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*N*²-Phenyldeoxyguanosine: Modulation of the Chemical Properties of Deoxyguanosine toward One-Electron Oxidation in DNA

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Owing to the remarkable structures produced by a π -stacked array of nucleotide base pairs, DNA has been envisaged as a novel medium of charge transport.¹ Experimental²⁻⁷ and theoretical⁸⁻¹¹ studies have shown that DNA actually mediates charge transport for some distances. A guanine radical cation (hole) produced by one-electron oxidation of DNA can migrate to a remote guanine site by successive hopping between neighboring guanines. The overall efficiency of the charge transport could be primarily determined by the rates of hole hopping and trapping. We have recently demonstrated the termination of charge transport by increasing the hole trapping rate at the site of N^2 -cyclopropyldeoxyguanosine (d^{CP}G),^{7a} in which cyclopropane ring opening functioned as a radical trapping device. These results imply that substitution of the exocyclic amino group of deoxyguanosine (dG) with functional groups could be an intriguing tool to modulate the reactivity of dG toward one-electron oxidation. We here report that incorporation of N²-phenyldeoxyguanosine¹² (d^{Ph}G) into duplex DNA dramatically suppressed oxidative decomposition, not only at dPhG, but more importantly, at dGs remote from the modified guanine.



The DNA oligomers containing dPhG and complementary strands used for the studies are listed in Table 1. All dPhG-containing ODNs were synthesized from corresponding ODNs containing 2-fluoroinosine by substitution of fluorine with aniline.13 dPhG was incorporated in the ODN PhGG1 probe by replacing G15 of the G15G doublet in the 21-mer ODN GG1. Complementary ODN CNBPU1 contains cyanobenzophenone-substituted uridine (d^{CNBP}U) as a photoinducible one-electron oxidant,7 whereas ODN T1 contained thymidine in place of d^{CNBP}U. In duplexes of GG1/^{CNBP}U1 and PhGG1/CNBPU1, the guanine radical cation was site selectively produced at G₈ by one-electron transfer to photoexcited d^{CNBP}U.⁷ ODN GG5 contained five GG doublets with identical neighboring sequences. The PhGG site was embedded in PhGG5(8) and **PhGG5(16)** by replacing G₈ and G₁₆ of **GG5**, respectively with ^{Ph}G. ODNs GG4 and PhGG4(8) lacked the G16G site of GG5 and **PhGG5(8)**, respectively, by replacing them with the T₁₆A sequence.

Table 1. Oligomers Used for the Studies^a

GG1: 5'-X	AG ₈ TGTGTA <u>G15</u> G V-3'
PhGG1: 5'-X	AG ₈ TGTGTA ^{Ph} G ₁₅ G V-3'
T1: 3'-Y	TC ACACAT C C W-5'
CNBPU1: 3'-Y CNB	PUC ACACAT C CW-5'
GG5:5'-X A	$\underline{\mathbf{G}}_{8}\underline{\mathbf{G}}TA\underline{\mathbf{G}}_{12}\underline{\mathbf{G}}TA \underline{\mathbf{G}}_{16}\underline{\mathbf{G}}TA\underline{\mathbf{G}}_{20}\underline{\mathbf{G}}TA\underline{\mathbf{G}}_{24}\underline{\mathbf{G}}V\text{-3}$
^{Ph} GG5(8): 5'-X A ^P	$^{h}\underline{G}_{8}\underline{G}TA\underline{G}_{12}\underline{G}TA \underline{G}_{16}\underline{G}TA\underline{G}_{20}\underline{G}TA\underline{G}_{24}\underline{G}V-3'$
^{Ph} GG5(16): 5'-X A	$\underline{G}_{8}\underline{G}TA\underline{G}_{12}\underline{G}TA^{Ph}\underline{G}_{16}\underline{G}TA\underline{G}_{20}\underline{G}TA\underline{G}_{24}\underline{G}V\text{-3}$
c-GG5 : 3'- Y T	C CATC CAT C CATC CATC CW-5'
GG4 :5'- X A	$\underline{\mathbf{G}}_{8}\underline{\mathbf{G}}\mathbf{T}\mathbf{A}\underline{\mathbf{G}}_{12}\underline{\mathbf{G}}\mathbf{T}\mathbf{A}$ TA $\mathbf{T}\mathbf{A}\underline{\mathbf{G}}_{20}\underline{\mathbf{G}}\mathbf{T}\mathbf{A}\underline{\mathbf{G}}_{24}\underline{\mathbf{G}}$ V-3'
^{Ph} GG4(8): 5'-X A ^P	$\mathbf{G}_{8}\mathbf{G}\mathbf{T}\mathbf{A}\mathbf{G}_{12}\mathbf{G}\mathbf{T}\mathbf{A}$ TA $\mathbf{T}\mathbf{A}\mathbf{G}_{20}\mathbf{G}\mathbf{T}\mathbf{A}\mathbf{G}_{24}\mathbf{G}\mathbf{V}$ -3'
c-GG4 : 3'-Y T	C CATC CAT AT ATC CATC C W-5'
$^{a}\mathbf{X} = ATTTAT; \mathbf{Y}$	X = TAAATA; V = TATTT; and W = ATAAA.

