



2-Amino-7-deazaadenine Forms Stable Base Pairs with Cytosine and Thymine

Akimitsu Okamoto, Kazuo Tanaka and Isao Saito*

Department of Synthetic Chemistry and Biological Chemistry, Faculty of Engineering, Kyoto University, and CREST, Japan Science and Technology Corporation, Kyoto 606-8501, Japan

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Abstract—2-Amino-7-deazaadenine (^{ADA}) was incorporated into oligodeoxynucleotides (ODN) and their base-pairing properties with natural nucleobases were investigated. In melting temperature (T_m) experiments, the duplex containing an ^{ADA}/C base pair showed a high stability comparable to that containing ^{ADA}/T base pair. Destabilization of the duplex usually observed for existing degenerate bases was not observed. However, the incorporation efficiency of dCTP was only 1.8% for TTP in single-nucleotide insertion reactions using polymerase. © 2001 Elsevier Science Ltd. All rights reserved.

Synthetic 7-deazapurine, a purine analogue in which the N7 is replaced by a carbon atom, is a promising nucleobase for studying the structure and function of nucleic acids.^{1–8} Several groups have incorporated these deazapurine analogues into oligodeoxynucleotides (ODNs), and their binding affinity toward DNA has been studied.^{9,10} In recent years, deazapurine has also been used as a tool for probing nucleic acid structures.^{11–13} While the thermodynamic parameters for the duplex formation of ODNs containing 7-deazapurine have been thoroughly studied, little is known about the base selectivity and fidelity of the base-pairing of 7-deazapurine. An investigation of the base-pairing properties of 7-deazapurine is important not only for structural studies of nucleic acids, but also for antisense and mutagenesis studies. We now wish to report that 2-amino-7-deazaadenine (**2**), a family of 2-aminoadenine (**3**) which is known to form a tight base pair with thymine by extra hydrogen bonding,^{14–16} acts as a superior degenerate base to form a stable base pair with both cytosine and thymine.

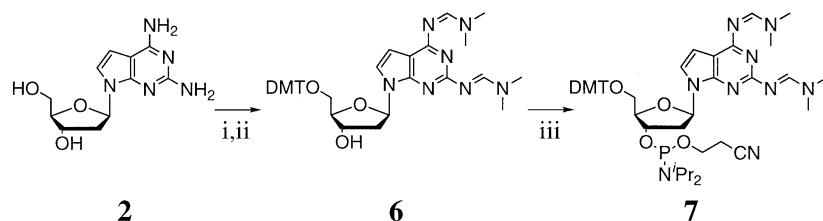
Nucleoside **2**¹⁷ was incorporated into ODNs using the conventional phosphoramidite method. The protected phosphoramidite of **2** was prepared as shown in Scheme 1.¹⁸ The thermal stabilities of the duplexes containing **2** and **3** were evaluated by determining the melting tem-

perature (T_m) of the duplexes, 5'-d(TTTGGTTXTTT)-3'/5'-(AAAYAACCAAA)-3' (X = **2** or **3**, Y = T or C), and from the change of A_{260} on heating. The normalized melting profile of the duplexes is shown in Figure 1. The duplex containing a **2**/T base pair showed a T_m value (13.8 °C) close to that of a **3**/T base pair (14.0 °C). The stability observed in the duplex containing a 'mismatched' **3**/C base pair (11.5 °C) remarkably decreased as compared with that for a **3**/T base pair, whereas the T_m of the duplex containing a **2**/C base pair (19.0 °C) was 5.2 °C higher than that containing a **2**/T base pair. The stable base pair with cytosine observed for **2** was striking, and in contrast to the destabilization of the **3**/C base pair.

The thermal stabilities of the duplexes containing **2**/C or **2**/T base pair were compared with those of the duplexes replaced by other base pairs. The T_m values of the duplexes are summarized in Table 1. The duplex containing a **2**/T base pair, which is expected to have extra hydrogen-bonding to thymine, showed a higher T_m than that of a 7-deazaadenine (**1**)/T base pair, as **3**/T duplex was more stable than A/T duplex. When guanine or adenine was incorporated opposite **2**, the duplex was remarkably destabilized. In contrast, the duplex containing a **2**/C base pair showed high duplex stability comparable to the G/C pair. It is worth noting that deazaadenine **2** can form a stable base pair not only with thymine, but also with cytosine.

