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Formation of DNA–Protein Cross-Links Between γ-Hydroxypropanodeoxyguanosine and EcoRI

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The toxicity of acrolein, an α , β -unsaturated aldehyde produced during lipid peroxidation, is attributable to its high reactivity toward DNA and cellular proteins. The major acrolein–DNA adduct, γ -hydroxypropano-2'-deoxyguanosine (γ -HOPdG), ring opens to form a reactive N^2 -oxopropyl moiety that crosslinks to DNA and proteins. We demonstrate the ability of γ -HOPdG in a duplex oligonucleotide to crosslink to a protein (EcoRI) that specifically interacts with DNA at a unique sequence. The formation of a cross-link to EcoRI was dependent on the intimate binding of the enzyme to its γ -HOPdG-modified recognition site. Interestingly, the cross-link did not restrict the ability of EcoRI to cleave DNA substrates. However, stabilization of the cross-link by reduction of the Schiff base linkage resulted in loss of enzyme activity. This work indicates that the γ -HOPdG–EcoRI cross-link is in equilibrium with free oligonucleotide and enzyme. Reversal of cross-link formation allows EcoRI to effect enzymatic cleavage of competitor oligonucleotides.

Introduction

Noncovalent interactions between DNA and proteins are essential for proper cellular function. The recognition of specific DNA sequences or structures by DNA-binding proteins is a crucial regulatory step during DNA replication, control of gene expression, and the response to and repair of DNA damage. Disruption of these interactions poses a significant threat to genome integrity and cell viability. Protein and DNA damage induced by endogenous electrophiles is associated with a number of human diseases, including cancer, cardiovascular disease, and neurological disorders (1-3). For example, oxidation products of polyunsaturated fatty acids comprise a series of cytotoxic α,β -unsaturated aldehydes, including malondialdehyde, 4-hydroxynonenal, and acrolein, that readily form covalent adducts with cellular macromolecules, contributing to their toxicity (reviewed in ref 4).

Acrolein is a mutagen and a tumor initiator produced endogenously via lipid peroxidation and myeloperoxidasecatalyzed amino acid oxidation (5-8). It is also an ubiquitous pollutant found in automobile exhaust and cigarette smoke (9). Acrolein exhibits facile reactivity with proteins (8) and represents a significant source of endogenous DNA damage by forming cyclic adducts with DNA bases and DNA-protein cross-links (10, 11). The major products of the reaction of acrolein with DNA are exocyclic adducts of deoxyguanosine, (8R/S)-3-(2'-deoxyribos-1'-yl)-5,6,7,8-tetrahydro-8-



Figure 1. Structures of DNA adducts and their ring-opening products.

hydroxypyrimido[1,2*a*]purin-10(3*H*)-one (γ -HOPdG)¹ and (6*R/S*)-3-(2'-deoxyribos-1'-yl)-5,6,7,8-tetrahydro-6-hydroxypyrimido[1,2*a*]purin-10(3*H*)-one (α -HOPdG) (10) (Figure 1). α -HOPdG is more mutagenic than γ -HOPdG and induces G \rightarrow T transversions in site-specific mutagenesis experiments (12). Both γ -HOPdG and α -HOPdG are present in human lung DNA at levels in excess of those typically reported for polycyclic aromatic hydrocarbon–DNA adducts (13).

While γ -HOPdG contributes to acrolein toxicity by directly affecting DNA replication (*12, 14*), the adduct itself is a reactive molecule. In duplex DNA, γ -HOPdG undergoes a spontaneous

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¹ Abbreviations: BSA, bovine serum albumin; EDTA, ethylenediaminetetraacetic acid; γ-HOPdG, γ-hydroxy-2'-propanodeoxyguanosine; M₁dG, pyrimido[1,2- α]purin-10(3*H*)-one; MOPS, 4-morpholinepropanesulfonic acid; MS, mass spectrometry; NER, nucleotide excision repair; SDS-PAGE, sodium dodecyl sulfate-polyacrylamide gel electrophoresis.

ring-opening reaction to generate N^2 -oxopropyl-2'-deoxyguanosine that can react with primary amines on cellular macromolecules (15). γ -HOPdG positioned within a Cp*G sequence context in duplex DNA undergoes ring opening and subsequent reaction with a guanine base in the complementary strand to form interstrand DNA-DNA cross-links (16). The ring-opened γ -HOPdG adduct is also reactive toward amino groups of peptides and is capable of forming reversible Schiff basemediated cross-links to proteins (12).