c-GG5 and **c-GG4** are complementary strands to **GG5** and **GG4**, respectively. The oxidation potential of $d^{Ph}G$ measured by cyclic voltammetry in DMF containing 0.1 M LiClO₄ was 0.70 V (vs Ag/Ag⁺, cf. G 0.67 V). The melting temperature of the $d^{Ph}G$ -containing 10-mer duplex d(GAT AGT ^{Ph}GGA C)/d(GTC CAC TAT C) was 3.7 °C higher than the corresponding normal G-containing duplex (50 μ M base concentration, 100 mM NaCl). CD spectra of the duplex showed a typical B-form structure (Figure S1).

Photoirradiation of duplex GG1/CNBPU1 at 312 nm for 60 min and subsequent piperidine treatment (90 °C, 20 min) produced a distinct cleavage band at the 5' side G of the G₁₅G doublet (Figure 1a, lane 2). In marked contrast, the cleavage at PhG15 in duplex PhGG1/CNBPU1 (lane 3) was considerably weaker than the cleavage at G_{15} in **GG1**/^{CNBP}U1 (lane 2). The band intensity of G_{15} relative to intact full length GG1 was 0.31, whereas the relative band intensity of PhG15 of PhGG1 was only 0.05. Suppression of the decomposition of dPhG by one-electron oxidation was further confirmed by riboflavin-sensitized oxidation of duplex PhGG1/T1. While strong cleavage occurred at G₁₅ of GG1/T1 (Figure 1b, lane 1), only a faint band was observed at PhG15 of PhGG1 (lane 2). We separately confirmed that strand cleavage of single-stranded dPhGcontaining oligomer d(ATT TAT AGT AGT AGT AGT ATT T) actually occurred at d^{Ph}G by the riboflavin-sensitized oxidation and subsequent piperidine treatment (Figure S2). The efficiency of the cleavage at d^{Ph}G was comparable to that at dG. Nucleoside analysis of d^{Ph}G-containing DNA by HPLC showed that 68% of d^{Ph}G remained intact in the duplex after 1 h of irradiation in the presence of riboflavin, whereas only 36% of dPhG remained intact in the single strand (Figure S3).

Oxidative decomposition of the d^{ph}G-containing duplex was suppressed not only at d^{ph}G but also remarkably at the GG sites

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Figure 1. Autoradiograms of denaturing sequencing gels for photoreactions of d^{Ph}G-containing oligomer duplexes. Photoirradiated ODNs were heated with piperidine and electrophoresed through a denaturing 15% polyacrylamide/7 M urea gel. Partial base sequences of ODNs were shown on the side. d^{CNBP}U was located opposite A shown with a box. (a) Duplexes were irradiated at 312 nm for 60 min. Lane 1, Maxam-Gilbert A+G sequencing reactions of GG1; lane 2, CNBPU1/GG1; lane 3, CNBPU1/PhGG1. (b) Duplexes were irradiated at 366 nm in the presence of riboflavin. Lane 1, T1/GG1; lane 2, T1/PhGG1; lane 3, A+G reactions of GG1. (c) Duplexes were irradiated at 366 nm in the presence of riboflavin. Lane 1, A+G reactions of GG5; lane 2, GG5/c-GG5; lane 3, PhGG5(16)/c-GG5; lane 4, PhGG5(8)/c-GG5; lane 5, PhGG4(8)/c-GG4; lane 6, GG4/c-GG4; lane 7, A+G reactions of GG4.



