Abstract: Many transition metal complexes mediate DNA oxidation in the presence of oxidizing radiation, photosensitizers, or oxidants. The final DNA oxidation products vary depending on the nature of metal complexes and the structure of DNA. Here we propose a mechanism of oxidation of a nucleotide, deoxyguanosine 5′-monophosphate (dGMP) by trans-△1,2-diaminocyclohexanetetrachloroplatinum (trans-Pt(dA)(1,2-(NH2)2C6H4)2Cl4, [PtIICl4(dach)]) to produce 7,8-dihydro-8-oxo-2′-deoxyguanosine 5′-monophosphate (8-oxo-dGMP) stoichiometrically. The reaction was studied by high-performance liquid chromatography (HPLC), 1H and 31P nuclear magnetic resonance (NMR), and electrospray ionization mass spectrometry (ESI-MS). The proposed mechanism involves PtIV binding to N7 of dGMP followed by cyclization via nucleophilic attack of a phosphate oxygen at C8 of dGMP. The next step is an inner-sphere, two-electron transfer to produce a cyclic phosphodiester intermediate, 8-hydroxyguanosine cyclic 5′,8′-(hydrogen phosphate). This intermediate slowly converts to 8-oxo-dGMP by reacting with solvent H2O.

Introduction

Oxidative damage of DNA plays a critical role in mutagenesis, carcinogenesis, aging, and lethality.1 DNA is oxidized by reactive oxygen species, ionizing radiation, and transition metal complexes. These oxidation reactions may induce modifications within the sugar or base moieties of DNA, leading to strand scission or base modification. Many transition metal complexes, including those of V, Cr, Mn, Re, Ru, Os, Co, Rh, Ni, Pd, and Cu, have been studied as potential probes of nucleic acid structure and for potential application as drugs.2 They mediate DNA oxidation in the presence of oxidizing radiation, photosensitizers, or oxidants. Complexes including CrVI, MnII, FeIII, CoII, NiII, and CuII in the presence of O2, H2O2, or O3− generate radicals that abstract one electron from guanine. FeIII-bleomycin, FeII-EDTA/H2O2, and photoinitiated charge-transfer complexes (RuIII) modify DNA by direct strand scission.3

High-valent oxometal (MnIV=O, RuIV=O, OsIV=O, CrV=O) complexes oxidize DNA at both sugar and base sites. The oxomanganese porphyrin, MnTMPyP/KHSO4, abstracts two electrons from guanine to produce the guanine cation, G+. The major oxidation product (>90%) from monomeric G is 2-amino-5-[(2-deoxy-β-D-erythro-pentofuranosyl)amino]-4H-imidazo[4,1-b]pyridine (imazodilone), but several other oxidation products are formed from double-stranded oligonucleotides.4,5 Oxidation of a dinucleoside monophosphate d(GpT) by MnIV=O, proposed to be a two-electron mechanism, produces dehydroguanidino-hydantoin as a major oxidation product.4c RuIII complexes coordinate to the N7 of deoxyguanosine5a and inosine5b and facilitate autoxidation of the nucleosides to yield the corresponding 8-ketonicotides. RuIV=O complexes oxidize guanine via an inner-sphere atom transfer to produce 7,8-dihydro-8-oxo-guanine (8-oxo-G).5 High-valent nickel complexes (NiII) formed from NiCR/KHSO4 bind to N7 of G and react by a one-electron mechanism.7 CrV=O complexes initiate DNA cleavage through a CrV—phosphate diester intermediate by hydrogen

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abstraction from Cl⁻. Some transition metals such as IrIV, MnO₂⁻, OsIII, and RuIII are reported to further oxidize 8-oxo-G, a major oxidation product of guanine in double-stranded DNA.²

