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# Propynyl Groups in Duplex, Hairpin and Triplex DNA: 7-Deazapurines and 9-Deazapurines

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### PROPYNYL GROUPS IN DUPLEX, HAIRPIN AND TRIPLEX DNA: 7-DEAZAPURINES AND 9-DEAZAPURINES

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- Laboratorium für Organische und Bioorganische Chemie, Institut für Chemie, Universität Osnabrück, Osnabrück, Germany and Center for Nanotechnology (CeNTech), Münster, Germany
  - Oligonucleotides containing 7-deazapurines or 9-deazapurines with propynyl groups at the 7- or 9-position were prepared. The stabilizing effect of the propynyl group was studied on DNA duplexes, hairpins and triplexes.

Keywords 7-Deazapurines, 9-Deazapurines, Oligonucleotides, Duplex, Hairpin, Triplex

#### INTRODUCTION

Among the various modifications carried out on the DNA nucleobases the introduction of the propynyl group has been shown to exert a stabilizing effect being useful for antisense technology and oligonucleotides hybridization used for diagnostic purposes. Recently, our laboratory has reported on 8-aza-7-deazapurines carrying 7-propynyl residues. Here, we compare the properties of oligonucleotides containing the 7-deazapurine nucleosides **1,2a-c** as well as the 9-deazapurine nucleosides **3a-c**. The phosphoramidites **4,5,6a-c** were prepared. The influence of the modified nucleosides on the stability of DNA duplexes, hairpins, and triplexes was studied (Scheme 1).

#### RESULTS AND DISCUSSION

#### Synthesis of the Phosphoramidites

The nucleosides **1c–3c** were synthesized from **1b–3b** using [(PPh<sub>3</sub>)<sub>4</sub>Pd(0)]/CuI catalyzed *Sonogashira* cross-coupling. The amino groups were blocked

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#### **SCHEME 1**

with either an acetamidine (1c), an isobutyryl or formamidine (2c), <sup>[2]</sup> or a formamidine residue (3c). The intermediates were converted into the corresponding DMT-derivatives and subsequently phosphitylated. This furnished the phosphoramidites 4c-6c, which were used in solid-phase synthesis (see Tables 1 and 2).

### The Effect of the 7-Propynyl Group on the DNA Duplex Stability

The displacement of four dG residues by 7-deaza-2'-deoxyguanosine (2a) within the duplex  $7 \cdot 8$  results in a decrease of the  $T_m$  value of  $3^\circ$  ( $9 \cdot 10$ ) (Table 1). The introduction of the 7-iodo substituents  $2b^{[3]}$  instead of 2a bring the stability to the level of the unmodified duplex (see  $11 \cdot 12$ , Table 1). Displacement of the 7-iodo by 7-propynyl groups enhances the  $T_m$  value further to  $53^\circ C$  ( $13 \cdot 14$ ) which equals to an increase of  $1.5^\circ$ /modification—a value which has been already found for 7-propynyl-substituted 8-aza-7-deaza-2'-deoxyguanosine within the same sequence. The incorporation of the 2'-deoxytubercidin derivative 1c as well as the 7-iodo compound  $1b^{[4]}$  results also in a stepwise stability increase, while the non-functionalized 1a shows the same behavior as dA (Table 1).

TABLE 1 T<sub>m</sub> Values and Thermodynamic Data of Oligonucleotides

| Duplex   | $T_{\rm m}$ (°C) | $\Delta H^{\circ}$ (kcal/mol) | $\Delta S^{\circ}$ (kcal/mol) | $\Delta G_{310}^{\circ}$ (kcal/mol) |
|--|------------------|-------------------------------|-------------------------------|-------------------------------------|
| 5'-d(TAG GTC AAT ACT) <b>7</b>                     | $47^a$           | -87                           | -243                          | -11.3                               |
| 3'-d(ATC CAG TTA TGA) 8                            | $50^b$           |                               |                               |                                     |
| 5'-d(TA <b>2a2a</b> TC AAT ACT) <b>9</b>           | $44^a$           | -92                           | -284                          | -3.6                                |
| 3'-d(ATC CA2a TTA T2aA) 10                         |                  |                               |                               |                                     |
| 5'-d(TA <b>2b2b</b> TC AAT ACT) <b>11</b>          | $48^a$           | -112                          | -348                          | -4.1                                |
| 3'-d(ATC CA <b>2b</b> TTA T <b>2b</b> A) <b>12</b> |                  |                               |                               |                                     |
| 5'-d(TA <b>2c2c</b> TC AAT ACT) <b>13</b>          | $53^{a}$         | -110                          | -314                          | -12.8                               |
| 3'-d(ATC CA2c TTA T2cA) 14                         |                  |                               |                               |                                     |
| 5'-d(TAG GTC AAT ACT) 15                           | $50^b$           | -77                           | -215                          | -10.7                               |
| 3'-d(ATC ClaG TTla TGA) 16                         |                  |                               |                               |                                     |
| 5'-d(TAG GTC AAT ACT) 17                           | $54^b$           | -85                           | -236                          | -12.3                               |
| 3'-d(ATC C1 <b>b</b> G TT1 <b>b</b> TGA) 18        |                  |                               |                               |                                     |
| 5'-d(TAG GTC AAT ACT) 19                           | $56^b$           | -85                           | -233                          | -12.5                               |
| 3'-d(ATC C1cG TT1c TGA) 20                         |                  |                               |                               |                                     |

<sup>&</sup>lt;sup>a</sup>10 mM Na-cacodylate, 10 mM MgCl<sub>2</sub>, 100 mM NaCl, pH 7.

