Direct Observation of the Hole Protonation State and Hole Localization Site in DNA-Oligomers

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Abstract: In this work, it is shown that the incorporation of an 8-deuteroguanine (G•+) moiety in DNA-oligomers allows for direct determination at 77 K of (i) the location of holes (i.e., the radical site) within dsDNA at specific base sites, even within stacks of G, as well as (ii) the protonation state of the hole at that site. These findings are based on our work and demonstrate that selective deuteration at C-8 on a guanine moiety in dGuo results in an ESR signal from the guanine cation radical (G•*) which is easily distinguishable from that of the undeuterated guanine cation radical (G•+). G•+ is also found to be easily distinguishable from its conjugate base, the N1-deprotonated radical, G•(−H•). Our ESR results clearly establish that at 77 K (i) one-electron oxidized guanine in double stranded DNA-oligomers exists as the deprotonated neutral radical G(−H•) as a result of facile proton transfer to the hydrogen bonded cytosine and (ii) the hole is preferentially located at the 5′-end in several ds DNA-oligomers with a GGG sequence.

Introduction

Hole transfer through DNA has fostered intensive cross disciplinary interest as a result of both its fundamental significance and growing practical uses.1–4 Experimental studies employing a variety of techniques including UV—visible excitation followed by gel electrophoresis,5–4 γ or X-ray irradiation followed by electron spin resonance (ESR) in DNA systems including single crystals of DNA-oligonucleotides,1,3,5,6 cyclic voltammetry,2,3,5,7,8 and time-resolved techniques such as pulse radiolysis and flash photolysis9–14 have established unequivocally that, during charge transfer processes in double stranded DNA, guanine is the primary site of “hole” localization, i.e., site of electron loss. These results are in agreement with the theoretically predicted ionization potential value of guanine in the G:C base pair system.15–17

The efficiency (rate and extent) of the hole transfer processes between guanine moieties in DNA is controlled by two important proton transfer processes: (i) the intrabase proton transfer process within the G•+:C base pair1,3,6,9–11,15–17 and (ii) the proton transfer processes between the G•+:C base pair and the surrounding water.12,13 Intrabase proton transfer leads to separation of charge from spin (see Scheme 1), and thus, these intrabase proton transfer processes have been proposed to slow or stop the hole transfer.12,13 Such proton transfer processes will strongly depend on the pKₐ of the ion-radical involved.9–13 The pKₐ of a cation radical is invariably found to be lower than the parent compound. For example, the pKₐ of guanine (N1−H) is reported as 9.6 in aqueous solution at ambient temperature, while the corresponding pKₐ of the


guanine cation radical (G\(\cdot^+\)) is ca. 3.9.\(^{9,10}\) Based upon the p\(K_a\) values of G\(\cdot^+\) (ca. 3.9) and C(H\(\cdot^+\)) (ca. 4.5), Steenken had estimated the equilibrium constant for this intrabase proton transfer, \(K_{eq}\), between (C:G\(\cdot^+\)) and (C(+H\(\cdot^+\))):G(-H\(\cdot^+\)) to be ca. 2.5.\(^{9-11}\) This value of \(K_{eq}\) suggests that, at ambient temperature, the base pair (G\(\cdot^+:C\)) in DNA would have only partial proton transfer and both cation radical (30%) and deprotonated neutral radical (70%) forms would exist. There exists extensive discussion in the literature\(^{1-6,9-17}\) regarding the protonation state of the base pair (C:G\(\cdot^+\)). Theoretical studies have treated this issue of intrabase proton transfer along the N=H bond (Scheme 1) in the gas phase at the standard state (298 K, 1 atm) and have shown that \(\Delta S\) (proton transfer) = ca. 0 cal/mol/K and \(\Delta G\) (proton transfer) = 1.4 kcal/mol.\(^{15-17}\) However, until now, no definitive experiments had been performed to show evidence of intrabase proton transfer in the C:G\(\cdot^+\) base pair as shown in Scheme 1.