A role for DNA—protein cross-links during acrolein toxicity has not been fully elucidated. The interaction of adducted oligonucleotides with restriction endonucleases represents a useful model system with which to evaluate the impact of DNA damage on recognition and catalysis of DNA-binding proteins. Previously, our group showed that exocyclic adducts in an EcoRI recognition site blocked binding and cleavage of DNA by that enzyme, although ring-opened adducts were substrates for endonucleolytic cleavage by EcoRI (*17*). Here, we examine the fate of γ -HOPdG and consequently its ring-opened analogue, when incorporated into duplex substrates of EcoRI. We describe the ability of γ -HOPdG to cross-link to EcoRI and affect enzyme activity. The results suggest that the DNA—protein cross-link is unstable and either undergoes hydrolysis to free oligonucleotide and enzyme or can be reduced to a stable, irreversible cross-link.

Experimental Procedures

Reagents. T4 polynucleotide kinase and PvuII were obtained from New England Biolabs (Beverly, MA). EcoRI was purified as described (*18*). EcoRI for some cross-linking experiments was purchased from MBI Fermenetas (Hanover, MD). NaC-NBH₃ and NaBH₄ were purchased from Sigma (St. Louis, MO). $[\gamma^{-32}P]$ -ATP was from ICN Biomedicals, Inc. (Costa Mesa, CA).

Oligonucleotides. The 21-mer oligonucleotides contained a unique EcoRI restriction site (5'-GAATTC-3') in the sequence 5'-TATCATGTCTNAATTCCCGGT-3', where N = dG, γ -HOPdG, or pyrimido[1,2- α]purin-10(3*H*)-one (M₁dG). The sequence of the complementary 27-mer oligonucleotide was 5'-AGAC-CGGGAATTCAGACATGATACGGA-3'. The scrambled EcoRI sequence was 5'-TATCATGTCTNTATACCCGGT-3', where N = γ -HOPdG; it was annealed to its complementary sequence. γ -HOPdG-modified oligonucleotides were synthesized and purified as previously described (*19*). The M₁dG-modified oligonucleotide was synthesized using the method previously described (*20*) and purified by polyacrylamide gel electrophoresis (*21*). Unmodified 21-mer and 27-mer oligonucleotides were synthesized by Integrated DNA Technologies (Coralville, IA).

Trapping of Covalent DNA-Protein Complexes Using NaCNBH₃. Modified and unmodified 21-mer oligonucleotides were 5'-end-labeled in a phosphorylation buffer of 50 mM 4-morpholinepropanesulfonic acid (MOPS) (pH 7.3), 10 mM MgCl₂, 100 µM EDTA, and 5 mM dithiothreitol containing 5 μ M oligonucleotide, 0.5 mM [γ^{32} P]-ATP, and 10 units of T4 polynucleotide kinase. Oligonucleotides were purified from free $[\gamma^{32}P]$ -ATP by passage through Bio-Spin P-6 columns (Bio-Rad, Hercules, CA). For restriction endonuclease studies, substrates were generated by annealing radiolabeled γ -HOPdGor M₁dG-modified 21-mer (2 pmol) to a complementary 27mer (10 pmol) in 50 mM MOPS (pH 7.4) and 100 mM NaCl (50 μ L total volume) by heating to 90 °C for 3 min, followed by slow cooling to room temperature. DNA-protein cross-links were detected essentially as described with slight modifications (12). ³²P-labeled substrate (160 pM) was incubated with EcoRI or PvuII restriction enzyme (1.6 nM) or bovine serum albumin (BSA) (5000 nM) in 50 mM MOPS (pH 7.4), 10 mM NaCl, and 0.1 mM ethylenediaminetetraacetic acid (EDTA), at room temperature for 2 h. NaCNBH₃ (50 mM) was added immediately following the addition of enzyme. Because Mg^{2+} is required for EcoRI cleavage activity but not required for the specific binding of enzyme to DNA, Mg^{2+} was omitted from the incubation. The reactions were terminated by addition of Laemmli sample buffer, then boiled for 5 min, and separated by 10% sodium dodecyl sulfate—polyacrylamide gel electrophoresis (SDS-PAGE) (22). Cross-linked bands were visualized using a FujiFilm FLA-5000 phosphorimager analyzer.