Figure 2. Band intensities (%) of GG sites relative to the intact full length bands obtained for the photoirradiated duplexes of (a) lanes 2, 3, and 4 and (b) lanes 5 and 6 shown in Figure 1c.

that were distant from d^{Ph}G (Figure 1c). Band intensities at GG sites of lanes 1-5 in Figure 1c were shown in Figure 2. While strand cleavage of photoirradiated GG5 in the presence of riboflavin occurred at all GG sites with comparable efficiency, the cleavage of GG sites of PhGG5(8) was suppressed at G12G and G16G in addition to ${}^{Ph}G_8G$ (Figure 2a). In contrast to $G_{12}G$, cleavage at $G_{24}G$ was only weakly suppressed compared with the cleavage in GG5, showing that the efficiency of suppression of strand cleavage decreased with an increase in distance from dPhG. Distance dependency of cleavage suppression was clearly shown in the oxidation of PhGG5(16) that contained dPhG in the middle of five GG sites. The efficiency of the cleavage was considerably reduced at all four GG sites in addition to the PhGG site. Significant insights into the mechanism of cleavage suppression by dPhG were obtained by the riboflavin-sensitized oxidation of GG4 and PhGG4(8). Strand cleavage of GG4 was observed at all four GG sites, whereas the cleavage of PhGG4(8) was strongly suppressed at G8G and G12G sites, but not at all at G₂₀G and G₂₄G sites (Figure 2b). The G₁₂G and G₂₀G sites in ^{Ph}GG4(8)/c-GG4 were separated by six intervening A-T base pairs, and the rate of hole transport between two sites was expected to be much smaller than that between G₈ and G_{12} (e.g., $2.5 \times 10^6 \text{ s}^{-1}$ for $G^{\bullet+}TTG \rightarrow GTTG^{\bullet+}$) and the estimated rate of hole trapping with water at the G site (e.g., $6 \times 10^4 \text{ s}^{-1}$).^{5b,6}

Thus, it is apparent that the efficiency of suppression of the oxidative decomposition at the given G sites in d^{Ph}G-containing duplex increased upon increasing the rate of hole transfer to d^{Ph}G.¹⁴

Our studies showed that (1) substitution of an exocyclic amino group of dG is effective in modulating the chemical properties of dG toward one-electron oxidation and (2) decomposition of the guanine radical cation was effectively suppressed near d^{Ph}G. These results indicate that d^{Ph}G is a prototype of nucleosides functioning as an intrinsic antioxidant of duplex DNA toward one-electron oxidation.

Supporting Information Available: CD spectra, HPLC, and PAGE analysis of dPhG-containing DNA (PDF). This material is available free of charge via the Internet at http://pubs.acs.org.

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- (14) These remarkable observations could be rationalized by assuming an annihilation process of the d^{Ph}G radical cation that was prevented from decomposing, leading to the formation of a piperidine-labile site, as shown in the scheme below. The rate of annihilation of $d^{Ph}G$ radical cation (k_{ann}) should be much faster than the hole trapping rate at GG (k_{trap1}) and ^{Ph}GG (k_{trap2}) to suppress the decomposition at these sites. When the hole transfer to d^{ph}G overrode the hole trapping at the given G site (i.e., $k_{HT} \gg k_{trap1})$, the hole was selectively depleted at d^{Ph}G by the annihilation process.

$$(GG)^{+} \xrightarrow{k_{HT}} (PhGG)^{+} \xrightarrow{k_{ann}} P_1 \qquad P_2 \xrightarrow{Ph} GG$$

P1, P2 denote pipeidine labile products.

Molecular modeling simulation showed that the phenyl group of d^{Ph}G in duplex DNA was located just in the middle of the minor groove, suggesting a considerable increase of solvent-accessible surface upon phenyl substitution. The calculated spin density of the $d^{Ph}G$ radical cation was delocalized on both the purine and phenyl rings. Since suppression of d^{ph}G decomposition was specific for duplex DNA, the annihilation process most likely occurred in the vicinity of the phenyl ring in the minor groove. One plausible mechanism of the putative annihilation process may involve a back-electron transfer from superoxide radical anion to a dPhG radical cation. In the case of riboflavin sensitization, the riboflavin radical anion is also conceivable as an electron donor.

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