Platinum coordination complexes are biologically important for their anticancer activities.³ PlatIV complexes bind preferentially to the N7 position of G.⁴ Cisplatin, cis-diaminedichloroplatinum(II), binds bifunctionally to adjacent Gs and kinks DNA.⁵ High mobility group proteins and p53 protein bind to the platinum-modified DNA, preventing DNA replication, and leading to cell death.⁶ Some PlatIV complexes have shown potential as powerful anticancer drugs, although most PlatIV anticancer drugs have been considered to be PlatII prodrugs.⁷ Because PlatIV complexes are kinetically inert,³ reaction with DNA is not expected until they have been reduced to PlatII in the cellular medium.¹² However, direct PlatIV binding to DNA has been observed in vitro in several labs.¹⁴⁻¹⁶ Choi et al. reported that PlatIV complexes with highly electron-withdrawing and bulky ligands have high reduction potentials¹⁵a and high reactivity with anticancer drugs have been considered to PlatII prodrugs. PlatIV complexes binding to DNA is not expected until they have been reduced to PlatII in the cellular medium.² However, direct PlatIV binding to DNA has been observed in vitro in several labs.¹⁴⁻¹⁶ Choi et al. reported that PlatIV complexes with highly electron-withdrawing and bulky ligands have high reduction potentials¹⁵a and high reactivity toward the monomeric nucleotide GMP.¹⁵b Furthermore, [PtIV, Cl₁(dach)], which has a high reduction potential, oxidizes guanine in GMP, dGMP, 2'-deoxyguanosinosine (dG), 2'-deoxyguanylyl(3'-5')-2'-deoxynosine [dGpG1], and a double-stranded oligonucleotide without any UV radiation or additional oxidants.¹⁵c The major oxidation product was identified as 8-oxo-G. We also observed that [PtIVCl₄(dach)] further oxidizes 8-oxo-G. It is important to study the mechanism in detail because the redox reaction between DNA and PlatIV complexes represents new chemistry. In this paper, we propose a detailed mechanism of the redox reaction between [PtIVCl₄(dach)] and dGMP based on UV-visible, HPLC, ¹H and 31P NMR, and ESI-MS studies. The mechanism involves PlatIV binding to N7 of dGMP followed by cyclization of a phosphate oxygen to C8 of dGMP. This study provides the most detailed insight to date of the mechanism of two-electron oxidation of guanine by a transition metal complex and provides structural characterization of a new cyclic nucleotide as the initial product.

Results

Identification of a PlatIV-dGMP Intermediate, I. The time course of the reaction of [PtIVCl₄(dach)]/dGMP (10 mM/20 mM) was monitored by ¹H NMR spectroscopy, and results paralleled those previously obtained with GMP.⁻¹⁸ Figure 1a shows the ¹H NMR spectra in the 9 ppm region. The multiplet peak at 9.3 ppm is assigned to H8 of PlatIV bound to N7 of dGMP.¹⁵b,¹⁶ The concentration of PlatIV—dGMP at various reaction times was calculated by multiplying 20 mM by the area under the 9.3 ppm peak relative to that at 8.3 ppm (H8 of free dGMP) at t = 0, and these are displayed in Figure 2 as a function of time. The concentration reaches a maximum at 4 h and gradually decreases, indicating that the PlatIV—dGMP adduct is an intermediate.

Intermediate I was also detected by HPLC (Figure 3). The HPLC chromatogram of the reaction at t = 0 shows two peaks with retention times of 4.6 and 6.2 min due to free dGMP and free [PtIVCl₄(dach)], respectively. After a 1-h reaction time (data not shown), the [PtIVCl₄(dach)]/dGMP (10 mM/20 mM) solution shows a new set of peaks with a retention time of 10.9 min. The UV/visible spectrum of this peak obtained directly from the HPLC eluate is different from that of free dGMP (Figure 4). It has a large UV absorption below 220 nm due to PlatIV and a λmax of 252 nm with a shoulder at 280 nm. In contrast, free dGMP has no absorption below 220 nm and a λmax of 254 nm with a shoulder at 272 nm. The intensity of this 10.9 min peak reaches a maximum around 4 h (Figure 3) and falls off afterward, which is the same trend as the peak at 9.2 ppm in ¹H NMR spectra (Figure 2). The UV absorption feature along with the ¹H NMR data support the proposal that I is the PlatIV—dGMP adduct.