It is well-established that GG and GGG sites are more easily oxidized than isolated guanine sites.\(^2-4\) Thus, during oxidative hole transfer processes in DNA, GG and GGG sites are hot spots for localization of the hole.\(^2-4\) A number of experimental and theoretical studies suggest that there is a marked preference for hole localization at the 5’-G site in GG and GGG sequences.\(^2-4,18-24\) In GGG sequences, there was evidence for a small extent of hole localization at the central G.\(^2,4,18-24\) However, these experimental studies were carried out in aqueous solution at ambient temperature. Therefore, the final site for the hole localization was determined after the temperature-dependent hopping of the hole and subsequent reactions at sites where the hole is stabilized, e.g., via the nucleophelic addition of \(\text{H}_2\text{O}\) at C8 in the guanine moiety followed by an one-electron oxidation leading to the formation of 8-oxo-guanine (8-oxo-G) moieties.\(^{1,5,6,16-25}\) On the basis of nuclear reorganization energy (NRE) values, it can be theoretically expected, in part, that the hole could be preferentially located to a single guanine in a stacked guanine environment. The formation G\(\cdot^+\) in the G:C base pair has been shown to have a higher NRE value (0.34 eV) than that for the formation of A\(\cdot^+\) in A:T (0.10 eV).\(^{15,17-26}\) Consequently, the G\(\cdot^+:C\) base pair has a significantly different structure than the neutral G:C; while the A\(\cdot^+:T\) base pair has a near identical structure to neutral A:T. Therefore, guanine stacks are predicted to readily undergo reorganization and localize the charge (i.e., hole) to a single G, whereas adenine stacks would tend to delocalize the hole. Of course, solvent polarization effects and the effects of surrounding ions limit the extent of delocalization within the stack.\(^{23,26-28}\) However, there are no direct experimental results identifying the actual location of the hole in stretches of guanine in a ds DNA oligomer prior to the temperature-dependent hopping and subsequent reactions, e.g., 8-oxo-G formation, while recent work in our laboratory has given evidence for hole delocalization in A stacks even at a low temperature range (77–154 K)\(^{25}\) where the temperature-dependent hopping of the hole does not occur.\(^{25}\)

Thus, in stacked environments, there are two very important questions regarding oxidative hole formation in DNA prior to any subsequent reactions: (i) What is the protonation state of the hole? (ii) Where is the site of radical localization in (G\(_n\)) sequences?

We have recently found that selective incorporation of deuterium (D) at C-8 on guanine in dGuo [8-D-dGuo = G\(\cdot^+\)] reduces the C-8:H proton hyperfine coupling of ca. 8 G to the C-8:D coupling of ca. 1.2 G, thereby narrowing the spectral width and improving spectral resolution.\(^{29,30}\) Owing to the collapse of the anisotropic C-8:H hyperfine coupling, the ESR signal from the 8-D-dGuo cation radical (G\(\cdot^+\)) became easily distinguishable from that of the cation radical (G\(\cdot^+\)) in the undeuterated dGuo.\(^{29,30}\) Furthermore, the ESR spectrum of G\(\cdot^+\) is also distinguishable from the N1-deprotonated neutral radical, G\((-H\cdot)^+\), owing to their different line shape, line width, and g values.\(^{31}\) Hence, in this work, we incorporate 8-D-dGuo (G\(\cdot^+\)) into sequences of ss and ds DNA (e.g., TG\(\cdot^+\)T and d[G\(\cdot^+\)CG\(\cdot^+\)CG\(\cdot^+\)CG\(\cdot^+\)]\(\perp\)) and, based on their ESR spectra, directly provide evidence of the protonation state of the hole at specifically labeled guanine sites in these sequences. Moreover, owing to clear and observable difference in the ESR spectra of G\(\cdot^+\) and G\(\cdot^+(\cdot-H\cdot)^+\) as well as that of G\(\cdot^+(\cdot-H\cdot)^+\) and G\((-H\cdot)^+\), we find that a hole on the G\(\cdot^+\) is easily distinguishable from a hole on G. Thus, we demonstrate that this novel method provides for the direct determination of the location of holes within dsDNA at specific guanine sites, even within stacks of G at 77 K (i.e.,