Restriction Digestion of Adducted Oligonucleotide Substrates. Radiolabeled substrates were generated by annealing adducted or unmodified 5'-32P-labeled 21-mer oligonucleotide with an equimolar amount of 5'-³²P-labeled 27-mer complement. Annealing of 2.5 pmol of each oligonucleotide was performed in buffer (50 mM MOPS, pH 7.4, and 100 mM NaCl) by heating to 90 °C for 3 min, followed by slow cooling. Nonradiolabeled substrates were generated by annealing adducted or unmodified 21-mer with complementary oligonucleotide under the conditions described above. Restriction digests (10 μ L) contained 5 nM ³²P-labeled substrate, 495 nM unlabeled substrate, 50 mM MOPS (pH 7.9), 10 mM MgCl₂, 100 mM NaCl, and 5 nM EcoRI restriction endonuclease. The reactions were carried out for specified times at 37 °C, then quenched by the addition of 10 mM EDTA in 90% formamide. The products of the reaction were resolved on a 20% denaturing gel using the Ultrapure Sequagel system (National Diagnostics, Atlanta, GA). The positions of the bands were visualized using a FujiFilm FLA-5000 phosphorimager analyzer. Cleavage of the EcoRI substrate generated two 5'-32P-labeled products of different sizes. The 21-mer oligonucleotide 5'-cleavage product was an 11-mer, whereas the 27-mer substrate's 5'-cleavage product was an 8-mer. Thus, by monitoring the appearance of the 11-mer and the 8-mer, cleavage of both strands of the substrate duplex could be monitored simultaneously. The extent of cleavage was determined by integrating the radioactivity of each band, then dividing the radioactivity of the band corresponding to cleavage product by the sum of the radioactivity of cleaved and uncleaved oligonucleotide to give the percent cleavage.

In some experiments, incubation mixtures were prepared in which 500 nM unlabeled adducted substrate or unmodified substrate was incubated for 2 h with 5 nM EcoRI. Then, 500 nM unmodified substrate, spiked with ³²P-labeled unmodified substrate, was added to the reaction mixture, and cleavage was monitored for 2 h. For other experiments, reactions were prepared under noncleavage conditions, that is, in the absence of Mg²⁺. γ -HOPdG-modified substrate or unmodified substrate (comprising 5 nM ³²P-labeled DNA and 495 nM nonradiolabeled DNA) was incubated in Mg²⁺-free buffer (50 mM MOPS, pH 7.9, and 100 mM NaCl) for 2 h with EcoRI (5 nM) in the presence or absence of NaCNBH₃. Following preincubation, 10 mM MgCl₂ was added to the reaction mixture, and cleavage was monitored for 1 h.

Results

Detection of a γ -HOPdG-Mediated Cross-Link to the Restriction Endonuclease, EcoRI. EcoRI binds the palindromic sequence, 5'-GAATTC-3', and sequentially cleaves both strands of the DNA duplex at the phosphodiester bond positioned 3' to deoxyguanosine. ³²P-labeled oligonucleotide duplexes were prepared containing a unique EcoRI recognition site that was modified with γ -HOPdG. Substrate DNA was incubated for 2 h with EcoRI (molar ratio of DNA duplex: EcoRI = 1:10) in the presence of NaCNBH₃, as described in the Experimental DNA-EcoRI Cross-Link Mediated by an Acrolein-DNA Adduct



Figure 2. γ -HOPdG forms a cross-link to EcoRI. ³²P-labeled γ -HOPdG- or M₁dG-modified or unmodified DNA substrates (160 pM) were incubated with EcoRI (1.6 nM) at room temperature for 2 h in the presence or absence of 50 mM NaCNBH₃. The products were resolved by SDS-PAGE and visualized by phosphorimager analysis. This is a representative autoradiogram from at least five independent experiments. The cross-linked product is indicated by the arrow.