Identification of a Guanosine-Cyclic Phosphodiester, II. Figure 1b displays the time course of the reaction of [PtIVCl₄(dach)]/dGMP (10 mM/20 mM) in the 6 ppm region. The peak at 6.46 ppm is assigned to H1' of free dGMP. There are three new peaks in this area upon reaction with PlatIV. The peak centered at 6.57 ppm grows in intensity, reaches a maximum, and gradually diminishes. The peaks at 6.36 and 6.42 ppm are assigned to the H1' of the final products, 8-oxo-dGMP and the PlatIV—dGMP adduct, respectively. The assignment was made by comparing the H1' region of ¹H NMR spectra of authentic 8-oxo-dGMP (6.36 ppm) and [PtIVCl₄(dach)]/dGMP) (6.42 ppm). The concentrations of the species giving the peak at 6.57 ppm at various reaction times were calculated by multiplying 20 mM.
by the area of the peak relative to that of 6.46 ppm at \( t = 0 \). These are overlaid with the concentration of \( \text{I} \) in Figure 2. The intermediate giving the 6.57 ppm peak reaches a maximum concentration of 3.5 mM after 10 h, while \( \text{I} \) reaches a maximum concentration of 1.9 mM after 4 h. Therefore we conclude that there is a second intermediate.

**Figure 1.** Time-dependent \(^1\text{H} \) NMR spectra of the reaction of 10 mM [Pt\(^{IV}\)Cl\(_4\)(dach)] with 20 mM dGMP at pH 8.6 (\( t = 0 \)) and 37 °C.

**Figure 2.** Time-dependent concentrations of \( \text{I} \) and \( \text{II} \) generated from the reaction of 10 mM [Pt\(^{IV}\)Cl\(_4\)(dach)] with 20 mM dGMP at pH 8.6 (\( t = 0 \)) and 37 °C. They were calculated from the \(^1\text{H} \) NMR peaks at 9.30 and 6.57 ppm, respectively.

**Figure 3.** Reversed-phase HPLC chromatograms (DAD, \( \lambda = 294 \text{ nm} \)) of the reaction of 10 mM [Pt\(^{IV}\)Cl\(_4\)(dach)] with 20 mM dGMP at pH 8.6 (\( t = 0 \)) and 37 °C.
Intermediate II was also detected by HPLC. A new peak with a retention time of 5.9 min is seen in the t = 4 and 12 h reaction chromatograms in Figure 3. The UV/visible spectrum of this peak obtained directly from the HPLC DAD has a λ_{max} at 252 and 280 nm and is different from those of free dGMP and I (Figure 4). The intensity of the 5.9 min peak reaches a maximum around 10 h and falls off afterward, which is the same trend as the peak at 6.57 ppm in the 1H NMR spectra (Figure 2).

A fraction containing II was collected directly from HPLC runs of 10-h reaction mixtures and analyzed by negative ion ESI-MS. The molecular ion of II occurs at m/z 344, which is 2 units lower than that of dGMP. The m/z of II was unchanged in a reaction conducted in 95% H2^{18}O vs H2^{16}O (vide infra).

The 31P NMR spectra of the reaction mixture at various times indicate the phosphorus atom in II is in a different environment than in dGMP (Figure 5). The peak at 5.0 ppm at t = 0 arises from 31P of free dGMP. It gradually shifts upfield as the reaction progresses, eventually appearing at 2.5 ppm at t = 16 h. Usually orthophosphate and nucleoside 5′- and 3′-phosphates appear in this chemical shift region, but this upfield shift is likely due to the lowering of the pH as the reaction proceeds. The pH of the reaction was adjusted to 8.6 at t = 0. It gradually drops to pH 4.0 at t = 16 h (Figure 6), which is due to the production of two protons in the two-electron redox reaction (vide infra).