<sup>&</sup>lt;sup>b</sup>60 mM Na-cacodylate, 100 mM MgCl<sub>2</sub>, 1 M NaCl, pH 7.

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**TABLE 2**  $T_{\rm m}$  Values of Oligonucleotide Hairpins

| Sequence  | <i>T</i> <sub>m</sub> (°C) |
|---|----------------------------|
| 5'-d(AGT AGG GAC CTA AGT AGG GAC CTA) 21                          | 70                         |
| 5'-d(AGT AGG G1aC CTA AGT AGG G1aC CTA) 22                        | 70                         |
| 5'-d(AGT AGG G1bC CTA AGT AGG G1bC CTA) 23                        | 76                         |
| 5'-d(AGT AGG G1cC CTA AGT AGG G1cC CTA) 24                        | 78                         |
| 5'-d(AGT AGG <b>2a</b> AC CTA AGT AGG <b>2a</b> AC CTA) <b>25</b> | 68                         |
| 5'-d(AGT AGG <b>2c</b> AC CTA AGT AGG <b>2c</b> AC CTA) <b>26</b> | 72                         |

 $^aMeasured$  at 260 nm in 100 mM NaCl, 10 mM MgCl<sub>2</sub>, and 10 mM Na-cacodylate buffer, pH = 7.0 with 2.5  $\mu M$  single-strand concentration.

# The Effect of 7-Substituted 7-Deazapurines on the dG-dA Hairpin Base Pairs

The oligonucleotide **21** (Table 2) forms a hairpin shown in Scheme 2. It contains three different d(GA) motifs: 1) a d(G-A) overhang, 2) an unpaired d(G-A) unit in the loop, and 3) two adjacent dG-dA pairs in the stem region. The replacement of dG by 7-deaza-2'-deoxyguanosine (**2a**) within the tandem d(G-A) pair leads to a slight destabilization while the replacement of dA by 7-deaza-2'-deoxyadenosine (**1a**) did not effect the stability (Table 2). Hoogsteen pairing is

SCHEME 2 Structure of the hairpin 21.

**TABLE 3**  $T_{\rm m}$  Values of Base-Modified Triplexes

| x  | $T_{\mathrm{m}}$ (°C) | X          | <i>T</i> <sub>m</sub> (°C) |
|----|-----------------------|------------|----------------------------|
| dC | 39, 65                | 3 <b>b</b> | 41, 64                     |
| 3a | 39, 65                | 3 <b>c</b> | 45, 65                     |

10 mM HEPES, 50 mM NaCl, 10 mM MgCl $_2$  0.5 mM Spermine, 1  $\mu M$  single-strand concentration, pH 6.5.

#### FIGURE 1

neither possible for dA nor for dG. Therefore, the most likely base pair motif is that of I. We now replaced the dA residues of the d(G-A) pairs by compounds **1b** and **1c**. The resulting hairpins **23** and **24** are significantly more stable than the hairpins **21** or **22**. Also dG was replaced by **2a** or **2c** leading to a slightly destabilized hairpin **25** (not functionalized at C7) and a stabilized hairpin (**26**) (Table 2). This means that the 7-iodo substituents as well as 7-propynyl groups have a positive effect on the stability of the non-canonical d(G-A) base pair which is represented by the bas pair motif II (propynyl modifications are possible on both sides).

# The Base Pairing Properties of 9-Deaza-2'-Deoxyguanosines in Triplex-DNA

DNA triplexes containing N<sup>7</sup>-2'-deoxyguanosine were described earlier. Here, we report on triplexes containing the nucleosides  $\bf 3a-c$ . The single strand  $\bf 27^{[6]}$  forms a double hairpin (motif III) with a triplex core. It shows two thermal transitions (Table 3). The lower one results from the melting of the 3'-terminus stabilized by Hoogsteen-pairs, the higher one is represented by the dissociation of the Watson-Crick base pair (duplex melting). Replacement of dC(28) (=X) by 9-deaza-2'-deoxyguanosine ( $\bf 3a$ )<sup>[7]</sup> does not change the stability of the triplex while the iodo compound  $\bf 3b$  or the propynyl derivative  $\bf 3c$  leads to a stepwise stabilization of the third strand without changing the  $\bf T_m$  value of the duplex structure (Figure 1).

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