Procedures. NaCNBH₃ does not react with EcoRI or γ -HOPdG but is capable of reducing an imine cross-link between the free aldehyde form of γ -HOPdG and a lysine residue. EcoRI reacted with γ -HOPdG-modified substrates to form a DNA-protein cross-link observable by SDS-PAGE analysis (Figure 2). Formation of the gel-shifted band corresponding to the protein-DNA complex was dependent on stabilization of the cross-link by reduction with NaCNBH₃. A weaker, second band was observed that migrated more slowly than the major γ -HOPdG-EcoRI band. The identity of this second band is unknown.

 γ -HOPdG ring opens to form an aldehydic moiety that is capable of reacting with proteins (15). The malondialdehyde– deoxyguanosine adduct, M₁dG, similarly ring opens in duplex DNA to form a potentially reactive oxopropenyl moiety, N^2 oxopropenyl-dG (23). The presence of M₁dG in the EcoRI recognition site does not prevent recognition of the site by EcoRI (17). Therefore, we tested whether M₁dG is also able to form cross-links to EcoRI. In contrast to γ -HOPdG, no cross-link was observed when M₁dG-modified substrates were incubated with EcoRI with or without NaCNBH₃ reduction (Figure 2).

Specificity of γ -HOPdG–EcoRI Cross-Link Formation. DNA substrates were prepared in which γ -HOPdG was positioned in a scrambled EcoRI recognition site (5'-GTATAC-3'). No EcoRI–DNA cross-link was observed when γ -HOPdG was in the scrambled sequence (Figure 3). When EcoRI was inactivated by heat (65 °C for 20 min) prior to the reaction with γ -HOPdG-modified substrate, only an extremely faint band at the position of the DNA-protein cross-link was detected after 2 h (Figure 4). Furthermore, γ -HOPdG did not cross-link to a restriction enzyme, PvuII, that does not specifically recognize the sequence of DNA where the adduct was positioned. Also, γ -HOPdG did not form cross-links with BSA, a common stabilizing agent added to commercial preparations of EcoRI, even at a DNA:BSA molar ratio = 1:5000 (data not shown). These results indicate that the cross-link formed between γ -HOPdG and EcoRI is most likely formed as a result of specific binding of the EcoRI recognition sequence and probably involves amino acids in close proximity to the adduct in the DNA-protein complex.

Time Course of γ **-HOPdG**–**EcoRI Cross-Link Formation.** Cross-link formation was monitored over 5 h (Figure 5). Reactions were prepared in the presence of NaCNBH₃, and then, a second reducing agent, NaBH₄, was added to rapidly quench



Figure 3. EcoRI cross-links to γ -HOPdG positioned within the EcoRI restriction site. ³²P-labeled γ -HOPdG-modified or unmodified DNA substrates (160 pM) were incubated with EcoRI (1.6 nM) at room temperature for 2 h in the presence or absence of NaCNBH₃. γ -HOPdG was positioned in either the EcoRI restriction site (GAATTC) or a scrambled sequence (GTATAC). The products were resolved by SDS-PAGE. A representative autoradiogram of the gel is shown.



Figure 4. Requirement of properly folded enzyme for cross-link formation. EcoRI (1.6 nM), native or denatured at 65 °C for 20 min, was incubated with γ -HOPdG-modified ³²P-end-labeled DNA substrate (160 pM) at room temperature for 2 h. NaCNBH₃ (50 mM) was added to the reaction to stabilize the Schiff base linkage between protein and DNA. The products were separated by 10% SDS-PAGE.