\(\text{Adhikary et al.}\))

\(\text{Scheme 1. Schematic Representation of Prototropic Equilibria in the G}\(\cdot^+\):C Pair}\n
\(\text{B J. AM. CHEM. SOC.} \text{ VOL. xxx, NO. xx, XXXX}\)
before the temperature-dependent hopping of the hole) as well as their state of protonation.

Materials and Methods

**Compounds.** 2'-Deoxyguanosine (dGuo) and lithium chloride (99% anhydrous, SigmaUltra) were obtained from Sigma Chemical Company (St Louis, MO). Potassium persulfate (crystal) was purchased Mallinckrodt, Inc. (Paris, KY). 8-DiGlu (G*)\(^{29,30}\) was synthesized according to the procedure of Huang et al.\(^{1a}\) by dissolving dGlu in D\(_2\)O in the presence of triethylamine (TEA) (Fischer Scientific, NJ) and by warming the solution at 55–60 °C for 16 h. Employing a Bruker 200 MHz NMR spectrometer, in G*, the degree of deuteration was found to be >96% by 1D signal integration.

The phosphoramidite of G* was synthesized by Glen Research, VA. Starting with this phosphoramidite of G*, the dsDNA oligomers dG and G* were prepared by dissolving earlier works with DNA-nucleosides and nucleotides, the solutions (Figure S1). These values were accounted for in the analyses of the fractions, which, upon annealing, soften to allow for molecular migration and solution phase chemistry.\(^{25,26,29,30,32}\) All of these glassy samples are stored at 77 K in Teflon containers in the dark.

**Glassy Sample Preparation.** According to our previous works, homogeneous solutions (7.5 M LiCl D\(_2\)O) of dGlu, G*, and the DNA-oligomers were thoroughly bubbled with nitrogen to remove oxygen.\(^{25,26,29,30,32,34}\) Transparent glassy samples were then prepared drawing the solution into 4 mm Suprasil quartz tubes (cat. no. 734-PQ-8, WILMAD Glass Co., Inc., Buena, NJ) followed by cooling to 77 K. We note that these samples at low temperatures are not crystalline solids but are glassy homogeneous supercooled solutions, which, upon annealing, soften to allow for molecular migration and solution phase chemistry.\(^{25,26,29,30,32,34}\)

**Formation of One-Electron Oxidized dGlu, G*, and DNA-Oligomers via Thermal Annealing and Storage of These Samples.** Following our previous works, the one-electron oxidized dGlu and the one-electron oxidized DNA-oligomers were produced via annealing in the dark in a variable temperature assembly (Air Products) in the temperature range 100–160 K employing cooled nitrogen gas.\(^{34}\) We have annealed samples for 15–30 min in the dark at 155 ± 2 K which softens the aqueous (D\(_2\)O) glass facilitating molecular migration of Cl\(^{+*}\), Cl\(^{+}\) one-electron oxidizes the solute with concomitant decrease of the ESR spectrum of Cl\(^{+*}\) with a simultaneous increase in ESR spectrum of a one-electron oxidized guanine moiety in dGlu, G*, or DNA-oligomers.\(^{1e,25,26,29,30,32,34}\) After annealing, these samples were immediately immersed in liquid nitrogen (77 K) and stored in Teflon containers at 77 K in the dark before ESR studies began.