It is known that both inorganic and nucleotide phosphates shift upfield at lower pH. The peak at 2.95 ppm on the shoulder of the 2.5 ppm peak on the t = 16 h spectrum is due to 8-oxo-dGMP, which was confirmed by the 31P NMR spectrum of authentic 8-oxo-dGMP. At t = 2 h (data not shown), a new peak at −5.5 ppm appears. It grows in intensity, reaches a maximum, and gradually diminishes as the reaction progresses. We believe that this peak is due to II. The resonance position of this peak does not change at different reaction times, indicating that it is pH-insensitive. It is known that the 31P NMR peak of cyclic phosphodiester appears upfield relative to that of phosphate and is pH-insensitive. For example, the 31P NMR peak of 3′,5′-cyclicAMP appears at −1 ppm compared to 2 ppm in AMP, and it is pH-insensitive.

Therefore, on the basis of UV, ESI-MS, and 31P NMR, we propose that the intermediate II is 8-hydroxyguanosine cyclic 5′,8-(hydrogen phosphate). It has the characteristic UV and 31P NMR spectra of 5′,8-cyclo-guanosine and two fewer hydrogens (2 amu) than dGMP.

Identification of the Final Oxidation Product of dGMP. The HPLC fraction containing the 6.2 min (Figure 3) or 9 min (Figure 7) peak was collected. Its UV spectrum shows λ_{max} at 252 and 280 nm and is different from those of free dGMP and I (Figure 4). The intensity of the 5.9 min peak reaches a maximum around 10 h and falls off afterward, which is the same trend as the peak at 6.57 ppm in the 1H NMR spectra (Figure 2).

Figure 4. UV spectra of dGMP, I, II, and 8-oxo-dGMP obtained directly from the diode array detector of the HPLC. The retention times of the intermediate correspond to the chromatograms in Figure 3.

Figure 5. Time-dependent 31P NMR spectra of the reaction of 10 mM [PtIVCl4(dach)] with 20 mM dGMP at pH 8.6 (t = 0) and 37 °C.

Figure 6. pH and [Cl−] vs time for the reaction of 5 mM [PtIVCl4(dach)] with 5 mM dGMP at pH 8.6 (t = 0) and 37 °C.

Two-Electron Oxidation of dGMP by a Pt(IV) Complex

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Figure 1. HPLC chromatograms (DAD λ = 280 nm) of the reaction of 10 mM [PtIVCl4(dach)] with 20 mM dGMP at pH 8.6 (t = 0) and 37 °C.

Figure 2. Positive-ion ESI-MS/MS analysis of fragment ions of 8-oxo-dGMP produced from the reaction of 10 mM [PtIVCl4(dach)] with 20 mM dGMP at pH 8.6 (t = 0) and 37 °C after 12 h. Black, reaction conducted in H216O; red, reaction conducted in H218O.

247 and 290 nm (Figure 4), coinciding with the literature values for 8-oxo-G.19 Fragmentation with positive-ion ESI-MS/MS displayed an m/z = 364 (Figure 8), the same m/z as 8-oxo-dGMP that is 16 units higher than dGMP (m/z 348). Furthermore, the 1H (6.37 ppm) and 31P NMR (3.0 ppm) spectra of this fraction were exactly the same as those of authentic 8-oxo-dGMP. Therefore, we conclude that the final oxidation product of dGMP is 8-oxo-dGMP.

Origin and Location of the Added Oxygen in 8-Oxo-dGMP. Experiments conducted under nitrogen in the absence of O2 also produced 8-oxo-dGMP. When 95% H218O was used as solvent, the molecular ion of 8-oxo-dGMP was 2 amu higher (366 amu) than that from the reaction in H216O (Figure 8). This indicates that the origin of the oxygen atom gained in the formation of 8-oxo-dGMP is solvent water.