Figure 5. Time course of γ -HOPdG cross-linking to EcoRI. γ -HOPdGmodified ³²P-end-labeled DNA substrate (160 pM) was incubated with EcoRI (1.6 nM) for up to 5 h at room temperature in the presence of 50 mM NaCNBH₃, and then, a second reducing agent, NaBH₄ (100 mM), was added to rapidly quench any remaining aldehydic substrate at specific time points. A representative autoradiogram from three independent experiments is presented, which displays the accumulation of the DNA-protein cross-linked band. Percent cross-linking is shown in the graph just below the gel expansion. The values are the means \pm standard deviations from three independent experiments.

any remaining aldehydic substrate at specific time points. The γ -HOPdG-EcoRI cross-link was observed within 10 min. The conversion of free DNA to cross-linked product reached a plateau within 3 h.

In Vitro Cleavage of γ -HOPdG-Modified DNA Substrate by EcoRI Restriction Endonuclease. Oligonucleotide duplexes containing a single adduct located within a unique EcoRI recognition site were 5'-³²P-end-labeled on both strands. The



Figure 6. Effect of γ -HOPdG on EcoRI activity. (A) 5'-Radiolabeled duplex containing a single EcoRI recognition site. The arrows depict cleavage sites. γ -HOPdG was contained on the 21-mer strand, as indicated by "N". (B) Cleavage reactions containing 5 nM ³²P-labeled oligonucleotide duplex, 495 nM nonlabeled oligonucleotide substrate, and 5 nM EcoRI were incubated at 37 °C for various times, then quenched, and subjected to gel electrophoresis and phosphorimager analysis as described in the Experimental Procedures. Unmodified substrate, 27-mer strand (**I**); γ -HOPdG-modified substrate, 21-mer strand (**A**). (C) Nonlabeled oligonucleotide duplexes modified with γ -HOPdG or unmodified duplexes (500 nM) were incubated with EcoRI endonuclease (5 nM) for 2 h at 37 °C, prior to the addition of fresh unmodified EcoRI substrate, spiked with ³²P-labeled unmodified substrate, to a final concentration of 500 nM. Cleavage of the radiolabeled substrate was monitored for 2 h. The data shown represent cleavage of the 27-mer strand of the radiolabeled duplex in the presence of the indicated competitor substrate. The 21-mer strand of each duplex was cleaved similarly to its complement (data not shown). Unmodified competitor substrate (**I**); γ -HOPdG-modified competitor substrate (**I**). The values are the means \pm standard deviations from three independent experiments.

adducted strand of the duplex was a different length than its complement, and the oligonucleotides were designed to release ³²P-end-labeled cleavage products of different lengths. Thus, simultaneous monitoring of cleavage of both strands of the duplex was achieved. The complement strand of the γ -HOPdG-modified substrate was cleaved nearly as efficiently as unmodified substrate (~90%) (Figure 6B). In contrast, only ~20% of the γ -HOPdG-modified strand was cleaved by EcoRI within 2 h.

To examine whether the reduced cleavage efficiency of modified substrate was due to a sequestration or inactivation of EcoRI by γ -HOPdG, nonradiolabeled γ -HOPdG-adducted substrates or unadducted substrates were incubated with EcoRI for 2 h. Then, ³²P-labeled unmodified substrate was added to the reaction mixture, and cleavage was monitored. The radiolabeled unmodified substrate was cleaved to a similar extent, regardless of whether unmodified substrate or γ -HOPdG-modified substrate was initially incubated with the enzyme (Figure 6C). Thus, EcoRI retained complete activity following exposure to the DNA adduct.

To examine whether a stabilized cross-link could affect EcoRI activity, γ -HOPdG-modified or unmodified substrates were incubated with EcoRI in the absence of Mg²⁺ and either the presence or the absence of NaCNBH₃ for 2 h; then, MgCl₂ was added to the reaction mixture to initiate cleavage. Following MgCl₂ addition, EcoRI cleavage of the substrate was substantially decreased in samples containing γ -HOPdG-modified DNA and NaCNBH₃ as compared to samples containing γ -HOPdG-modified DNA substrate in the absence of reducing agent. There was no effect of NaCNBH₃ on cleavage of the unmodified DNA substrate by EcoRI (Figure 7).