**Electron Spin Resonance.** The ESR spectra of these γ-irradiated and subsequently annealed samples were recorded at 77 K and at 40 dB (20 μW) using a Varian Century Series ESR spectrometer operating at 9.2 GHz with an E-4531 dual cavity, 9-in. magnet and with a 200 mW klystron. For field calibration, we have used Fresny’s salt (g (center of the spectrum) = 2.0056, As = 13.09 G) as per our previous works.\(^{1e,25,26,29,30,32,34}\)

**Analyses of ESR Spectra.** Following our previous work,\(^{26}\) each spectrum has been stored in a 1000-point array with field calibration marks from the three ESR lines of the Fresny’s salt. We have estimated the fraction that a particular radical contributes to the overall ESR spectrum by employing doubly integrated areas of first derivative ESR spectra. The fractional contribution of the radicals in each experimental ESR was obtained by use of programs developed in our laboratory (ESRADSUB, ESRPLAY) by fitting benchmark spectra to the ESR signal of interest via linear least-squares analysis.\(^{25,26,34}\) In these analyses an error function (R) is employed as a test for goodness of fit from the sum of the differences between the squares in heights of the experimentally obtained spectrum and that of the simulated spectrum obtained employing the benchmark spectra (R = \(\frac{1}{2} \sum (\text{exp}(i)-(\Sigma \text{exp}(i))^{2})\)). To estimate the error limits in our analyses, nonoptimized fits were generated and the value of the error (R) was plotted against the fraction of the contributing radical benchmark spectra (see Supporting Information Figure S2 as an example). On this basis, we have estimated the error limit of our analyses as ±10%.

**Results and Discussion**

The organization of this paper is as follows: (i) We first present evidence that the ESR spectra of pairs G* + vs G* +.

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These differences in ESR spectra allow the determination of the central marker is at and in the other figures represent positions of Fremy’s salt resonances (the one-electron oxidation by Cl$_2^*$ was carried out via thermal annealing at 155 K in the dark. Parts A and B clearly show that deuteration at C8 replaces the C-8-H proton hyperfine coupling from ca. 8 G in G*^• in its N1-deprotonated neutral radical, G(−H)^• with a ca. 1.2 G (C-8-D coupling) thereby narrowing the ESR spectral width and improving the spectral resolution in the deuterated species, G*^• and G(−H)^•. Part C shows that, in dGuo, G^• (black) is not clearly distinguishable from G(−H)^•, whereas part D represents that, in G*, owing to the smaller C-8-D coupling, G^• (black) is readily distinguishable from its N1-deprotonated neutral radical, G(−H)^• (red), because of their different line shapes, line widths, and g$_{||}$ values. All spectra were recorded at 77 K. The three triangular calibration markers in this figure and in the other figures represent positions of Fremy’s salt resonances (the central marker is at g = 2.0056, and each of three markers is separated from one another by 13.09 G).

**Figure 1.** ESR spectra found for (A) guanine cation radicals (G^•*, G^••*) and (B) deprotonated guanine neutral radicals (G(−H)^•, G(−H)^•). These radicals were produced by Cl$_2^*$ oxidation of dGuo (3 mg/mL) (blue) and of G^• (3 mg/mL) [G^• = 8-D-dGuo, 96% D] (red) at pH’s ca. 3 and 9 in 7.5 M LiCl glasses in D$_2$O in the presence of K$_2$SO$_4$ (5 mg/mL) as an electron scavenger. The one-electron oxidation by Cl$_2^*$ was carried out via thermal annealing at 155 K in the dark. Parts A and B clearly show that deuteration at C8 replaces the C-8-H proton hyperfine coupling from ca. 8 G in G*^• and in its N1-deprotonated neutral radical, G(−H)^• with a ca. 1.2 G (C-8-D coupling) thereby narrowing the ESR spectral width and improving the spectral resolution in the deuterated species, G*^• and G(−H)^•. Part C shows that, in dGuo, G^• (black) is not clearly distinguishable from G(−H)^•, whereas part D represents that, in G*, owing to the smaller C-8-D coupling, G^• (black) is readily distinguishable from its N1-deprotonated neutral radical, G(−H)^• (red), because of their different line shapes, line widths, and g$_{||}$ values. All spectra were recorded at 77 K. The three triangular calibration markers in this figure and in the other figures represent positions of Fremy’s salt resonances (the central marker is at g = 2.0056, and each of three markers is separated from one another by 13.09 G).