The location of the added oxygen from water could be deduced by comparing the positive-ion ESI-MS/MS spectra of the fragments of 8-oxo-dGMP from the reaction in H216O to those in H218O. Only 1% of the fragments of m/z = 168 (assigned as 7,8-dihydro-8-oxoguanine) is formed from 8-oxo-dGMP (m/z = 166 amu) than that from the reaction in H216O (Figure 8). This indicates that the added oxygen has not been incorporated into the glycosidic bond. The HPLC chromatogram (Figure 3) of the 4 h reaction mixture at pH 5.8 was analyzed immediately by HPLC. The region around 8, 6, and 2 ppm of the final reaction mixture display the same peaks at 8.65, 6.42, and 2.23 ppm as authentic 8-oxo-dGMP. Therefore, we conclude that the new oxygen atom gained from water resides on the phosphate group of 8-oxo-dGMP. However, a minor pathway (±1%) involves direct incorporation of 18O into 8-oxo-dGMP from solvent.

Identification of [PtIIICl2(dach)] and [PtIII(dach)(dGMP)]. The final reduction products of [PtIVCl4(dach)] consist of [PtIICl2(dach)] and [PtIII(dach)(dGMP)]. Since the solubility of [PtIIICl2(dach)] is very low (~1 mM), most of the [PtIICl2(dach)] precipitates as orange crystals. The FTIR spectrum of the orange crystals is identical to that of an authentic sample of [PtIIICl2(dach)].15c The existence of [PtIII(dach)(dGMP)] in solution was shown by 1H NMR and HPLC. The 1H NMR spectra of the regions around 8, 6, and 2 ppm of the final reaction mixture display the same peaks at 8.65 and 2.23 ppm (Figure 9) as those of authentic 8-oxo-dGMP. The peak at 6.42 ppm (Figure 1b) is due to H1 of PtII(dGMP) adduct. The HPLC chromatogram (Figure 3) of the reaction mixture of [PtIVCl4(dach)]/dGMP (10 mM/20 mM) at t = 24 h shows the peak at 7.5 min that is due to [PtIII(dach)(dGMP)].

Stoichiometry of the Reaction. Figure 6 displays the pH and the concentration of chloride for the reaction of [PtIVCl4(dach)] with dGMP (both 5 mM) as a function of time. It clearly shows the production of both H+ and Cl− during the course of the reaction. The concentration of dGMP at various time intervals was obtained by comparison of calibration curves of authentic dGMP with HPLC chromatograms obtained with DAD monitoring at 280 nm. Because of the hydrolysis of [PtIVCl4(dach)], it was difficult to quantify [PtIVCl4(dach)] by HPLC. However, the chemical shifts of the 1H NMR peaks (3.31, 2.42, 1.82, and 1.43 ppm) of aqueous [PtIVCl4(dach)] stay the same over several days. Since the peak at 2.42 ppm was sensitive to the oxidation state of platinum [2.42 ppm for PtIV and 2.23 ppm for PtII], we used the relative intensity of this peak compared to the internal standard, 3-(trimethylsilyl)propionic acid, to calculate the concentration of the unreacted [PtIVCl4(dach)].

The concentration of 8-oxo-dGMP was obtained by comparison of a calibration curve of authentic 8-oxo-dGMP with HPLC chromatograms obtained with DAD at 280 nm. To obtain the overall total concentration of 8-oxo-dGMP generated from the reactants, the solution was acidified to pH 4 in order to quickly convert all of II to 8-oxo-dGMP. When the 4 h reaction mixture at pH 5.8 was analyzed immediately by HPLC, II appeared at 8.1 min and then disappeared in the acidified solution (pH 4.0) (Figure 7). The intensity of the 9.0 min peak in the pH 4.0 solution increases compared to that of the pH 5.8 solution, confirming that all of II was converted to 8-oxo-dGMP. We obtained the total concentration of 8-oxo-dGMP from the intensity of the 9.0 min peak of the acidified solution. The concentrations of reacted dGMP, [PtIVCl4(dach)], and total 8-oxo-dGMP are displayed in Figure 10.

