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Stability and structural features of the duplexes containing nucleoside analogues with a fixed N-type conformation, 2'-0,4'-C-methyleneribonucleosides

Satoshi Obika, Daishu Nanbu, Yoshiyuki Hari, Jun-ichi Andoh Ken-ichiro Morio, Takefumi Doi, and Takeshi Imanishi*

Graduate School of Pharmaceutical Sciences, Osaka University, 1-6 Yamadaoka, Suita, Osaka 565-0861, Japan.

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Abstract

Bicyclic nucleoside analogues with a fixed N-type conformation, 2'-0,4'-C-methyleneuridine and -cytidine, were incorporated into oligonucleotides, and the binding efficiency of the modified oligonucleotides to the complementary DNA and RNA as well as the CD spectra of the modified DNA-DNA and modified DNA-RNA duplexes were studied. © 1998 Elsevier Science Ltd. All rights reserved.

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Extensive efforts have been directed toward developing novel DNA and RNA analogues for the practical use of antisense and antigene technologies [1-5]. In these studies, several types of conformationally restricted nucleoside analogues were synthesized by our laboratory [6,7] and other groups [8]. In the previous paper [6], we reported the first synthesis of bicyclic nucleoside analogues with a fixed N-type conformation, 2'-O,4'-C-methyleneuridine **1a** and -cytidine **1b**.¹ It is well known that the duplex formation of the preorganized oligonucleotides (ONs) with complementary DNA and RNA are entropically favorable [11]. In addition, for the reason that the A-form RNA duplex possesses the N-type sugar conformation while the B-form DNA duplex possesses the S-conformation [12], the nucleoside analogues with a rigid N-type conformation, such as **1**, are readily expected to enhance the hybridization ability towards complementary RNA. Here, we wish to demonstrate the duplex stability of the ONs containing the conformationally fixed nucleoside analogues **1**, and the structural features of these duplexes.

As shown in Scheme 1, the phosphoramidite building block 3 was prepared from the corresponding 5'-O-dimethoxytrityl derivative 2 [6] by the usual method [13]. The modified ONs 5-13 as well as unmodified RNA and DNA strands 16-20 were synthesized by standard

^{*} e-mail: imanishi@phs.osaka-u.ac.jp

¹ Just after our publication of the synthesis of 1 [6], Wengel and co-workers reported the synthesis and some properties of the same compounds [9,10].



Scheme 1. (a) 2-cyanoethyl N,N,N',N'-tetraisopropylphosphordiamidite, diisopropylammonium tetrazolide, MeCN-THF (3:1), r.t., 1.5 h, 81%; (b) 1,2,4-triazole, POCl₃, Et₃N, MeCN, 0 °C, 4 h, 89%.

Table	1
1 44 10 10	

Tm Values (°C) of the modified ONs 5-12 towards complementary RNA 17 and DNA 18. *

Oligonucleotides		RNA Complement 17 5'-r (AGCAAAAAACGC)-3'	DNA Complement 18 5'-d (AGCAAAAAACGC)-3'
5'-d (GCGTTTTTTGCT)-3'	16	45	47
5'-d (GCGXTTTTTGCT)-3'	5	49 (+4)	50 (+3)
5'-d (GCGTTXTTTGCT)-3'	6	49 (+4)	49 (+2)
5'-d (GCGTTTXTTGCT)-3'	7	50 (+5)	49 (+2)
5'-d (GCGTTTTTXGCT)-3'	8	51 (+6)	52 (+5)
5'-d (GCGXXTTTTGCT)-3'	9	53 (+4)	51 (+2)
5'-d (GCGTTXXTTGCT)-3'	10	53 (+4)	50 (+1.5)
5'-d (GCGTTTTXXGCT)-3'	11	55 (+5)	54 (+3.5)
5'-d (GCGXXXXXXGCT)-3'	12	71 (+4.3)	58 (+1.8)

a. Duplex concentration: 4 μ M. Buffer: 100 mM NaCl,10 mM sodium phosphate buffer (pH 7.2); The values in parentheses are Δ Tm/modifications.