Base pairs with both thymine and cytosine can be formed by the well-known degenerate bases O6-methyl-

*Corresponding author. Tel.: +81-75-753-5656; fax: +81-75-753-5676; e-mail: saito@sbchem.kyoto-u.ac.jp



Scheme 1. Synthesis of **2** phosphoramidite. Reagents and conditions: (i) *N,N*-dimethylformamide dimethylacetal, DMF, 55 °C, 8 h, 81%; (ii) 4,4'-dimethoxytrityl chloride (DMTCI), DMAP, pyridine, rt, 3 h, 86%; (iii) $(^i\text{Pr}_2\text{N})_2\text{PO}(\text{CH}_2)_2\text{CN}$, tetrazole, acetonitrile, rt, 4 h, quant.

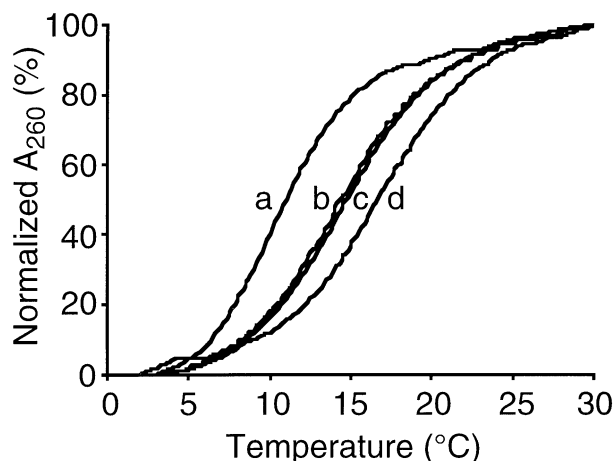


Figure 1. Normalized melting profile of the duplex 5'-d(TTTGGTTXTTT)-3'/5'-d(AAAYAACCAAA)-3' (X = **2** or **3**, Y = T or C). (a) **3**/C, (b) **2**/T, (c) **3**/T, (d) **2**/C.

guanine (**4**)^{19,20} and 2-amino-6-methoxyaminopurine (**5**)^{21,22} (Fig. 2). We measured the T_m of the ODNs containing these degenerate bases under the same conditions. Indeed, the degenerate bases **4** and **5** formed base pairs with both thymine and cytosine with similar stabilities (Table 1, entries 11–14). However, these base pairs remarkably destabilized the duplexes when compared to the duplexes containing normal A/T and G/C base pairs. Thus, the T_m experiments demonstrated that **2** is a superior degenerate base which can form a stable base pair with both cytosine and thymine without the duplex destabilization commonly observed for known degenerate bases like **4** and **5**.

High stability of the **2**/C duplex comparable to that of the **2**/T duplex was observed also in the T_m experiments

using GC-rich sequences. We measured the T_m of 5'-d(GCGATG2GTAGCG)-3'/5'-d(CGCTACNCATCG-C)-3' (N = C and T) in 10 mM PIPES (pH 7.0). The T_m of the duplex containing a **2**/C base pair was 44.4 °C, showing that the **2**/C duplex stability was nearly equal to the **2**/T duplex stability (T_m = 44.7 °C for **2**/T). In contrast, when T_m was measured in high salt concentration, the **2**/C duplex was less stable than that observed for the **2**/T duplex. In 10 mM PIPES in the presence of 100 mM sodium chloride and 10 mM magnesium chloride,²³ the T_m s of the duplexes containing the **2**/C and **2**/T base pairs were 56.1 and 62.0 °C, respectively. Although the T_m of the **2**/C duplex was risen under high salt condition, the stabilization of the **2**/C duplex by high salt was not so great as that observed for the **2**/T duplex.

In order to find out the efficiency of the base-pairing of 7-deazapurines with four natural nucleobases, we examined a single-nucleotide insertion reaction mediated by polymerase using a template containing **2**. The primer–template duplex was incubated with one of four natural deoxynucleotide triphosphates in the presence of a Klenow fragment (*exo*[−]) at 37 °C, and the incorporation efficiency of these four bases opposite **2** was determined by gel electrophoresis (Table 2).²⁴ The steady-state kinetic parameters, V_{\max} and K_m of the single-nucleotide insertion reaction were derived from a Lineweaver–Burk plot calculated from the intensities of the gel bands.²⁵ As shown in Table 2, the efficiency of dCTP insertion was considerably low: only 1.8% of that for TTP insertion, although the formation of a stable **2**/C base pair was observed in the T_m experiments. Table 2 shows that the low dCTP insertion efficiency was due to its large K_m value. The K_m of dCTP insertion was 25 times larger than that observed in TTP insertion,

Table 1. The melting temperatures (T_m) of duplexes containing various base pairs^a

5' -TTTGGTT X TTT-3'					
3' -AAACCA Y AAA-5'					
Entry	X/Y	T_m (°C)	Entry	X/Y	T_m (°C)
1	1 /T	12.2	8	A/T	12.8
2	2 /T	13.8	9	G/C	19.8
3	3 /T	14.0	10	A/C	6.8
4	2 /C	19.0	11	4 /T	9.3
5	3 /C	11.5	12	4 /C	8.5
6	2 /G	7.0	13	5 /T	6.7
7	2 /A	— ^b	14	5 /C	7.8

^aConditions: 2.5 μM duplex, 10 mM sodium cacodylate, pH 7.0.