The concentration of reacted dGMP and [PtIVCl4(dach)] are the same within experimental error, indicating that 1 equiv of dGMP reacts with 1 equiv of [PtIVCl4(dach)]. The concentration-time profile of 8-oxo-dGMP is symmetrical to those of reactants, indicating that 1 equiv of [PtIVCl4(dach)] converts 1 equiv of dGMP to 1 equiv of 8-oxo-dGMP. For the 10-h reaction mixture of 10 mM [PtIVCl4(dach)] and 20 mM dGMP (pH 8.6 shifted to 199 when the reaction was conducted in H218O. Therefore, we conclude that the new oxygen atom gained from water resides on the phosphate group of 8-oxo-dGMP. However, a minor pathway (±1%) involves direct incorporation of 18O into 8-oxo-dGMP from solvent.
at $t = 0$), approximately 5 mM $\left[\text{Pt}^{\text{IV}} \text{Cl}_4 (\text{dach})\right]$ and dGMP were consumed to produce 5 mM 8-oxo-dGMP. Therefore, the overall reaction can be written as

$$\left[\text{Pt}^{\text{IV}} \text{Cl}_4 (\text{dach})\right] + \text{dGMP} + \text{H}_2\text{O} \rightarrow \left[\text{Pt}^{\text{II}} \text{Cl}_2 (\text{dach})\right] + 8\text{-oxo-dGMP} + 2\text{H}^+ + 2\text{Cl}^- \quad (1)$$

For the reaction of equimolar $\left[\text{Pt}^{\text{IV}} \text{Cl}_4 (\text{dach})\right]$ and dGMP (10 mM), a small amount (~5% relative to 8-oxo-dGMP) of guanidinohydantoin was observed by negative-ion LC-ESI-MS ($m/z = 352$, or +6 relative to dGMP) after 20 h. Guanidinohydantoin is a further oxidation product of 8-oxo-dGMP by $\left[\text{Pt}^{\text{IV}} \text{Cl}_4 (\text{dach})\right]$.

Discussion

According to eq 1, the overall reaction generates two protons, one 8-oxo-dGMP, and two chlorides when one Pt IV and one dGMP are allowed to react. To our knowledge, the present study reports the only system in which 8-oxo-G is generated cleanly and stoichiometrically from the transition metal oxidation of a mononucleotide. One-electron photooxidation of the dG nucleoside generates imidazolone and its hydrolysis product oxazolone as major products.20 Two-electron oxidation of dG $^{4a,b}$ and the dinucleotide d(GpT) $^{4c}$ by Mn V produces imidazolone and dehydroguanidinohydantoin, respectively. Although it is reported that Ru IV complexes oxidize guanine via an inner-sphere, oxygen atom transfer to produce 8-oxo-G, the detailed mechanism and stoichiometry have not been reported.

The HPLC, UV-visible, 1H and 31P NMR, and ESI-MS studies lead us to propose the mechanism in Scheme 1. The first step is loss of Cl$^-$ from $\left[\text{Pt}^{\text{IV}} \text{Cl}_4 (\text{dach})\right]$, followed by binding to G N7. In a previous study, we reported that the decay of $\left[\text{Pt}^{\text{IV}} \text{Cl}_4 (\text{dach})\right]$ is sigmoidal in shape, indicating that the reaction is autocatalyzed. Although we show the axial chloride substituted with dGMP, it is also possible that the equatorial chloride

could be replaced. The next step involves nucleophilic attack at C8 by an oxygen anion of the phosphate group, producing a cyclic phosphodiester. The next step is an inner-sphere, two-electron transfer from bound dGMP to PtIV. At this point, the H8 proton and one more chloride are released to form the 8-hydroxyguanosine 5′,8 cyclic phosphodiester and [Pt II Cl2-(dach)]. The cyclic phosphodiester slowly hydrolyzes to form 8-oxo-dGMP, a process that is faster at lower pH. Indeed, higher quantities of intermediate II, now assigned as the cyclic phosphodiester, could be isolated by HPLC if the pH was kept at 7 or above. The [PtII Cl2(dach)] produced can also bind to another dGMP to form [Pt II Cl(dach)(dGMP)] with release of a chloride ion.