Table 2

Tm Values (°C) of the modified ONs 13-15 towards complementary DNA 20. a

Oligonucleotides	DNA Complement 20 5'-d (AAAAGGGAGAGAGA)-3'	
5'-d (TCTCTCTCCCTTTT)-3' 19	46	
5'-d (XCXCXCXCCCXXXT)-3' 13	54 (+1.1)	
5'-d (TYTYTYTYYYYTTTT)-3' 14	65 (+3.2)	
5'-d (XYXYXYXYYYXXXT)-3' 15	74 (+2.2)	



X: Base = uracil-1-yl Y: Base = cytosin-1-yl

a. See footnote in Table 1.

phosphoramidite protocol on the DNA synthesizer.² In addition, the modified ONs 14 and 15, in which 2'-0,4'-C-methylenecytidine 1b was introduced, were prepared by the postelongation-modification procedure [14]. The phosphoramidite 3 was treated with triazole, POCl₃ and Et₃N to afford the triazolouridine derivative 4 in good yield, which can be converted into the cytidine derivative by treatment with conc. ammonia after ON synthesis.²

The binding efficiency of the modified ONs to the complementary sequences was assessed by an analysis of the UV melting curve. The melting temperatures (Tm) for the modified ONs 5-12 and 13-15 are summarized in Tables 1 and 2, respectively. Incorporation of the nucleoside analogues 1 into ONs significantly enhances hybridization ability towards the complementary strand. Especially, in the case of the modified ONs 5-12, the remarkable thermal stability of the duplexes with complementary RNA 17 was observed (Δ Tm/modifications towards 17 = ca. +5 °C, Δ Tm/modifications towards 18 = ca. +3 °C), which was derived from the rigid N-form structure of nucleoside analogues 1.

In order to investigate a thermodynamic contribution of the conformationally restricted nucleoside analogue to duplex stability, the parameters (ΔH° , ΔS° and $\Delta G^{\circ}_{37^{\circ}C}$) for the duplexes **16**•17 and **12**•17 were determined by van't Hoff plots [15]. The results are shown in Table 3. The free energy ($\Delta G^{\circ}_{37^{\circ}C}$) of the modified duplex **12**•17 is increased when compared with that of the unmodified DNA-RNA duplex **16**•17. Furthermore, it is noteworthy that the modified duplex **12**•17 showed a quite favorable ΔS° , relative to the unmodified duplex. From these results, the difference in binding ability towards complementary RNA between the modified ON **12** and the unmodified one **16** ($\Delta Tm = +26$ °C) translates to a difference in binding entropy ($\Delta \Delta S^{\circ}$) of *ca*. 28 cal•mol⁻¹•K⁻¹.

Table 3

Thermodynamic parameters for duplex formation between modified ON 12 and Complementary RNA 17. a

Duplexes	ΔH° (kcal \cdot mol ⁻¹)	ΔS° (cal · mol ⁻¹ · K ⁻¹)	$\Delta G^{\circ}_{37^{\circ}C}$ (kcal · mol ⁻¹)
16 • 17	-98.38	-282.0	-10.95
12 • 17	-96.98	-254.2	-18.18

a. Thermodynamic parameters were determined in 10 mM sodium phosphate buffer (pH 7.2) with 100 mM NaCl. Duplex concentrations ranged from 0.5 to 6.0 μ M.

CD spectra of the modified (12•18) and unmodified DNA-DNA duplexes 16•18 were measured as well as those of the modified (12•17) and unmodified DNA-RNA duplexes 16•17, which indicated the structural features of the modified duplexes (Fig. 1). The CD spectrum of the modified DNA-DNA duplex 12•18 had a relatively strong positive band at 271 nm and a relatively weak negative band at 245 nm, while the spectrum of the unmodified duplex 16•18 had roughly equal intensity of positive and negative bands at 283 and 250 nm, respectively. Similar tendency was observed in the spectrum of the modified DNA-RNA duplex 12•17. The strong positive band for the modified duplex 12•17 was shifted to shorter wave lengths, and the

² The modified ONs were synthesized on the DNA synthesizer (Gene Assembler[®] Plus, Pharmacia, 0.2 μmol scale, 5'-dimethoxytrityl on). After treatment with conc. ammonia, removal of the 5'-dimethoxytrytyl group and purification were performed on NENSORBTM PREP reversed-phase columns. The purity of the modified ONs was verified using reversed-phase HPLC and the compositions were determined by MALDI-MS.

negative band around 240 nm was reduced in comparison with these bands in the spectra of unmodified duplex 16.17. These observations supported that the incorporation of the nucleoside analogue 1a into ONs transform the duplexes into A-like conformation.



Fig. 1. CD Spectra of the modified and unmodified duplexes. A, modified DNA-DNA duplex 12•18 (solid line); unmodified duplex 16•18 (dashed line). B, modified DNA-RNA duplex 12•17 (solid line); unmodified duplex 16•17 (dashed line). Duplex concentration: 4 μ M. Buffer: 100 mM NaCl, 10 mM sodium phosphate buffer (pH 7.2): Temp 18 °C.

We have demonstrated here that the conformationally restricted nucleoside analogues 1 have a significant hybridization ability with complementary nucleic acids, especially with complementary RNA, which arises from the entropically favorable character of 1. These results should reveal a promising route to development of antisense methodology.

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