G(−H)^• vs G^•(−H)^•, and G^•* vs G^•(−H)^• are each distinguishable from each other in nucleotides. (ii) We then show these differences in ESR spectra allow the determination of the protonation state of one-electron oxidized guanine in ss and ds DNA-oligomers. (iii) Finally, recognizing that the hole in ds DNA is in the form of G(−H)^•, we then use the spectra of G(−H)^• and G^• (−H)^• obtained from the ds DNA-oligomers as benchmarks to analyze the spectra of the ds DNA-oligomers containing GGG sequences and assign the position of the hole in these oligomers.

**Figure 2.** ESR spectra of (A) G^•* (black) and G^•(−H)^• (red) obtained from glassy (7.5 M LiCl in D$_2$O) samples of G^• (G^• = 8-D-dGuo, 96% D] (3 mg/mL). (B) One-electron oxidized ss oligomer, TG*T, (4.5 mg/mL) in blue after subtraction of 20% of the G(−H)^• spectrum (to account for 80% 8-deuteration level (see Supporting Information Figure S1). In part B, the match between the experimentally observed spectrum of one-electron oxidized TG*T (blue) with that of the simulated (pink) spectrum obtained using the spectra of G^•* and G^•(−H)^• shown in (A) as benchmarks indicate that, for one-electron oxidized TG*T at pH 5, G^•* and G^•(−H)^• are present in equal amounts. (C) Spectrum of the one-electron oxidized ds DNA oligomer d(G*CG*CG*CG*C) (2 mg/mL) after subtraction of 20% of the G(−H)^• spectrum (to account for 80% 8-deuteration level (see Supporting Information Figure S1), again with G^•(−H)^• from A in red superimposed. The match of green and red spectra in (C) clearly shows that one-electron oxidized guanine in ds DNA oligomer d(G*CG*CG*CG*C) exists as G^•(−H)^•. One-electron oxidation of the monomer and the DNA-oligomers were carried out via thermal annealing at 155 K. All spectra were recorded at 77 K. Line shape and g value differences between G^•* and G^•(−H)^• allow for this analysis.

**Protonation State of G^•+ in ss and in ds DNA Oligomers.** The ESR spectra of G^•* (black) and its N1-deprotonated radical, G^•(−H)^• (red), both from the nucleoside, dGuo, from Figure 1D are repeated in Figure 2A and are employed as benchmarks for the analyses of the subsequent spectra of ssDNA-oligomer (Figure 2B) and dsDNA-oligomer (Figure 2C).

In Figure 2B and C, we present the ESR spectra of one-electron oxidized DNA-oligomers TG*+T (blue) (a model for ss DNA) and d[G^•CG*CG*C]$_2$ (green) (a model for ds DNA), respectively, at pH ca. 5 in 7.5 M LiCl in D$_2$O after reaction with Cl$_2^*$. One-electron oxidation of the DNA-oligomers by Cl$_2^*$ was carried out via annealing the glassy aqueous samples in the dark at ca. 155 K.

In Figure 2B, the ESR spectrum (blue) of one-electron oxidized TG*+T at the native pH ca. 5 of 7.5 M LiCl glass/ D$_2$O after accounting for the fact that TG*+T had incorporation of G^• at an 80% level (see Supporting Information Figure S1) is presented. Over this spectrum, we superimpose the best fit
In the dark at 155 K for 20 min. oxidized oligomers shown in (A) to (E) were obtained by thermal annealing. The relative error (see Materials and Methods and Supporting Information Figure S3) for in our analyses (see Supporting Information Figure S1). We deuterated dropped to 50% for these oligos which was accounted. The fact that the totals of G*(-H)* on the three sites vary from 95 to 105% (vs 100% expected) gives some confidence in the hole localization mainly at the 5'-G end (ca. 60%) with lesser amounts at the middle and the 3'-end (ca. 15 – 20%).