This 2e−, 2H+ oxidation process contrasts with the mechanism of autooxidation of a series of 6-oxopurines by [RuIII(NH3)5] complex in basic media. The RuII complex binds to N7 of inosine, leading to deprotonation of H8 and transfer of 1e− to RuIII, and generates a radical at C8. This radical is attacked by water and undergoes a second 1e−, 1H+ oxidation to produce 8-ketonucleosides.

The formation of a cyclic 5′,8-phosphodiester upon purine oxidation is a novel finding, although it is related to the observation of cyclic ethers formed in guanosine oxidation by nucleophilic attack of the 5′-hydroxyl at C8. In the case of cyclic guanosine ethers, the product is stable toward further hydrolysis, while the cyclic phosphodiester obtained from PtIV oxidation of dGMP hydrolyzes over the course of several hours. The instability of the cyclic phosphodiester, compared to a 3′,5′-phosphodiester linkage in DNA, could be due to ring strain in the nine-membered ring as well as the greater leaving group ability of 8-hydroxyguanine, which may be similar to a phenolate.

That a phosphate oxygen serves as the nucleophile suggests that the ring strain cannot be terribly high, since the high concentration of solvent would otherwise compete. If H2O were the attacking nucleophile, 8-oxo-dGMP would be formed directly at this stage, and use of H218O would lead to incorporation of the isotopically labeled oxygen at C8 of the base. In fact, careful analysis of the isotope ratios in the base-containing fragments (e.g., m/z = 168) from the ESI-MS/MS study of 8-oxo-dGMP formed in H218O indicate that about 1% of the product derives from direct attack of water on the platinated intermediate rather than phosphodiester hydrolysis. This indicates that the intramolecular phosphate oxyanion is by far the best available nucleophile for the reaction of the mononucleotide. On the other hand, it is not known if a phosphodiester anion could play this role during PtIV oxidation of double-stranded DNA due to conformational restraints and the reduced nucleophilicity of a phosphodiester anion. However, other nucleophiles may be associated with duplex DNA in the cellular milieu, including polyamines and proteins containing nucleophilic side chains. For those cases where the PtIV complex has a sufficient intracellular concentration and lifetime, these studies suggest that PtIV oxidation of guanine in DNA could potentially be responsible for both mutagenic lesions such as 8-oxo-dG as well as formation of adducts or DNA–protein cross-links.

In summary, we have demonstrated that [PtIVCl2(dach)] binds to N7 of guanine followed by an inner-sphere, two-electron redox reaction to form a novel intermediate, 8-hydroxyguanosine cyclic 5′,8-(hydrogen phosphate), that slowly hydrolyzes to 8-oxo-dGMP. The biological implications of this mechanism and its applicability to dG in oligomers are currently under investigation.
Experimental Section

**Materials.** [PtIVCl4(dach)] (tetraplatin) was obtained from the National Cancer Institute, Drug Synthesis and Chemistry Branch, Developmental Therapeutics Program, Division of Cancer Treatment. 8-oxo-dGMP was a gift from Professor Wallace at the University of Vermont.

**Reaction of [PtIVCl4(dach)] with dGMP.** Buffers were not used in order to avoid complications arising from buffer coordination to platinum.21 Stock solutions (11 mM) of [PtIVCl4(dach)] was prepared by dissolving 11 µmol of the PtIV compound in 1 mL of H2O in an amber vial. The solubility of [PtIVCl4(dach)] is low, so several heating (maximum 50 °C) and vortexing cycles over 5 or 6 h were necessary for complete dissolution. After the PtIV compound was dissolved, an appropriate amount of dGMP stock solution (100 mM) was added to give the desired final concentration. The pH of the solution was adjusted to pH 8.6 with 0.1 M NaOH. The time when the pH was adjusted to pH 8.6 was recorded as the beginning of the reaction. All the samples for NMR analysis were prepared in D2O (>98%). Most reaction solutions were filtered through a syringe-end filter disk (Gelman Acrodisk 0.45 µm pore size) before HPLC and NMR analysis.