Discussion

The identification of cellular targets of endogenous reactive aldehydes has garnered significant attention in an attempt to understand the genotoxic and cytotoxic mechanisms of these molecules. The products of acrolein's reaction with cellular



Figure 7. EcoRI cleavage following reduction of the cross-link. ³²Pradiolabeled γ -HOPdG-modified substrate or unmodified substrate (500 nM) was incubated with EcoRI (5 nM) in the absence of Mg²⁺ and either the presence or the absence of NaCNBH₃ for 2 h. Following preincubation, MgCl₂ (10 mM) was added to initiate cleavage. Reactions were incubated at 37 °C for various times. Cleavage profiles shown are for the 21-mer strand of the DNA duplex. Unmodified substrate, -NaCNBH₃ (**II**); unmodified substrate, +NaCNBH₃ (**II**); γ -HOPdGmodified substrate, -NaCNBH₃ (**O**); and γ -HOPdG-modified substrate, +NaCNBH₃ incubation (O).

macromolecules have been suggested to participate in diseases associated with oxidative stress (24). In addition to forming adducts to protein and adducts to DNA, acrolein induces Schiff base-mediated cross-link formation between DNA and protein (11), and the major acrolein–DNA adduct γ -HOPdG is itself reactive with both DNA and proteins (12, 16, 25-27). Thus, γ -HOPdG may represent a mode for acrolein's DNA-protein cross-link formation in vivo. Although a mechanistic link between DNA-protein cross-links and disease has not been elucidated, the formation of cross-links can be detrimental to cell survival. The bulky nature of the cross-links is likely to disrupt DNA metabolism by posing significant blocks to replication and transcription. Consistent with this proposal, aldehyde-derived DNA-protein cross-links block SV40 DNA replication (28). Although little is known about the mutagenic properties of DNA-peptide cross-links, Minko et al. recently described the mutagenic potential of NaCNBH₃-reduced $-\gamma$ -HOPdG-mediated cross-links to peptides (29).

In the present study, we demonstrated the ability of γ -HOPdG to cross-link to a protein that forms intimate interactions with DNA, the restriction endonuclease, EcoRI. Previous work has demonstrated that γ -HOPdG-containing oligonucleotides form cross-links to histone and to T4 pyrimidine dimer DNA glycosylase (*12*). It appears that γ -HOPdG forms cross-links to EcoRI more rapidly and at a considerably reduced excess of protein to DNA. The differences between the two studies are most likely attributable to the fact that EcoRI has high affinity and specificity for binding duplex oligonucleotides that contain a recognition site and that the γ -HOPdG adduct is present at the first dG residue in the recognition site.

Given that EcoRI was able to recognize and digest γ -HOPdGmodified DNA substrates, γ -HOPdG is presumed to be accommodated in the recognition interface in a manner similar to physiological substrates. Furthermore, the native structure of EcoRI was essential for rapid DNA-protein cross-link formation, and no cross-link was detected when γ -HOPdG was not located within the enzyme's recognition site. These results suggest that upon binding to the γ -HOPdG-modified cognate site, the reactive N^2 -oxopropyl moiety forms a cross-link between the 2-amino position of the modified G (NAATTC) and a proximal amino acid at the recognition interface.

Lysine residues have been implicated in DNA-protein crosslink formation by acetaldehyde (30) and may also participate in γ -HOPdG-mediated cross-links. The stabilization of the EcoRI-DNA cross-links by NaCNBH₃ is consistent with this in that NaCNBH3 preferentially reduces imine bonds formed by reaction of carbonyl groups with lysine residues (31). There are several amino acids involved in the interaction of EcoRI with its recognition sequence, which contain primary amines that could be involved in cross-link formation. Although Lys-148, Lys-117, and Lys-89 are in the vicinity, a likely candidate is Lys-113, which not only polarizes one of the oxygens of the scissile phosphate but also, along with Arg-145, may stabilize the phosphorane intermediate in the transition state (32). The Lys-113 side chain is disordered in the apoenzyme but becomes ordered upon binding DNA substrate (33). In molecular dynamics simulations, Lys-113 moves to different positions depending on the presence or absence of Mg^{2+} (32).