We note here that the percentages of G*(-H)* and G*+(-H)* (see Table 1 and Figure 3) are obtained independently and separately for each DNA-oligomer. Thus the hole localization on G* in each of the three positions in GGG (G*GG, GG*G, and GGG*) give independent assessments of the distribution. The site of hole localization can be identified in such ds DNA-oligomers.

Hole Protonation State/Localization Site in DNA-Oligomers

In Figure 2C is the spectrum obtained after subtraction of 20% of the G(-H)* spectrum (Figure 3A) of the one-electron oxidized ds DNA-oligomer d[GGGCGGCGC]2 to G(-H)*. This assignment is clearly established by the very close similarities in spectral parameters (overall hyperfine splitting, anisotropic nitrogen hyperfine couplings, line shape, and the center of the spectrum) in the one-electron oxidized d[GGGCWWCGGC]2 and G(-H)* spectrum35 from dGuo (see Supporting Information Figure S3 for evidence). We also find that the ESR spectra of one-electron oxidized ds DNA-oligomers d[GGGCCCGC]2 and d[GGGC-CC]2 are identical (see Supporting Information Figure S4). Thus, it can be concluded that irrespective of the occurrence of the guanine moiety in the ds DNA sequence as a continuous stretch (e.g., GGG) or as a single G unit, one-electron oxidation of the guanine moiety in ds DNA would lead to G(-H)* formation. Figure 3A and B, therefore, show ESR spectra of G(-H)* and G*+(-H)* spectra obtained in the glassy aqueous samples of d[GGGCWCGGC]2 and d[GGGCWCGC]2, respectively. These spectra are used below as benchmarks for G*(-H)* and G*+(-H)* to analyze the fractional compositions of G(-H)* and G*(+H)* in the ESR spectra shown in Figure 3C to E.

In Figure 3C to E, we show the ESR spectra of one-electron oxidized d[GGGCCCGC]2 with G* moieties at the 5'-end, the middle, and the 3'-end of the GGG sequence. Analyses of spectra in Figure 3C to 3E shown in Table 1 demonstrate that the hole localizes mainly at the 5'-G end (ca. 60%) with lesser amounts at the middle and the 3'-end (ca. 15 – 20%).

We note here that the percentages of G*(-H)* and G*+(-H)* (see Table 1 and Figure 3) are obtained independently and separately for each DNA-oligomer. Thus the hole localization on G* in each of the three positions in GGG (G*GG, GG*G, and GGG*) give independent assessments of the distribution. The fact that the totals of G*(-H)* on the three sites vary from 95 to 105% (vs 100% expected) gives some confidence in the method.

One-electron oxidation of the isotopically substituted ds DNA oligomers, viz. d[TGG*GCGCCA], d[TGG*GCGCCA], and d[TGG*GCGCCA]2 that were HPLC purified, were carried out as described above. The ESR spectra of these one-electron oxidized ds DNA oligomers were recorded at 77 K, and these spectra were found to be very similar to the corresponding ESR spectra shown in Figure 3C to E (see Supporting Information Figure S5). Using the G*+(-H)* spectrum from HPLC-purified d[G*+CG*CG*CG*C]2 and the G*+(-H)* spectrum from dGGGCGGGCGC2 as benchmarks, the analyses of radical percentages in these oligomers show a clear preference for localization of the hole at the 5'-G end (Table 1). Note that owing to the further HPLC purification the level of deuteration dropped to 50% for these oligos which was accounted for in our analyses (see Supporting Information Figure S1). We also note here that by employing the G*+(-H)* spectrum from G* (96% deuterated) and the G*(+H)* spectrum from dGuo as benchmarks, we have obtained percentages of G*(+H)* and G*+(-H)*.