**Preparation of [PtIICl(dach)(dGMP)].** For reference purposes, [PtIV-Cl(dach)(dGMP)] was prepared by dissolving an appropriate amount of dGMP to give a final concentration of 1 mM dGMP in the 1 mM [PtIVCl4(dach)]. It is noted that [PtIV(dach)(dGMP)] was not produced even in the presence of excess (100-fold) dGMP for up to 3 days. [PtIV-Cl2(dach)] was obtained by reducing [PtIVCl4(dach)] with 10-fold excess of ascorbic acid.22 The reaction was allowed to proceed for 2 h, after which time a bright yellow-orange powder was obtained: IR (3267, 3190, 3066, 2936, 2863, 1564 cm−1).

**pH and Chloride Concentration.** pH and chloride concentration were measured with pH and chloride (Orion Model 94-17B) electrodes, respectively, connected to a pH meter (Orion Research 960).

**HPLC.** HPLC measurements were performed on either a Waters or Beckman liquid chromatograph equipped with a diode-array detector. Isocratic (1 mL/min) conditions with a Waters symmetry shield RP18 (5 µm, 4.6 mm × 250 mm) column and solvent consisting of 95% triethylammonium acetate (50 mM, pH 7.0) and 5% methanol were the best for the elution of 1 and [PtIVCl(dach)(dGMP)] at reasonable retention times (10.9 and 7.5 min, respectively). Isocratic conditions (1 mL/min) with a RESTEK Ultra IBD (5 µm, 4.6 mm × 250 mm) column and solvent consisting of 98% ammonium acetate (20 mM, pH 5.8) and 2% acetonitrile gave the best resolution for the components in the reaction mixture (Figure 7). However, one drawback for these conditions was that the retention times of 1 and [PtIVCl(dach)(dGMP)] were long (>40 min).

**NMR.** NMR spectra were recorded on a Bruker NMR spectrometer equipped with a broad-band inverse tunable probe operating at 400.13 MHz for 1H and 161.97 MHz for 31P. Chemical shifts for 1H and 31P are reported relative to tetramethylsilane (TMS) and 85% H3PO4, respectively, each at 0.00 ppm. Samples were prepared in D2O and the residual water signal was further suppressed by the watergate pulse sequence.23 Time-course experiments were performed at 37 °C by recording spectra at 1 h intervals by use of a multiple acquisition program (total acquisition time + delay between experiments = 1 h) with 256 scans for 1H or 1000 scans for 31P with broad-band 1H decoupling.

**ESI-MS.** Labeled and unlabeled samples were analyzed by positive-ion electrospray ionization (ESI) on a Micromass Quattro II tandem mass spectrometer equipped with a Zspray API source. Samples were dissolved in acetonitrile and water (1:1) and introduced via infusion at a flow rate of 0.30 mL/h. The source and desolvation temperatures were 80 and 120 °C, respectively. The capillary voltage was set to 3.1 kV, sampling cone voltage to 58 V, and the extractor cone to 3 V. The collision energy was set to 14 eV. Argon, used as a collision gas for the CID experiments, was adjusted to a pressure of 1.7 × 10−4 mBar. In a representative experiment, MS-1 was set to the mass for the nucleotide and the precursor ion was subjected to CID in the static quadrupole. The resulting spectrum of the products recorded by scanning the second scanning analyzer (MS-2) between 50 and 370 Da. The scan duration and the interscan delay time were 3.0 and 0.1 s, respectively. The instrument was operated and data were accumulated with Micromass Masslynx software (version 3.2).

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