescence of the corresponding hairpin or duplex with or without the dabcyl quencher.

fluorescent tags were able to exchange energy by resonance with the guanines at the opposite ends.

Oligothiophene fluorophores can be electronically coupled to a uridine base to make it sensible to the correct base pairing, then used to detect single nucleotide mutations [43,44] (Figure 5).

In this case, the uridine derivatives (**20-23**) can be seen as conjugate nucleosides having an oligothiophene joined to the uridine base with a conductive linker. The formation of a hydrogen bonding following the correct pairing (U:dA) modifies the molecular orbitals of the uridine and of the conjugate label, resulting in a different fluorescence. We found that the phosphoramidite of the 5'-dimethoytrityl modified

uridines (20-23) were suitable for the automated synthesis of oligonucleotides and were not affected by the following deblocking procedures. The resulting oligonucleotides containing the modified uridine at the middle of the sequence, showed appreciable fluorescence variations when faced with an oligonucleotide containing the correct or a mismatched base (Figure 5 right panel).

Compounds **20-23** were prepared via Sonogashira's coupling (**20**) or reverse coupling (**21**) or from Stille's coupling reaction (**22**, **23**) starting from 5'-dimethoxytrityl-5-iodouridine **24** or 5'-dimethoxytrityl-5etinyl-uridine (**26**) according Scheme 5. The amidites were prepared by reaction of the corresponding precursors with 2-cyanoethyl-*N*,*N*-



Figure 5: Left panel: modified uridines used to detect single nucleotide polymorphisms. Right panel: hybridization experiments; the probe sequence containing the modified uridine in the middle, showed a different fluorescence pattern when hybridized with a complementary strand containing a deoxy-adenosine, or any other base in the facing position.



(yields: 17a 97%, 17b 40%).

diisopropylchlorophosphoramidite in dichloromethane and DIPEA (in case of compound **25**) or with the bis-*N*,*N*-diisopropyl(2-*O*-cyanoethyl)phosphite in dichloromethane/acetonitrile with one eq. of tetrazole (compounds **27** and **28a** and **b**), in these cases isolation of the derivatives was not necessary.

Adenosine Derivatives

The possibility to bind a functional molecule to a nucleoside, as demonstrated above, offer the chance of precisely locating that molecule along an oligonucleotide, therefore enhancing the possibility of tuning a possible biochemical effect. Looking for a convenient way to use oligonucleotide conjugated with psoralene having the paper of Sproat [45] as a starting model, we settled on a versatile synthesis based on the modification of the C-8 position of deoxyadenosine. Psoralene is a planar molecule able to intercalate preferentially among TA base pairs to give, after UV irradiation, a reversible [2+2] cycloadduct with the C5-C6 atoms of the thymidine rings on the two facing strands. This molecule had been conjugated to antisense DNA since the beginning of this methodology [46,47]. Our original methodology was based on the efficient C8 alkylation of the adenine ring by a sulfur atom at the end of a bifunctional dithiol alkyl linker (Scheme 6). The 8-Brdeoxyadenosine 29 was reacted with an alkyl dithiol in hot water and few equivalents of TEA to give, after extraction with ethyl acetate, the derivative 30. This was readily converted to the psoralen nucleoside 31, then, through reactions c-e, it was 5' protected with DMTrCl in pyridine to give compound 32, converted to 33 with diphenylacetyl chloride, and eventually to the corresponding amidite 34 by reaction with 2-cyanoethyl-N,N-diisopropylchlorophosphoramidite in dichloromethane and TEA. The psoralen nucleoside was finally inserted in the middle of a sequence by conventional solid phase synthesis.

The conjugated oligonucleotide containing the psoraleneadenosine nucleoside **31** was demonstrated to be able to bind to its complementary sequences and to do cross-linking upon UV irradiation by HPLC experiments (Figure 6) [48].

Utilizing the same chemistry, we were able to prepare a conjugate with a different intercalator, i.e., acridine (Scheme 7).

In this case the 2,9-dichloro-6-mehoxyacridine was reacted with 8-dithiopentenyldeoxyadenosine (compound **30** from the previous scheme), in DMF and TEA at 80°C, for 17 h to give **35** (70% yield). With the same reaction pattern **c-e** already seen this compound was tritylated to **36**, protected with diphenylacetyl to **37** and eventually converted to the phosphoramidite **38** in analogy with the synthesis of the psoralene phosphoramidite **34** with similar yields. The oligonucleotide conjugated with this fluorescent molecule was found to have a reversible binding toward gold nanoparticles, possibly opening the way for a fast purification methodology for every oligonucleotides bearing at least two different sulfur atoms [49].

One interesting way to prepare a large variety of oligonucleotide conjugates is based on the orthogonal chemoselectivity of the clickchemistry. This kind of reaction can be performed with two partners, one having an azido group and the other a terminal alkyne. The reaction can be carried on in an aqueous solution and is largely independent from the presence of many chemical groups, so it is ideal to further elaborate complicate (bio)molecules. Again, our modification at the C-8 of the 2'deoxyadenosine can be utilized to prepare oligonucleotides bearing an alkyne moiety as in Scheme 8.





Figure 6: Left panel: HPLC profile of the UV irradiated mixture of the oligonucleotide-psoralen conjugate with its complementary DNA strand at different times. Right panel: interpretation of the chromatographic peaks.





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Figure 7: Preparation of a multiple fluorescent oligonucleotide using the click chemistry with commercially available reagents.

The synthesis of the compound proceeded as following: the 8-Brdeoxyadenosine **29** was reacted with the hex-5-yne-1-thiol **39** in water and TEA at 100 °C, 2h, (30%). Steps **c-e** were performed as in previous schemes. At the best of our knowledge the amidite of this derivative is the only alternative to the C6-pyrimidine analogues commercially available [50]. Through the click-chemistry these oligonucleotides can be further processed to the desired compound, connecting the desired tag to an azido-tether [48] even with multiple groups (Figure 7) [43].

Conclusions

The conjugation of small molecules to nucleic acids allows affording several techniques with the best aptly derivatives. The only limits are the imagination of scientists and the feasibility of the involved chemical steps. We hope that the reported applications can be of inspiration to scholars facing biochemical challenges.

Conflicts of Interest

The authors declare no conflict of interest.

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