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Synthesis and duplex stability of oligonucleotides containing 7-vinyl-7-deazaguanine as a strong electron-donating nucleobase

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Abstract

We synthesized 7-vinyl-7-deazaguanine (${}^{vz}G$), an N7-modified guanine base, which has a smaller oxidation potential than G and 7-deazaG. The incorporation of ${}^{vz}G$ into oligonucleotides was achieved by phosphoramidite chemistry on a DNA synthesizer. The T_m of the duplex containing ${}^{vz}G$ was higher than that of the corresponding duplex containing G or 7-deazaG. The stabilization of the duplex containing ${}^{vz}G$ is due to the favorable enthalpy change by increased π -stacking between the vinyl group and the 5'-flanking base. © 2000 Published by Elsevier Science Ltd.

The π -stacking of DNA bases plays an important role in duplex stabilization. Recently, the importance of π -stacking of hydrophobic nucleobases in the stabilization of duplex was noted by Kool¹⁻³ and Schultz.^{4,5} For example, Kool et al. suggested that stacking surface area would strongly affect the duplex stability of oligonucleotides containing hydrophobic dangling bases.³ Extended and well-overlapped π -stacking is also important in long-range charge transport through DNA duplex.⁶⁻¹⁰ Efficiency of charge transport is strongly affected by π -stacking in DNA structure such as A-form DNA,¹¹ DNA containing bulge sequence¹² and mismatched DNA.¹³ Furthermore, the expansion of π -stacking area lowers the ionization potentials of stacked bases.¹⁴ The extension of the π -conjugate system of 7-deazaguanine, which has a lower oxidation potential than that of guanine (G), would result in a further lowering of the oxidation potential. We therefore designed a new base, 7-vinyl-7-deazaguanine (^{vz}G) functionalized at guanine N7 where the vinyl group is extruded into the major groove upon duplex formation. Herein, we report the synthesis of ^{vz}G and the incorporation of ^{vz}G into oligonucleotides. We also measured the oxidation potential of ^{vz}G and examined the stability of ^{vz}G-containing DNA duplex.

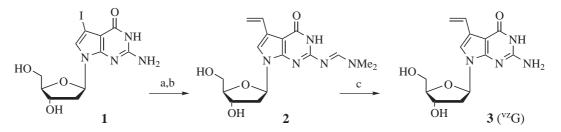
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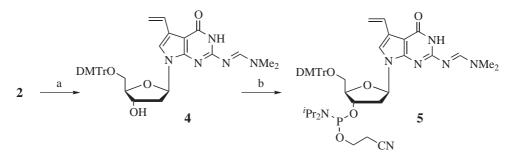
The synthetic route of ^{vz}G is shown in Scheme 1. The nucleoside 7-iodo-7-deaza-2'deoxyguanosine (1) was prepared according to the synthetic protocol previously reported.¹⁵ The exocyclic amino group of 1 was protected with DMF–diethylacetal (83%), and vinyl group was subsequently introduced into C7 by Stille coupling with tributyl(vinyl)tin to afford 2 in 60% yield.^{16,17} Nucleoside 2 was deprotected under alkaline conditions to afford ^{vz}G monomer 3 (83%). The UV absorption band for 3 showed a pronounced bathochromic shift by the substitution of the vinyl group at the 7-position (3, $\lambda_{max} = 264$ nm, $\varepsilon = 2820$; $\varepsilon_{320} = 550$; 7-deazaG, $\lambda_{max} = 259$ nm, $\varepsilon = 3760$; $\varepsilon_{320} = \sim 0$ in 10% acetonitrile–10 mM sodium cacodylate, pH 7.0).

In order to estimate the ability of **3** as an electron donating base, the oxidation potential of **3** was compared with those of 7-deazaG and G by means of cyclic voltammetry. The order of oxidation potential was as follows: G (1.07 V)>7-deazaG (0.74 V)>**3** (0.67 V) versus SCE, suggesting that **3** can serve as a highly electron-donating nucleobase in DNA duplex.



Scheme 1. (a) Dimethylformamide diethylacetal, DMF, 50°C, 2 h (83%); (b) tributyl(vinyl)tin, 5 mol% Pd(PPh₃)₄, DMF, 100°C, 5 h (60%); (c) 28% aqueous ammonia, methanol (1:1), rt, 5 h (83%)

For the preparation of phosphoramidite of v^z G, **2** was protected by a 4,4'-dimethoxytrityl group (67%), and then quantitatively converted to cyanoethyl phosphoramidite **5** (Scheme 2). Amidite **5** was employed in solid-phase synthesis of oligonucleotides on an automated DNA synthesizer. The time for the iodine oxidation of phosphite in each step was only 15 seconds, because v^z G incorporated into DNA was readily damaged by long exposure to iodine. Synthesized oligomers were deprotected overnight by aqueous ammonia at 37°C, and then purified by HPLC. The composition of oligomer was proved by MALDI-TOF mass spectrometry¹⁸ and enzymatic digestion by snake venom phosphodiesterase, nuclease P1, and alkaline phosphatase. An example of an HPLC chart after enzymatic digestion is shown in Fig. 1.



Scheme 2. (a) 4,4'-Dimethoxytritylchloride, DMAP, pyridine, rt, 15 h (67%); (b) ($^{1}Pr_{2}N)_{2}PO(CH_{2})_{2}CN$, tetrazole, acetonitrile, rt, 1 h (quant.)