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<tr>
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<tr>
<td>d[TGG*GCGCCA]2</td>
<td>G*+(-H)* 30</td>
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<tr>
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Figure 3. Spectra A and B are the benchmark spectra of G(−H)* and G*+(−H)* (G* = 8-D-dGuo). These spectra are for the glassy samples of d[GGGCCCGC]2 and d[G*CG*CG*CG*C]2 and are assigned to G(−H)* and G*(+H)*. Note that the spectrum in B is 80% 8-deuterated. This spectrum is used as the basis function as it self-corrects in our analysis. Spectra C to E are the one-electron oxidized d[GGGCCCGC]2 with G* moieties at the 5′-end, middle, and 3′-end of the G stack, respectively. Analyses using G(−H)* and G*(+H)* spectra show that the site of hole localization is ~60% at the 5′-end. We have expressed the radical percentages to ±10% relative error (see Materials and Methods and Supporting Information Figure S2). All the spectra were recorded at 77 K, and the spectra of one-electron oxidized oligomers shown in (A) to (E) were obtained by thermal annealing in the dark at 155 K for 20–30 min.

The ESR spectra of these one-electron oxidized ds DNA oligomers were recorded as described above. The ESR spectra of these one-electron oxidized ds DNA oligomers were recorded at 77 K, and these spectra were found to be very similar to the corresponding ESR spectra shown in Figure 3C to E (see Supporting Information Figure S5). Using the G*+(-H)* spectrum from HPLC-purified d[G*+CG*CG*CG*C]2 and the G*+(-H)* spectrum from dGGGCGGGCGC2 as benchmarks, the analyses of radical percentages in these oligomers show a clear preference for localization of the hole at the 5'-G end (Table 1). Note that owing to the further HPLC purification the level of deuteration dropped to 50% for these oligos which was accounted for in our analyses (see Supporting Information Figure S1). We also note here that by employing the G*+(-H)* spectrum from G* (96% deuterated) and the G*(+H)* spectrum from dGuo as benchmarks, we have obtained percentages of G*(+H)* and G*+(-H)*.
G(−H•) in these ds DNA-oligomers (see Supporting Information) and these percentages are found to be very similar to those shown in Table 1.

Therefore, the radical percentages of G(−H•) and G•*(−H•) in the GGG sequence are similar for ds DNA-oligomers with and without single nucleotide extension at the ends, 5′-T and 3′-A (see Table 1). As a result, we conclude that the presence of 5′-T and 3′-A units in the ds DNA oligomers with GGG stretches did not alter the preferential location of the hole at the 5′-G site in these sequences.

Conclusion

(i) Sites of Hole Localization and Protonation States of One-Electron Oxidized Guanine in DNA-oligomers: Our work directly finds that the hole is preferentially localized at the 5′-end in GGG sequences at 77 K. Our results also show that the hole is not fully localized to the 5′ G site. Previous experimental results obtained in aqueous solution at room temperature had previously suggested that in GGG sequences the 5′-G would likely be the preferential site of hole localization. In those experiments, it was the preferential formation of 8-oxo-G at the 5′-end of the GGG sequences in ds DNA-oligomers after photo-oxidation that suggested the preferential hole localization at the 5′-end. In agreement theoretical calculations point out that in a GGG sequence, the 5′-end is the most stable site. Further, owing to significant NRE values, guanine stacks are not predicted to share a charge and are expected to readily undergo reorganization so as to localize the charge (i.e., hole) on a single G. Our experiments show no evidence for delocalization of charge as it would be expected to markedly alter the ESR spectra. Any comparison of 77 K hole localization with 8-oxo-G formation of course depends on the kinetics of hole conversion to 8-oxo-G formation being independent of the base sequence. Experiments show no such dependence of 8-oxo-G formation on the base sequence. As a consequence, the agreement between our results regarding sites of hole localization at 77 K with those results found at room temperature for the distribution of the formation of 8-oxo-G is reasonable.