The impact of cross-link formation on EcoRI catalytic activity was unexpected. The conditions used for the enzymatic cleavage assay provided DNA substrate in 100-fold molar excess to EcoRI. Seemingly all of the endonuclease is involved in crosslink formation. Nevertheless, no loss of activity was observed without NaCNBH₃ reduction and stabilization of the cross-link; treatment of EcoRI alone with NaCNBH₃ did not reduce endonucleolytic activity. The loss of activity observed following treatment of the EcoRI $-\gamma$ -HOPdG-DNA complex with NaC-NBH₃ correlates to the appearance of the stable cross-link band on gel electrophoresis. Thus, it appears that the formation of the putative imine-mediated protein-DNA cross-link is readily reversible so that the protein-DNA cross-link hydrolyzes to EcoRI and γ -HOPdG-containing oligonucleotide during the time course of the incubation. Thus, no loss of enzyme activity is apparent without reductive stabilization.

The proposed mechanism for γ -HOPdG-mediated formation of DNA-protein cross-links involves ring opening of γ -HOPdG to N^2 -(γ -oxopropyl)-dG, followed by reaction with a primary amine. Kurtz and Lloyd observed that γ -HOPdG cross-links occurred only with the NH₂ termini of peptides containing lysine and arginine residues (27). This is consistent with the lower pK_a and stronger nucleophilic nature of an α-amine as compared to the ε-amine of lysine. In related chemistry, the aldehydic form of an abasic site cross-linked to the catalytically active N-terminal threonine of T4 PDG and the imine was trapped by NaCNBH₃ reduction (34). It is unlikely that the γ-HOPdG– EcoRI cross-link occurs at the NH₂ terminus of the protein. A revised model of EcoRI in solution places the N-terminal regions at a subunit interface distal to the DNA, and removal of the first four residues has no effect on catalytic activity, suggesting that the N termini are spatially remote from the EcoRI active sites (35, 36). It is more likely that cross-linking occurs to a Lys residue in the interior of the protein. Exclusion of water from the recognition interface could lower the pK_a of a Lys side chain, thereby increasing the fraction of deprotonated nucleophilic amine.

 γ -HOPdG readily formed cross-links to EcoRI, which were trappable by NaCNBH3 reduction, but the M1dG adduct did not. Although M₁dG undergoes rapid hydrolytic ring opening in duplex DNA, the aldehyde of the product, N^2 -(3-oxopropenyl)-dG, is conjugated to the exocylic amino group, which reduces its reactivity [i.e., N^2 -(3-oxopropenyl)-dG could be considered a vinylogous amide] (37). Furthermore, N^2 -(3oxopropenyl)-dG exhibits a pK_a of 6.9 so that at physiological pH it exists predominantly as the negatively charged conjugate base (37, 38). This is anticipated to significantly reduce its reactivity to nucleophiles. Thus, M1dG does not appear to be a good candidate to form cross-links with DNA binding proteins. Malondialdehyde-mediated DNA-protein cross-links have been documented in vitro and in cells treated with MDA (39). In that study, cross-linking only occurred if the protein-MDA adduct was formed first and then exposed to DNA. This behavior is consistent with the present observation that a DNA molecule containing the malondialdehyde adduct, M₁dG, does not form cross-links to EcoRI.

Although y-HOPdG-protein cross-links can be generated and stabilized in vitro, the reversibility of the cross-link may limit its effects in vivo. In COS-7 cells, replication of plasmid DNA adducted with a NaCNBH₃-reduced- γ -HOPdG-peptide crosslink resulted in approximately 8.4% mutation frequency, consisting of primarily G to T transversions, which is a higher frequency of mutation than that observed following replication of the γ -HOPdG adduct (29). The excision of DNA-protein cross-links by the UvrABC system was shown to proceed with moderate efficiency (40). Thus, secondary modifications to the adduct structure are capable of modulating the mutagenicity of γ -HOPdG and underscore the likelihood that DNA replication, transcription, or repair may be disrupted dependent on the longevity of the γ -HOPdG-protein cross-link. To date, the stability of Schiff base-mediated cross-links in vivo has not been elucidated. It will be interesting to determine what biological role, if any, y-HOPdG-mediated DNA-protein cross-links play in acrolein-mediated toxicity in vivo.

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