^bNo sigmoidal melting curve was observed.

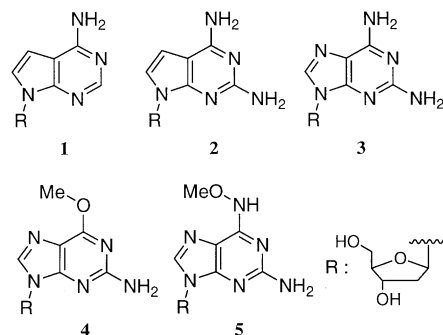


Figure 2. Structures of the synthetic 2'-deoxynucleosides used in this study.

Table 2. The steady-state kinetic parameters for the insertion of a single nucleotide into a template-primer duplex mediated by a Klenow fragment (*exo*[−])^a

5'–TAATACGACTCACTATAGGGAGA (2) –3' 3'–ATTATGCTGAGTGATATCCCTCT 2 GTCA–5'				
dNTP	V_{\max} (pmol U ^{−1} min ^{−1})	K_m (μM)	Efficiency (V_{\max}/K_m)	Relative frequency
TTP	45.3	1.14	39.7	100
dCTP	18.7	25.6	0.730	1.84
dGTP	13.1	62.0	0.211	0.531
dATP	0.994	22.0	0.0452	0.114

^aPrimer–template duplex was incubated with dNTP in the presence of a Klenow fragment (*exo*[−]) as a polymerase at 37 °C (see ref 23 for details). V_{\max} and K_m were calculated from a Lineweaver–Burk plot after three trials.

indicating that **2**/C base-pairing was inefficient in the interior of the enzyme pocket.

In order to confirm the T_m of the duplex elongated by single-nucleotide insertion, the T_m of the duplex consisting of the template and a 12-mer, 5'-d(CTA-TAGGGAGAPy)-3', possessing thymine or cytosine at the 3' end of the truncated primer (opposite **2** on a template strand) was measured. The **2**/C duplex showed a slightly higher T_m (28.0 °C) than the **2**/T duplex (26.2 °C), which was consistent with the T_m results shown in Table 1. Although the **2**/C base pair was as stable as the **2**/T base pair, the insertion of the dCTP opposite **2** was much slower than the TTP insertion. These results suggest that the **2**/C base pair formation in the polymerase reaction is more complicated when compared with the **2**/T base pair formation.

In summary, 7-deazaadenine **2** formed a stable base pair not only with thymine but also with cytosine without any duplex destabilization. While **2** was an effective degenerate base, only TTP was incorporated opposite **2** in a single-nucleotide insertion reaction, and the incorporation of dCTP was inefficient. Owing to its unique base-pairing property, 7-deazaadenine **2** is useful not only as a superior degenerate base that can be used for sequence-alternative hybridization, but also as a potent mutagen for site-directed mutagenesis which can induce the G/C to A/T transition during the DNA replication step.

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- Selected data for compound **6**. ¹H NMR (400 MHz, CDCl₃) δ 8.91 (br, 1 H), 8.67 (br, 1 H), 6.96 (d, 1 H, $J=3.7$ Hz), 7.42–6.78 (m, 14 H), 6.51 (d, 1 H, $J=3.8$ Hz), 4.55 (m, 1 H), 3.76 (s, 6 H), 3.26 (m, 2 H), 3.14 (d, 6 H, $J=10.5$ Hz), 3.09 (d, 6 H, $J=9.3$ Hz), 3.05 (m, 1 H), 2.41 (m, 2 H). ¹³C NMR (100 MHz, CDCl₃) δ 158.54, 158.49, 157.52, 157.08, 153.84, 144.66, 135.83, 135.81, 135.71, 135.69, 130.07, 128.17, 127.87, 126.84, 120.64, 113.16, 107.76, 101.52, 86.44, 84.76, 77.21, 72.84, 64.24, 64.08, 55.23, 55.22, 45.77, 41.17, 41.00, 40.57, 35.18, 34.79. MS (FAB, NBA/CH₂Cl₂) m/z (%) 678 [(M+H)⁺], HRMS (FAB) calcd for C₃₈H₄₃N₇O₅ [(M+H)⁺] 678.3401, found 678.3390.
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