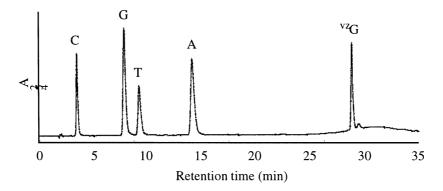


Figure 1. C18 reverse phase HPLC profile of 5'-d(CG^{vz}GAT)-3' after enzymatic digestion by snake venom phosphodiesterase, nuclease P1 and alkaline phosphatase (100 mM triethylammonium acetate, pH 7.0, 3–8% acetonitrile over 20 min, and then 8–35% over another 15 min)

The stability of the ^{vz}G-containing duplex 5'-d(TTTGGTT^{vz}GTTT)-3' (ODN1)/5'-d(AAA-CAACCAAA)-3' was examined by monitoring the melting temperature (T_m) . The T_m of the ^{vz}G-containing duplex was 2.1°C higher than that of the corresponding duplex containing G and 7-deazaG. Thermodynamic parameters for the duplex formation, ΔH , ΔS and ΔG , were obtained from van't Hoff plots. As shown in Table 1, the stabilization of ^{vz}G-containing duplex was due to the favorable enthalpy change in comparison with the duplexes replaced by G and 7-deazaG. This enthalpy change seems to be due to the increase of π -stacking between the vinyl group of ^{vz}G and the flanking base. The large favorable enthalpy change by the substitution of the vinyl group at the 7-position would compensate unfavorable entropy changes induced by the exclusion of water molecules in the major groove. B-form structures of the duplexes examined in T_m measurements (ODN1, 2 and 3) were confirmed by means of circular dichroism.

Table 1						
$T_{\rm m}$ values and	thermodynamic data	of oligonucleotide	duplexes			

5'-TTTGGTT**X**TTT-3' 3'-AAACCAACAAA-5'

	X	$T_{\rm m}^{\ a}$ (°C)	$-\Delta H^{\rm b}$ (kcal mol ⁻¹)	$-\Delta S^{\rm b}$ (kcal mol ⁻¹ K ⁻¹)	$-\Delta G_{298}$ (kcal mol ⁻¹)
ODN1	vzG	20.3	279	0.920	4.40
ODN2	G	18.2	232	0.765	3.52
ODN3	7-deazaG	18.2	204	0.670	4.14

^a Conditions: 1.25 µm duplex, 10 mM sodium cacodylate, pH 7.0.

^b Thermodynamic data were determined by plotting $(T_m)^{-1}$ versus $\ln(C_t/4)$. C_t (total concentration); 0.30, 0.60, 1.25, and 2.50 µm DNA duplex.

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The stacking surface area of ^{vz}G and 5'-flanking thymine in B-form DNA was calculated by the use of molecular modeling. The top views of TG/CA and T^{vz}G/CA where the sugar–phosphate backbone was deleted are shown in Fig. 2. The vinyl group of ^{vz}G overlapped considerably with the pyrimidine ring of 5'-thymine. Stacking surface area was calculated according to the method reported previously by Kool et al.^{3,19} The increase of stacking surface area by the incorporation of the 7-vinyl group was 14.9 Å² which corresponds to 12% of the total stacking surface area of TG/CA.

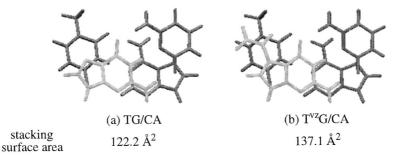


Figure 2. Molecular modeling of stacked geometries in B-form DNA. Models were built using canonical B-form geometry. DNA backbones were removed

In summary, we synthesized DNA duplex containing 7-vinyl-7-deazaguanine (^{vz}G) functionalized at guanine N7 where the vinyl group is extruded into the major groove. The extension of the π -conjugate system resulted in a lowering of the oxidation potential and a stabilization of the duplex by increased π -stacking between the vinyl group of ^{vz}G and the 5'-flanking base. Thus, ^{vz}G would be used as an effective electron-donating nucleobase for DNA-mediated electrontransfer chemistry.

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- 17. Selected data for compound **2**: ¹H NMR (400 MHz, CD₃OD) δ 8.65 (s, 1H), 7.23 (s, 1H), 6.87 (dd, J=11.2, 6.8 Hz, 1H), 6.56 (t, J=7.2 Hz, 1H), 6.03 (dd, J=16.0, 1.6 Hz, 1H), 5.09 (dd, J=9.2, 2.0 Hz, 1H), 4.49–4.45 (m, 1H), 3.96–3.91 (m, 1H), 3.73 (dd, J=7.2, 4.0 Hz, 1H), 3.71 (dd, J=7.2, 4.0 Hz, 1H), 3.19 (s, 3H), 3.11 (s, 3H), 2.58–2.51 (m, 1H), 2.30–2.23 (m, 1H); HRMS (FAB) calcd for C₁₆H₂₂N₅O₄ [(M+H)⁺] 348.3839, found 348.1667.
- 18. MALDI-TOF MS for 5'-GC^{vz}GAT-3': calcd for $C_{52}H_{65}N_{19}O_{28}P_4$ [(M–H)[–]] 1527.32, found 1527.12 and 5'-TTTGGTT^{vz}GTTT-3': calcd for $C_{113}H_{143}N_{30}O_{72}P_{10}$ [(M–H)[–]] 3383.28, found 3383.60.
- 19. Surface areas were calculated with molecular mechanics simulations using the MM2* force field as implemented in MacroModel 5.0.