(ii) Protonation State of One-Electron Oxidized Guanine in DNA-oligomers: We find that one-electron oxidized guanine is entirely the proton transferred radical (C(+H•):G(−H•)) at 77 K in ds DNA. At ambient temperatures, a prototropic equilibrium between G•*:C and C(+H•):G(−H•) would be established that would still favor the proton transferred species, as it is the thermodynamically stable state (see Introduction). This equilibrium has been proposed by a number of workers as a gating switch to hole transfer along the DNA strand. The presence of G•*:C would also allow for nucleophilic addition of H2O that leads to the formation of 8-oxo-G. It is well established that one-electron oxidized guanine must remain in its cation radical form to undergo water addition. Thus, Steenken’s proposition that (C(+H•):G(−H•)) would be the dominant species at ambient temperature is supported by our work.

(iii) Prototropic Equilibrium of One-Electron Oxidized Guanine in DNA-oligomers: Previous theoretical calculations carried out for the base pair (G•*:C) in the gas phase at standard state (298 K, 1.0 atm) predict that ΔS (proton transfer) = ca. 0 cal/mol/K and ΔG (proton transfer) = 1.4 kcal/mol. However, our ESR studies at low temperature clearly demonstrate that the base pair (G•*:C) exists as (C(+H•):G(−H•)) i.e., the free energy, ΔG, for proton transfer within the base pair (G•*:C) is negative. This disagreement between our experimental results and previous theoretical work is small in magnitude and could easily arise from the neglect of stacking and water interactions in the gas phase calculations.

(iv) Why is not the hole fully localized at the 5′-end? It is clear from the results reported here and others that the 5′-end is the dominant site for hole localization, although a significant fraction of the holes are also localized at the middle and 3′-end (Table 1). This indicates that the energetic preference for the 5′-end is small and that variations in the local environment such as counterion placement or structural variations are sufficient to stabilize the hole at the other sites. We note that there is no evidence for delocalization of the hole in the GGG stacks. This is in contrast to stacks of A for which some delocalization of hole is theoretically predicted via charge-resonance interaction. This difference in the behavior of holes in guanine vs adenine stacks has been theoretically attributed to the lack of significant nuclear reorganization on formation of the adenine cation radical. This allows holes in A stacks to be shared, whereas in G stacks the larger reorganization energy causes localization to a single base.

(v) Advantage of the Technique: The most important feature of incorporation of G* (8-deuteroguanine) in oligos is the clear resolution of the ESR spectra of one-electron oxidized ds DNA oligomers with G* at specific sites in GGG. Thus we are able to identify directly not only the protonation state of the G moiety but also the site of localization of the hole in DNA sequences having stretches of G’s.

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Supporting Information Available: Figure S1: Analyses for deuteration levels in the G* labeled ds DNA-oligomers. Figure S2: Estimation of relative error regarding fitting of the experimentally observed spectrum using benchmarks. Figure S3: ESR spectrum of the guanine neutral radical (G(−H•)) from dGuo and d[GCGCGCGC]2. Figure S4: ESR Spectra of the one-electron oxidized dsDNA-oligomers: d[GCCGCCGGC]1 and d[GCGCGCGC]1. Figure S5: ESR Spectra of the one-electron oxidized ds DNA-oligomers with and without single nucleotide extension at the ends, 5′-T and 3′-A. Analyses: Analyses of the ESR spectra of ds DNA-oligomers shown in Figure S5 using guanine neutral radical (G(−H•)) from dGuo and guanine neutral radical (G*(−H•)) from G* [G* = 8-D-dGuo, 96% deuterated]. This material is available free of charge via the Internet at http://pubs.acs.org.

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